Supplementary Table 1: Association of Gene Ontology Categories with Decay Rate for HepG2 Experiments

These tables show details for all Gene Ontology categories. Inferences for manual classification scheme shown at the bottom.

Those categories used in Figure 1A are highlighted in bold. Standard Deviations are shown in parentheses.

P-values less than 1E-20 are indicated with a "0".

			Rate *r* (hour^-1)				Half-life < 2hr. Decay %			
GO Number	Category Name	Probe Sets	Group	Non-Group	Distribution	p-value	In-Group	Non-Group	Representation	p-value
GO:0006350	**transcription**	**1523**	**0.221 (0.009)**	**0.127 (0.002)**	**FASTER**	**0**	**13.1 (0.4)**	**4.5 (0.1)**	**OVER**	**0**
GO:0006351	transcription, DNA-dependent	1498	0.220 (0.009)	0.127 (0.002)	FASTER	0	13.0 (0.4)	4.5 (0.1)	OVER	0
GO:0006355	regulation of transcription, DNA-dependent	1163	0.230 (0.011)	0.128 (0.002)	FASTER	5.00E-21	14.2 (0.5)	4.6 (0.1)	OVER	0
GO:0006366	transcription from Pol II promoter	845	0.225 (0.012)	0.130 (0.002)	FASTER	1.88E-14	13.0 (0.5)	4.8 (0.1)	OVER	0
GO:0006139	nucleobase, nucleoside, nucleotide and nucleic acid r	3004	0.173 (0.006)	0.127 (0.002)	FASTER	1.28E-12	8.4 (0.2)	4.5 (0.1)	OVER	0
GO:0006357	regulation of transcription from Pol II promoter	487	0.231 (0.016)	0.132 (0.002)	FASTER	6.05E-10	13.5 (0.6)	4.9 (0.1)	OVER	0
GO:0008283	cell proliferation	625	0.189 (0.014)	0.132 (0.002)	FASTER	1.95E-05	10.1 (0.6)	5.0 (0.1)	OVER	1.50E-20
GO:0006513	monoubiquitination	36	0.305 (0.049)	0.134 (0.002)	FASTER	2.69E-04	25.4 (4.4)	5.1 (0.1)	OVER	2.04E-06
GO:0007050	cell cycle arrest	57	0.311 (0.054)	0.133 (0.002)	FASTER	4.83E-04	16.1 (0.7)	5.1 (0.1)	OVER	0
GO:0000074	**regulation of cell cycle**	**575**	**0.178 (0.014)**	**0.133 (0.002)**	**FASTER**	**6.18E-04**	**7.7 (0.5)**	**5.0 (0.1)**	**OVER**	**2.29E-07**
GO:0007178	transmembrane receptor protein serine/threonine kina	48	0.316 (0.057)	0.134 (0.002)	FASTER	6.58E-04	21.8 (3.8)	5.1 (0.1)	OVER	4.44E-06
GO:0007179	TGFbeta receptor signaling pathway	44	0.301 (0.055)	0.134 (0.002)	FASTER	1.19E-03	19.7 (3.8)	5.1 (0.1)	OVER	7.31E-05
GO:0042127	regulation of cell proliferation	295	0.190 (0.020)	0.133 (0.002)	FASTER	2.23E-03	12.2 (0.9)	5.0 (0.1)	OVER	2.26E-15
GO:0006596	polyamine biosynthesis	36	0.239 (0.038)	0.134 (0.002)	FASTER	3.07E-03	27.2 (1.9)	5.1 (0.1)	OVER	0
GO:0008285	negative regulation of cell proliferation	167	0.215 (0.031)	0.133 (0.002)	FASTER	3.73E-03	14.1 (1.2)	5.1 (0.1)	OVER	3.06E-15
GO:0000079	regulation of CDK activity	100	0.225 (0.035)	0.134 (0.002)	FASTER	4.81E-03	11.5 (1.4)	5.1 (0.1)	OVER	3.01E-06
GO:0006367	transcription initiation from Pol II promoter	78	0.254 (0.046)	0.133 (0.002)	FASTER	4.89E-03	14.1 (2.8)	5.1 (0.1)	OVER	5.36E-04
GO:0006915	**apoptosis**	**428**	**0.175 (0.017)**	**0.133 (0.002)**	**FASTER**	**6.35E-03**	**7.1 (0.6)**	**5.1 (0.1)**	**OVER**	**2.30E-04**
GO:0012501	programmed cell death	428	0.175 (0.017)	0.133 (0.002)	FASTER	6.35E-03	7.1 (0.6)	5.1 (0.1)	OVER	2.30E-04
GO:0006352	transcription initiation	86	0.241 (0.043)	0.134 (0.002)	FASTER	6.65E-03	12.8 (2.5)	5.1 (0.1)	OVER	0.001
GO:0000086	G2/M transition of mitotic cell cycle	112	0.214 (0.032)	0.134 (0.002)	FASTER	6.75E-03	11.9 (1.3)	5.1 (0.1)	OVER	9.23E-08
GO:0006512	ubiquitin cycle	184	0.183 (0.020)	0.134 (0.002)	FASTER	6.98E-03	7.4 (1.2)	5.1 (0.1)	NO SIG	0.025
GO:0007275	**development**	**1082**	**0.156 (0.009)**	**0.133 (0.002)**	**FASTER**	**9.16E-03**	**6.7 (0.3)**	**5.0 (0.1)**	**OVER**	**8.01E-07**
GO:0007049	cell cycle	1204	0.155 (0.009)	0.133 (0.002)	NO SIG	0.011	6.7 (0.4)	5.0 (0.1)	OVER	8.41E-06
GO:0006595	polyamine metabolism	40	0.215 (0.035)	0.134 (0.002)	NO SIG	0.011	24.5 (1.7)	5.1 (0.1)	OVER	0
GO:0019748	secondary metabolism	40	0.215 (0.035)	0.134 (0.002)	NO SIG	0.011	24.5 (1.7)	5.1 (0.1)	OVER	0
GO:0009653	morphogenesis	613	0.161 (0.012)	0.133 (0.002)	NO SIG	0.012	7.4 (0.4)	5.1 (0.1)	OVER	1.49E-07
GO:0009887	organogenesis	613	0.161 (0.012)	0.133 (0.002)	NO SIG	0.012	7.4 (0.4)	5.1 (0.1)	OVER	1.49E-07
GO:0007397	histogenesis and organogenesis	101	0.220 (0.038)	0.134 (0.002)	NO SIG	0.012	14.8 (1.3)	5.1 (0.1)	OVER	6.88E-14
GO:0008219	cell death	448	0.169 (0.016)	0.133 (0.002)	NO SIG	0.013	6.7 (0.5)	5.1 (0.1)	OVER	0.001
GO:0042401	biogenic amine biosynthesis	56	0.213 (0.036)	0.134 (0.002)	NO SIG	0.013	20.3 (1.6)	5.1 (0.1)	OVER	0
GO:0006468	protein amino acid phosphorylation	729	0.160 (0.012)	0.133 (0.002)	NO SIG	0.014	6.0 (0.5)	5.1 (0.1)	NO SIG	0.028
GO:0016265	death	455	0.167 (0.016)	0.133 (0.002)	NO SIG	0.017	6.6 (0.5)	5.1 (0.1)	OVER	0.002
GO:0016071	mRNA metabolism	43	0.228 (0.045)	0.134 (0.002)	NO SIG	0.019	5.8 (2.7)	5.1 (0.1)	NO SIG	0.409

GO:0008099	synaptic vesicle endocytosis	37	0.237 (0.052)	0.134 (0.002)	NO SIG	0.023		18.2 (2.8)	5.1 (0.1)	OVER	2.00E-06
GO:0042398	amino acid derivative biosynthesis	60	0.201 (0.034)	0.134 (0.002)	NO SIG	0.023		18.9 (1.5)	5.1 (0.1)	OVER	0
GO:0006298	mismatch repair	37	0.242 (0.055)	0.134 (0.002)	NO SIG	0.024		11.2 (3.4)	5.1 (0.1)	NO SIG	0.036
GO:0045005	maintenance of fidelity during DNA dependent DNA r	37	0.242 (0.055)	0.134 (0.002)	NO SIG	0.024		11.2 (3.4)	5.1 (0.1)	NO SIG	0.036
GO:0000278	mitotic cell cycle	719	0.156 (0.012)	0.133 (0.002)	NO SIG	0.029		6.6 (0.5)	5.1 (0.1)	OVER	0.002
GO:0000122	negative regulation of transcription from Pol II promot	100	0.201 (0.036)	0.134 (0.002)	NO SIG	0.032		15.0 (1.5)	5.1 (0.1)	OVER	1.73E-11
GO:0016481	negative regulation of transcription	100	0.201 (0.036)	0.134 (0.002)	NO SIG	0.032		15.0 (1.5)	5.1 (0.1)	OVER	1.73E-11
GO:0016310	phosphorylation	765	0.155 (0.012)	0.133 (0.002)	NO SIG	0.033		5.8 (0.5)	5.1 (0.1)	NO SIG	0.083
GO:0000082	G1/S transition of mitotic cell cycle	165	0.182 (0.026)	0.134 (0.002)	NO SIG	0.034		7.9 (1.0)	5.1 (0.1)	OVER	0.004
GO:0006511	ubiquitin-dependent protein degradation	320	0.159 (0.014)	0.134 (0.002)	NO SIG	0.039		6.1 (0.8)	5.1 (0.1)	NO SIG	0.104
GO:0000087	M phase of mitotic cell cycle	257	0.167 (0.019)	0.134 (0.002)	NO SIG	0.041		7.8 (0.9)	5.1 (0.1)	OVER	8.30E-04
GO:0007067	mitosis	253	0.166 (0.019)	0.134 (0.002)	NO SIG	0.047		8.0 (0.9)	5.1 (0.1)	OVER	6.04E-04
GO:0006796	phosphate metabolism	1021	0.150 (0.010)	0.133 (0.002)	NO SIG	0.050		5.4 (0.4)	5.1 (0.1)	NO SIG	0.207
GO:0007369	gastrulation	27	0.244 (0.070)	0.134 (0.002)	NO SIG	0.057		10.9 (4.8)	5.1 (0.1)	NO SIG	0.113
GO:0009790	embryonic development	27	0.244 (0.070)	0.134 (0.002)	NO SIG	0.057		10.9 (4.8)	5.1 (0.1)	NO SIG	0.113
GO:0009792	embryonic development (sensu Animalia)	27	0.244 (0.070)	0.134 (0.002)	NO SIG	0.057		10.9 (4.8)	5.1 (0.1)	NO SIG	0.113
GO:0019941	protein-ligand dependent protein degradation	328	0.156 (0.014)	0.134 (0.002)	NO SIG	0.058		6.0 (0.8)	5.1 (0.1)	NO SIG	0.137
GO:0006360	transcription from Pol I promoter	48	0.193 (0.038)	0.134 (0.002)	NO SIG	0.060		10.1 (2.0)	5.1 (0.1)	OVER	0.006
GO:0007093	mitotic checkpoint	32	0.204 (0.046)	0.134 (0.002)	NO SIG	0.065		4.3 (2.7)	5.1 (0.1)	NO SIG	0.383
GO:0000280	nuclear division	304	0.162 (0.018)	0.134 (0.002)	NO SIG	0.065		7.2 (0.8)	5.1 (0.1)	OVER	0.004
GO:0007020	microtubule nucleation	50	0.209 (0.050)	0.134 (0.002)	NO SIG	0.067		7.8 (1.6)	5.1 (0.1)	NO SIG	0.048
GO:0000279	M phase	320	0.160 (0.018)	0.134 (0.002)	NO SIG	0.072		6.8 (0.7)	5.1 (0.1)	NO SIG	0.011
GO:0000075	cell cycle checkpoint	71	0.191 (0.039)	0.134 (0.002)	NO SIG	0.072		5.7 (2.1)	5.1 (0.1)	NO SIG	0.390
GO:0007276	gametogenesis	140	0.173 (0.028)	0.134 (0.002)	NO SIG	0.082		8.6 (0.9)	5.1 (0.1)	OVER	9.62E-05
GO:0009888	histogenesis	86	0.184 (0.037)	0.134 (0.002)	NO SIG	0.086		7.2 (1.7)	5.1 (0.1)	NO SIG	0.109
GO:0007021	tubulin folding	32	0.224 (0.067)	0.134 (0.002)	NO SIG	0.088		10.2 (1.6)	5.1 (0.1)	OVER	7.76E-04
GO:0007323	pheromone processing	43	0.195 (0.046)	0.134 (0.002)	NO SIG	0.092		8.4 (1.5)	5.1 (0.1)	NO SIG	0.015
GO:0016485	protein processing	43	0.195 (0.046)	0.134 (0.002)	NO SIG	0.092		8.4 (1.5)	5.1 (0.1)	NO SIG	0.015
GO:0006879	iron homeostasis	40	0.199 (0.049)	0.134 (0.002)	NO SIG	0.093		12.5 (0.0)	5.1 (0.1)	OVER	0
GO:0006383	transcription from Pol III promoter	42	0.211 (0.061)	0.134 (0.002)	NO SIG	0.104		4.8 (2.3)	5.1 (0.1)	NO SIG	0.436
GO:0006974	response to DNA damage	83	0.197 (0.052)	0.134 (0.002)	NO SIG	0.111		9.9 (1.4)	5.1 (0.1)	OVER	4.96E-04
GO:0007167	enzyme linked receptor protein signaling pathway	173	0.161 (0.023)	0.134 (0.002)	NO SIG	0.113		7.6 (1.2)	5.1 (0.1)	NO SIG	0.021
GO:0009611	response to wounding	147	0.169 (0.029)	0.134 (0.002)	NO SIG	0.114		8.8 (1.2)	5.1 (0.1)	OVER	0.001
GO:0008625	induction of apoptosis via death domain receptors	35	0.206 (0.060)	0.134 (0.002)	NO SIG	0.116		6.4 (2.1)	5.1 (0.1)	NO SIG	0.283
GO:0007089	start control point of mitotic cell cycle	32	0.216 (0.069)	0.134 (0.002)	NO SIG	0.117		6.2 (3.4)	5.1 (0.1)	NO SIG	0.374
GO:0001501	skeletal development	122	0.167 (0.029)	0.134 (0.002)	NO SIG	0.124		10.6 (1.0)	5.1 (0.1)	OVER	5.89E-08
GO:0000003	reproduction	165	0.162 (0.025)	0.134 (0.002)	NO SIG	0.125		7.4 (0.8)	5.1 (0.1)	OVER	0.003
GO:0007324	a-factor processing (proteolytic)	35	0.197 (0.055)	0.134 (0.002)	NO SIG	0.127		10.3 (1.9)	5.1 (0.1)	OVER	0.003
GO:0009292	genetic exchange	55	0.178 (0.040)	0.134 (0.002)	NO SIG	0.130		6.6 (1.2)	5.1 (0.1)	NO SIG	0.111
GO:0007283	spermatogenesis	94	0.175 (0.037)	0.134 (0.002)	NO SIG	0.131		10.5 (1.0)	5.1 (0.1)	OVER	2.23E-08
GO:0008371	obsolete	436	0.152 (0.017)	0.134 (0.002)	NO SIG	0.135		6.4 (0.6)	5.1 (0.1)	NO SIG	0.019
GO:0006464	**protein modification**	**1436**	**0.143 (0.008)**	**0.133 (0.002)**	**NO SIG**	**0.135**		**4.7 (0.3)**	**5.2 (0.1)**	**NO SIG**	**0.086**

GO:0006261	DNA dependent DNA replication	171	0.161 (0.025)	0.134 (0.002)	NO SIG	0.141	5.0 (1.1)	5.1 (0.1)	NO SIG	0.437
GO:0006576	biogenic amine metabolism	75	0.165 (0.029)	0.134 (0.002)	NO SIG	0.141	15.2 (1.2)	5.1 (0.1)	OVER	4.43E-18
GO:0008284	positive regulation of cell proliferation	128	0.157 (0.023)	0.134 (0.002)	NO SIG	0.153	9.7 (1.5)	5.1 (0.1)	OVER	8.06E-04
GO:0006947	cell-cell fusion	47	0.177 (0.043)	0.134 (0.002)	NO SIG	0.161	7.7 (1.4)	5.1 (0.1)	NO SIG	0.032
GO:0007322	mating (sensu Saccharomyces)	47	0.177 (0.043)	0.134 (0.002)	NO SIG	0.161	7.7 (1.4)	5.1 (0.1)	NO SIG	0.032
GO:0030461	mating (sensu Fungi)	47	0.177 (0.043)	0.134 (0.002)	NO SIG	0.161	7.7 (1.4)	5.1 (0.1)	NO SIG	0.032
GO:0006954	inflammatory response	98	0.169 (0.036)	0.134 (0.002)	NO SIG	0.166	8.1 (1.3)	5.1 (0.1)	NO SIG	0.013
GO:0045087	innate immune response	98	0.169 (0.036)	0.134 (0.002)	NO SIG	0.166	8.1 (1.3)	5.1 (0.1)	NO SIG	0.013
GO:0000244	snRNP recycling	31	0.190 (0.059)	0.134 (0.002)	NO SIG	0.168	5.9 (3.2)	5.1 (0.1)	NO SIG	0.402
GO:0006950	stress response	719	0.146 (0.013)	0.134 (0.002)	NO SIG	0.172	6.9 (0.5)	5.1 (0.1)	OVER	6.01E-05
GO:0007345	embryogenesis and morphogenesis	129	0.165 (0.033)	0.134 (0.002)	NO SIG	0.174	7.0 (1.5)	5.1 (0.1)	NO SIG	0.101
GO:0007398	ectoderm development	49	0.172 (0.042)	0.134 (0.002)	NO SIG	0.182	6.5 (1.3)	5.1 (0.1)	NO SIG	0.133
GO:0007257	activation of JUN kinase	29	0.197 (0.071)	0.134 (0.002)	NO SIG	0.188	2.5 (2.7)	5.1 (0.1)	NO SIG	0.170
GO:0006397	**mRNA processing**	**314**	**0.148 (0.016)**	**0.134 (0.002)**	**NO SIG**	**0.196**	**3.9 (0.7)**	**5.1 (0.1)**	**NO SIG**	**0.026**
GO:0006470	protein amino acid dephosphorylation	344	0.148 (0.017)	0.134 (0.002)	NO SIG	0.198	5.2 (0.6)	5.1 (0.1)	NO SIG	0.434
GO:0016311	dephosphorylation	344	0.148 (0.017)	0.134 (0.002)	NO SIG	0.198	5.2 (0.6)	5.1 (0.1)	NO SIG	0.434
GO:0016181	synaptic vesicle transport	46	0.178 (0.053)	0.134 (0.002)	NO SIG	0.204	14.7 (2.3)	5.1 (0.1)	OVER	1.63E-05
GO:0006401	RNA catabolism	35	0.166 (0.041)	0.134 (0.002)	NO SIG	0.221	2.4 (2.1)	5.1 (0.1)	NO SIG	0.093
GO:0006270	DNA replication initiation	39	0.169 (0.046)	0.134 (0.002)	NO SIG	0.224	6.9 (2.6)	5.1 (0.1)	NO SIG	0.241
GO:0007254	JNK cascade	55	0.175 (0.054)	0.134 (0.002)	NO SIG	0.224	4.5 (2.3)	5.1 (0.1)	NO SIG	0.393
GO:0007399	neurogenesis	275	0.147 (0.018)	0.134 (0.002)	NO SIG	0.227	6.4 (0.5)	5.1 (0.1)	OVER	0.006
GO:0006514	deubiquitination	54	0.165 (0.042)	0.134 (0.002)	NO SIG	0.229	4.6 (2.0)	5.1 (0.1)	NO SIG	0.392
GO:0006897	endocytosis	118	0.152 (0.027)	0.134 (0.002)	NO SIG	0.249	10.1 (1.2)	5.1 (0.1)	OVER	7.09E-06
GO:0007088	regulation of mitosis	68	0.161 (0.040)	0.134 (0.002)	NO SIG	0.250	2.6 (1.5)	5.1 (0.1)	NO SIG	0.049
GO:0006726	eye pigment biosynthesis	28	0.168 (0.051)	0.134 (0.002)	NO SIG	0.253	8.3 (3.8)	5.1 (0.1)	NO SIG	0.205
GO:0042441	eye pigment metabolism	28	0.168 (0.051)	0.134 (0.002)	NO SIG	0.253	8.3 (3.8)	5.1 (0.1)	NO SIG	0.205
GO:0008632	apoptotic program	74	0.160 (0.040)	0.134 (0.002)	NO SIG	0.256	11.3 (1.0)	5.1 (0.1)	OVER	6.99E-10
GO:0006281	DNA repair	355	0.145 (0.018)	0.134 (0.002)	NO SIG	0.262	6.4 (0.7)	5.1 (0.1)	NO SIG	0.026
GO:0008544	epidermal differentiation	42	0.159 (0.040)	0.134 (0.002)	NO SIG	0.263	3.0 (1.3)	5.1 (0.1)	NO SIG	0.042
GO:0042254	ribosome biogenesis and assembly	60	0.153 (0.030)	0.134 (0.002)	NO SIG	0.264	6.4 (1.6)	5.1 (0.1)	NO SIG	0.198
GO:0006626	protein-mitochondrial targeting	34	0.158 (0.041)	0.134 (0.002)	NO SIG	0.278	0.3 (1.0)	5.1 (0.1)	UNDER	3.23E-07
GO:0006628	mitochondrial translocation	31	0.158 (0.040)	0.134 (0.002)	NO SIG	0.278	0.3 (1.0)	5.1 (0.1)	UNDER	3.29E-07
GO:0007046	ribosome biogenesis	57	0.150 (0.028)	0.134 (0.002)	NO SIG	0.278	6.6 (1.5)	5.1 (0.1)	NO SIG	0.171
GO:0016070	RNA metabolism	553	0.141 (0.012)	0.134 (0.002)	NO SIG	0.286	3.9 (0.5)	5.2 (0.1)	UNDER	0.005
GO:0006968	cellular defense response	40	0.158 (0.047)	0.134 (0.002)	NO SIG	0.304	9.8 (2.9)	5.1 (0.1)	NO SIG	0.052
GO:0030006	heavy metal ion homeostasis	61	0.152 (0.036)	0.134 (0.002)	NO SIG	0.306	8.2 (0.0)	5.1 (0.1)	OVER	0
GO:0000080	G1 phase of mitotic cell cycle	57	0.157 (0.050)	0.134 (0.002)	NO SIG	0.318	5.8 (2.1)	5.1 (0.1)	NO SIG	0.381
GO:0000084	S phase of mitotic cell cycle	295	0.143 (0.019)	0.134 (0.002)	NO SIG	0.321	5.0 (0.8)	5.1 (0.1)	NO SIG	0.419
GO:0006283	transcription-coupled nucleotide-excision repair	29	0.165 (0.070)	0.134 (0.002)	NO SIG	0.329	2.0 (2.4)	5.1 (0.1)	NO SIG	0.096
GO:0008624	induction of apoptosis by extracellular signals	73	0.149 (0.038)	0.134 (0.002)	NO SIG	0.347	4.2 (1.5)	5.1 (0.1)	NO SIG	0.260
GO:0006913	nucleocytoplasmic transport	162	0.142 (0.023)	0.134 (0.002)	NO SIG	0.352	5.2 (0.9)	5.1 (0.1)	NO SIG	0.473
GO:0016482	cytoplasmic transport	162	0.142 (0.023)	0.134 (0.002)	NO SIG	0.352	5.2 (0.9)	5.1 (0.1)	NO SIG	0.473

GO:0007017	microtubule-based process	197	0.141 (0.020)	0.134 (0.002)	NO SIG	0.357	4.0 (0.8)	5.1 (0.1)	NO SIG	0.072
GO:0009314	radiation response	85	0.146 (0.036)	0.134 (0.002)	NO SIG	0.366	5.7 (1.0)	5.1 (0.1)	NO SIG	0.283
GO:0006951	response to heat shock	116	0.143 (0.027)	0.134 (0.002)	NO SIG	0.369	4.7 (0.8)	5.1 (0.1)	NO SIG	0.286
GO:0019236	pheromone response	71	0.145 (0.033)	0.134 (0.002)	NO SIG	0.370	5.1 (0.9)	5.1 (0.1)	NO SIG	0.481
GO:0006396	RNA processing	505	0.138 (0.013)	0.134 (0.002)	NO SIG	0.373	3.9 (0.5)	5.2 (0.1)	UNDER	0.005
GO:0007420	brain development	28	0.153 (0.062)	0.134 (0.002)	NO SIG	0.379	3.5 (2.0)	5.1 (0.1)	NO SIG	0.202
GO:0006917	induction of apoptosis	156	0.143 (0.030)	0.134 (0.002)	NO SIG	0.380	2.7 (1.0)	5.1 (0.1)	UNDER	0.006
GO:0012502	induction of programmed cell death	156	0.143 (0.030)	0.134 (0.002)	NO SIG	0.380	2.7 (1.0)	5.1 (0.1)	UNDER	0.006
GO:0006620	post-translational membrane targeting	51	0.146 (0.043)	0.134 (0.002)	NO SIG	0.390	6.1 (0.9)	5.1 (0.1)	NO SIG	0.126
GO:0000226	microtubule cytoskeleton organization and biogenesis	102	0.141 (0.028)	0.134 (0.002)	NO SIG	0.396	5.1 (1.1)	5.1 (0.1)	NO SIG	0.491
GO:0006607	NLS-bearing substrate-nucleus import	38	0.149 (0.056)	0.134 (0.002)	NO SIG	0.396	9.0 (3.2)	5.1 (0.1)	NO SIG	0.109
GO:0006291	pyrimidine-dimer repair\, DNA damage excision	33	0.146 (0.061)	0.134 (0.002)	NO SIG	0.422	1.8 (2.1)	5.1 (0.1)	NO SIG	0.054
GO:0007156	homophilic cell adhesion	26	0.140 (0.052)	0.134 (0.002)	NO SIG	0.450	4.9 (3.3)	5.1 (0.1)	NO SIG	0.469
GO:0006289	nucleotide-excision repair	62	0.140 (0.046)	0.134 (0.002)	NO SIG	0.452	2.1 (1.6)	5.1 (0.1)	NO SIG	0.028
GO:0007417	central nervous system development	83	0.138 (0.037)	0.134 (0.002)	NO SIG	0.457	5.1 (1.0)	5.1 (0.1)	NO SIG	0.475
GO:0007269	neurotransmitter release	59	0.139 (0.045)	0.134 (0.002)	NO SIG	0.458	11.5 (1.8)	5.1 (0.1)	OVER	0.00021
GO:0008380	RNA splicing	246	0.136 (0.019)	0.134 (0.002)	NO SIG	0.464	2.9 (0.8)	5.2 (0.1)	UNDER	0.001
GO:0000165	MAPKKK cascade	86	0.137 (0.030)	0.134 (0.002)	NO SIG	0.465	2.6 (1.1)	5.1 (0.1)	NO SIG	0.013
GO:0016051	carbohydrate biosynthesis	74	0.136 (0.031)	0.134 (0.002)	NO SIG	0.471	1.4 (1.2)	5.1 (0.1)	UNDER	0.001
GO:0001505	neurotransmitter maintenance	67	0.137 (0.041)	0.134 (0.002)	NO SIG	0.475	10.9 (1.8)	5.1 (0.1)	OVER	5.38E-04
GO:0006612	protein-membrane targeting	83	0.136 (0.032)	0.134 (0.002)	NO SIG	0.480	4.3 (0.9)	5.1 (0.1)	NO SIG	0.165
GO:0030005	di-\, tri-valent inorganic cation homeostasis	71	0.135 (0.033)	0.134 (0.002)	NO SIG	0.482	7.0 (0.0)	5.1 (0.1)	OVER	0
GO:0007126	meiosis	53	0.136 (0.057)	0.134 (0.002)	NO SIG	0.482	3.2 (1.5)	5.1 (0.1)	NO SIG	0.102
GO:0007606	chemosensory perception	80	0.135 (0.029)	0.134 (0.002)	NO SIG	0.492	4.5 (0.8)	5.1 (0.1)	NO SIG	0.225
GO:0009593	perception of chemical substance	80	0.135 (0.029)	0.134 (0.002)	NO SIG	0.492	4.5 (0.8)	5.1 (0.1)	NO SIG	0.225
GO:0007031	peroxisome organization and biogenesis	42	0.135 (0.047)	0.134 (0.002)	NO SIG	0.492	0.4 (0.9)	5.1 (0.1)	UNDER	9.98E-08
GO:0006875	metal ion homeostasis	75	0.134 (0.032)	0.134 (0.002)	NO SIG	0.500	6.7 (0.1)	5.1 (0.1)	OVER	0
GO:0006575	amino acid derivative metabolism	94	0.134 (0.026)	0.134 (0.002)	NO SIG	0.496	12.1 (0.9)	5.1 (0.1)	OVER	4.00E-14
GO:0016337	cell-cell adhesion	27	0.133 (0.052)	0.134 (0.002)	NO SIG	0.495	4.7 (3.2)	5.1 (0.1)	NO SIG	0.445
GO:0006953	acute-phase response	47	0.133 (0.027)	0.134 (0.002)	NO SIG	0.491	8.5 (0.0)	5.1 (0.1)	OVER	0
GO:0006471	protein amino acid ADP-ribosylation	52	0.133 (0.039)	0.134 (0.002)	NO SIG	0.487	1.5 (1.6)	5.1 (0.1)	NO SIG	0.010
GO:0007292	oogenesis	27	0.132 (0.048)	0.134 (0.002)	NO SIG	0.484	1.6 (2.3)	5.1 (0.1)	NO SIG	0.061
GO:0000067	DNA replication and chromosome cycle	333	0.133 (0.019)	0.134 (0.002)	NO SIG	0.484	5.6 (0.8)	5.1 (0.1)	NO SIG	0.290
GO:0006371	mRNA splicing	228	0.133 (0.020)	0.134 (0.002)	NO SIG	0.483	2.8 (0.8)	5.2 (0.1)	UNDER	0.001
GO:0007166	cell surface receptor linked signal transduction	519	0.133 (0.014)	0.134 (0.002)	NO SIG	0.483	5.6 (0.6)	5.1 (0.1)	NO SIG	0.181
GO:0006260	DNA replication	272	0.133 (0.020)	0.134 (0.002)	NO SIG	0.478	4.6 (0.8)	5.1 (0.1)	NO SIG	0.251
GO:0000209	polyubiquitination	41	0.131 (0.039)	0.134 (0.002)	NO SIG	0.473	2.3 (1.4)	5.1 (0.1)	NO SIG	0.021
GO:0006935	chemotaxis	64	0.130 (0.042)	0.134 (0.002)	NO SIG	0.467	4.0 (1.2)	5.1 (0.1)	NO SIG	0.174
GO:0042330	taxis	64	0.130 (0.042)	0.134 (0.002)	NO SIG	0.467	4.0 (1.2)	5.1 (0.1)	NO SIG	0.174
GO:0006869	lipid transport	38	0.131 (0.029)	0.134 (0.002)	NO SIG	0.461	0.5 (1.1)	5.1 (0.1)	UNDER	1.14E-05
GO:0006960	antimicrobial humoral response (sensu Invertebrata)	68	0.129 (0.043)	0.134 (0.002)	NO SIG	0.455	3.7 (1.5)	5.1 (0.1)	NO SIG	0.172
GO:0016065	humoral defense mechanism (sensu Invertebrata)	68	0.129 (0.043)	0.134 (0.002)	NO SIG	0.455	3.7 (1.5)	5.1 (0.1)	NO SIG	0.172

GO:0019730	antimicrobial humoral response	68	0.129 (0.043)	0.134 (0.002)	NO SIG	0.455	3.7 (1.5)	5.1 (0.1)	NO SIG	0.172
GO:0006333	chromatin assembly/disassembly	96	0.131 (0.026)	0.134 (0.002)	NO SIG	0.453	4.6 (0.9)	5.1 (0.1)	NO SIG	0.270
GO:0006975	DNA damage induced protein phosphorylation	30	0.126 (0.063)	0.134 (0.002)	NO SIG	0.447	0.5 (1.3)	5.1 (0.1)	UNDER	1.22E-04
GO:0008629	induction of apoptosis by intracellular signals	39	0.125 (0.063)	0.134 (0.002)	NO SIG	0.442	2.0 (2.0)	5.1 (0.1)	NO SIG	0.062
GO:0009266	temperature response	128	0.130 (0.025)	0.134 (0.002)	NO SIG	0.436	4.2 (0.7)	5.1 (0.1)	NO SIG	0.109
GO:0007165	**signal transduction**	**1834**	**0.133 (0.008)**	**0.134 (0.002)**	**NO SIG**	**0.431**	**4.8 (0.3)**	**5.2 (0.1)**	**NO SIG**	**0.122**
GO:0007267	cell-cell signaling	284	0.129 (0.019)	0.134 (0.002)	NO SIG	0.404	7.7 (0.7)	5.1 (0.1)	OVER	2.90E-04
GO:0007004	telomere maintenance	35	0.110 (0.094)	0.134 (0.002)	NO SIG	0.401	2.0 (2.1)	5.1 (0.1)	NO SIG	0.066
GO:0007188	G-protein signaling\, coupled to cAMP nucleotide sec	43	0.123 (0.043)	0.134 (0.002)	NO SIG	0.398	1.9 (1.8)	5.1 (0.1)	NO SIG	0.037
GO:0019933	cAMP-mediated signaling	43	0.123 (0.043)	0.134 (0.002)	NO SIG	0.398	1.9 (1.8)	5.1 (0.1)	NO SIG	0.037
GO:0006873	ion homeostasis	83	0.126 (0.030)	0.134 (0.002)	NO SIG	0.395	6.0 (0.1)	5.1 (0.1)	OVER	3.60E-18
GO:0019725	homeostasis	83	0.126 (0.030)	0.134 (0.002)	NO SIG	0.395	6.0 (0.1)	5.1 (0.1)	OVER	3.60E-18
GO:0030003	cation homeostasis	83	0.126 (0.030)	0.134 (0.002)	NO SIG	0.395	6.0 (0.1)	5.1 (0.1)	OVER	3.60E-18
GO:0006606	protein-nucleus import	71	0.123 (0.036)	0.134 (0.002)	NO SIG	0.385	4.9 (1.7)	5.1 (0.1)	NO SIG	0.456
GO:0006916	anti-apoptosis	81	0.124 (0.033)	0.134 (0.002)	NO SIG	0.384	3.8 (1.5)	5.1 (0.1)	NO SIG	0.195
GO:0007010	cytoskeleton organization and biogenesis	330	0.129 (0.015)	0.134 (0.002)	NO SIG	0.383	2.8 (0.6)	5.2 (0.1)	UNDER	1.63E-05
GO:0006613	co-translational membrane targeting	31	0.119 (0.047)	0.134 (0.002)	NO SIG	0.378	1.3 (1.9)	5.1 (0.1)	NO SIG	0.024
GO:0007610	behavior	60	0.122 (0.038)	0.134 (0.002)	NO SIG	0.376	3.3 (1.2)	5.1 (0.1)	NO SIG	0.060
GO:0000245	spliceosome assembly	33	0.114 (0.056)	0.134 (0.002)	NO SIG	0.357	4.3 (2.4)	5.1 (0.1)	NO SIG	0.356
GO:0007059	chromosome segregation	28	0.104 (0.080)	0.134 (0.002)	NO SIG	0.352	8.8 (3.6)	5.1 (0.1)	NO SIG	0.155
GO:0007127	meiosis I	31	0.102 (0.082)	0.134 (0.002)	NO SIG	0.349	0.8 (1.6)	5.1 (0.1)	UNDER	0.004
GO:0007128	meiotic prophase I	31	0.102 (0.082)	0.134 (0.002)	NO SIG	0.349	0.8 (1.6)	5.1 (0.1)	UNDER	0.004
GO:0006259	**DNA metabolism**	**875**	**0.130 (0.011)**	**0.134 (0.002)**	**NO SIG**	**0.347**	**4.6 (0.4)**	**5.2 (0.1)**	**NO SIG**	**0.106**
GO:0006505	GPI anchor metabolism	28	0.114 (0.048)	0.134 (0.002)	NO SIG	0.340	1.8 (1.8)	5.1 (0.1)	NO SIG	0.033
GO:0006506	GPI anchor biosynthesis	28	0.114 (0.048)	0.134 (0.002)	NO SIG	0.340	1.8 (1.8)	5.1 (0.1)	NO SIG	0.033
GO:0042079	GPI/GSI anchor metabolism	28	0.114 (0.048)	0.134 (0.002)	NO SIG	0.340	1.8 (1.8)	5.1 (0.1)	NO SIG	0.033
GO:0042080	GPI/GSI anchor biosynthesis	28	0.114 (0.048)	0.134 (0.002)	NO SIG	0.340	1.8 (1.8)	5.1 (0.1)	NO SIG	0.033
GO:0007028	cytoplasm organization and biogenesis	552	0.129 (0.012)	0.134 (0.002)	NO SIG	0.325	3.2 (0.4)	5.2 (0.1)	UNDER	1.59E-06
GO:0007601	vision	58	0.112 (0.046)	0.134 (0.002)	NO SIG	0.320	1.5 (1.4)	5.1 (0.1)	UNDER	0.006
GO:0009583	perception of light	58	0.112 (0.046)	0.134 (0.002)	NO SIG	0.320	1.5 (1.4)	5.1 (0.1)	UNDER	0.006
GO:0006996	organelle organization and biogenesis	480	0.128 (0.013)	0.134 (0.002)	NO SIG	0.315	2.9 (0.4)	5.2 (0.1)	UNDER	1.02E-07
GO:0009057	macromolecule catabolism	692	0.129 (0.010)	0.134 (0.002)	NO SIG	0.305	3.8 (0.4)	5.2 (0.1)	UNDER	6.91E-04
GO:0009416	light response	73	0.114 (0.039)	0.134 (0.002)	NO SIG	0.301	1.2 (1.1)	5.1 (0.1)	UNDER	3.45E-04
GO:0042440	pigment metabolism	61	0.117 (0.032)	0.134 (0.002)	NO SIG	0.296	3.8 (1.7)	5.1 (0.1)	NO SIG	0.222
GO:0000183	chromatin silencing at ribosomal DNA (rDNA)	50	0.107 (0.050)	0.134 (0.002)	NO SIG	0.294	2.5 (1.8)	5.1 (0.1)	NO SIG	0.068
GO:0006342	chromatin silencing	50	0.107 (0.050)	0.134 (0.002)	NO SIG	0.294	2.5 (1.8)	5.1 (0.1)	NO SIG	0.068
GO:0016440	transcriptional gene silencing	50	0.107 (0.050)	0.134 (0.002)	NO SIG	0.294	2.5 (1.8)	5.1 (0.1)	NO SIG	0.068
GO:0016458	gene silencing	50	0.107 (0.050)	0.134 (0.002)	NO SIG	0.294	2.5 (1.8)	5.1 (0.1)	NO SIG	0.068
GO:0009613	response to pest/pathogen/parasite	364	0.125 (0.017)	0.134 (0.002)	NO SIG	0.293	5.2 (0.7)	5.1 (0.1)	NO SIG	0.431
GO:0009315	drug resistance	37	0.102 (0.056)	0.134 (0.002)	NO SIG	0.284	0.3 (0.8)	5.1 (0.1)	UNDER	2.18E-09
GO:0016194	non-selective vesicle exocytosis	27	0.101 (0.057)	0.134 (0.002)	NO SIG	0.283	3.1 (2.2)	5.1 (0.1)	NO SIG	0.180
GO:0006813	potassium transport	32	0.109 (0.041)	0.134 (0.002)	NO SIG	0.275	0.2 (0.8)	5.1 (0.1)	UNDER	5.50E-09

GO:0006418	amino acid activation	85	0.119 (0.023)	0.134 (0.002)	NO SIG	0.262	0.5 (0.7)	5.1 (0.1)	UNDER	5.04E-10
GO:0007266	Rho protein signal transduction	46	0.106 (0.042)	0.134 (0.002)	NO SIG	0.256	5.7 (1.7)	5.1 (0.1)	NO SIG	0.379
GO:0005976	polysaccharide metabolism	68	0.112 (0.033)	0.134 (0.002)	NO SIG	0.249	0.6 (0.8)	5.1 (0.1)	UNDER	2.12E-08
GO:0007600	sensory perception	154	0.117 (0.024)	0.134 (0.002)	NO SIG	0.248	2.9 (0.7)	5.1 (0.1)	UNDER	6.96E-04
GO:0009066	aspartate family amino acid metabolism	30	0.095 (0.057)	0.134 (0.002)	NO SIG	0.248	2.1 (2.3)	5.1 (0.1)	NO SIG	0.093
GO:0006323	DNA packaging	271	0.122 (0.018)	0.134 (0.002)	NO SIG	0.248	4.0 (0.7)	5.1 (0.1)	NO SIG	0.045
GO:0006508	proteolysis and peptidolysis	657	0.127 (0.010)	0.134 (0.002)	NO SIG	0.247	3.8 (0.4)	5.2 (0.1)	UNDER	0.002
GO:0030163	protein degradation	657	0.127 (0.010)	0.134 (0.002)	NO SIG	0.247	3.8 (0.4)	5.2 (0.1)	UNDER	0.002
GO:0009582	perception of abiotic stimulus	158	0.117 (0.024)	0.134 (0.002)	NO SIG	0.243	2.9 (0.7)	5.1 (0.1)	UNDER	3.87E-04
GO:0009100	glycoprotein metabolism	120	0.117 (0.024)	0.134 (0.002)	NO SIG	0.243	3.6 (0.4)	5.1 (0.1)	UNDER	1.51E-04
GO:0006650	glycerophospholipid metabolism	33	0.099 (0.050)	0.134 (0.002)	NO SIG	0.240	1.9 (1.5)	5.1 (0.1)	NO SIG	0.020
GO:0007186	G-protein coupled receptor protein signaling pathway	207	0.117 (0.023)	0.134 (0.002)	NO SIG	0.239	5.6 (0.8)	5.1 (0.1)	NO SIG	0.288
GO:0006605	protein targeting	253	0.121 (0.018)	0.134 (0.002)	NO SIG	0.234	3.0 (0.6)	5.2 (0.1)	UNDER	2.31E-04
GO:0006997	nuclear organization and biogenesis	324	0.121 (0.019)	0.134 (0.002)	NO SIG	0.234	4.0 (0.6)	5.1 (0.1)	NO SIG	0.033
GO:0008643	carbohydrate transport	28	0.092 (0.058)	0.134 (0.002)	NO SIG	0.233	0.1 (0.6)	5.1 (0.1)	UNDER	4.40E-19
GO:0007265	RAS protein signal transduction	47	0.094 (0.053)	0.134 (0.002)	NO SIG	0.226	4.4 (2.1)	5.1 (0.1)	NO SIG	0.354
GO:0006461	protein complex assembly	145	0.114 (0.026)	0.134 (0.002)	NO SIG	0.222	2.4 (0.7)	5.1 (0.1)	UNDER	6.23E-05
GO:0006486	protein amino acid glycosylation	108	0.115 (0.023)	0.134 (0.002)	NO SIG	0.210	3.8 (0.3)	5.1 (0.1)	UNDER	2.87E-06
GO:0009101	glycoprotein biosynthesis	108	0.115 (0.023)	0.134 (0.002)	NO SIG	0.210	3.8 (0.3)	5.1 (0.1)	UNDER	2.87E-06
GO:0007517	muscle development	112	0.111 (0.029)	0.134 (0.002)	NO SIG	0.208	3.4 (1.0)	5.1 (0.1)	NO SIG	0.050
GO:0006919	caspase activation	45	0.086 (0.059)	0.134 (0.002)	NO SIG	0.207	3.0 (1.7)	5.1 (0.1)	NO SIG	0.104
GO:0007338	fertilization (sensu Animalia)	29	0.087 (0.056)	0.134 (0.002)	NO SIG	0.201	0.3 (0.9)	5.1 (0.1)	UNDER	1.74E-07
GO:0009566	fertilization	29	0.087 (0.056)	0.134 (0.002)	NO SIG	0.201	0.3 (0.9)	5.1 (0.1)	UNDER	1.74E-07
GO:0040029	regulation of gene expression\, epigenetic	29	0.099 (0.042)	0.134 (0.002)	NO SIG	0.201	0.0 (0.3)	5.1 (0.1)	UNDER	0
GO:0006325	establishment and/or maintenance of chromatin archi	230	0.118 (0.019)	0.134 (0.002)	NO SIG	0.197	3.8 (0.7)	5.1 (0.1)	NO SIG	0.024
GO:0007422	peripheral nervous system development	30	0.089 (0.051)	0.134 (0.002)	NO SIG	0.189	0.4 (1.1)	5.1 (0.1)	UNDER	4.25E-06
GO:0007243	protein kinase cascade	196	0.113 (0.023)	0.134 (0.002)	NO SIG	0.185	2.4 (0.7)	5.2 (0.1)	UNDER	6.04E-05
GO:0007125	invasive growth	29	0.083 (0.056)	0.134 (0.002)	NO SIG	0.182	0.1 (0.7)	5.1 (0.1)	UNDER	8.73E-13
GO:0007149	colony morphology	29	0.083 (0.056)	0.134 (0.002)	NO SIG	0.182	0.1 (0.7)	5.1 (0.1)	UNDER	8.73E-13
GO:0006814	sodium transport	44	0.098 (0.040)	0.134 (0.002)	NO SIG	0.180	0.2 (0.6)	5.1 (0.1)	UNDER	9.72E-15
GO:0007001	chromosome organization and biogenesis (sensu Euk	308	0.116 (0.019)	0.134 (0.002)	NO SIG	0.173	3.7 (0.6)	5.2 (0.1)	UNDER	0.007
GO:0006730	one-carbon compound metabolism	30	0.078 (0.059)	0.134 (0.002)	NO SIG	0.168	0.3 (0.9)	5.1 (0.1)	UNDER	1.27E-07
GO:0008203	cholesterol metabolism	90	0.109 (0.025)	0.134 (0.002)	NO SIG	0.166	6.3 (1.1)	5.1 (0.1)	NO SIG	0.151
GO:0009129	pyrimidine nucleoside monophosphate metabolism	45	0.104 (0.031)	0.134 (0.002)	NO SIG	0.165	4.9 (1.6)	5.1 (0.1)	NO SIG	0.448
GO:0009309	amine biosynthesis	117	0.111 (0.024)	0.134 (0.002)	NO SIG	0.165	9.7 (0.8)	5.1 (0.1)	OVER	5.06E-10
GO:0016043	cell organization and biogenesis	886	0.124 (0.010)	0.134 (0.002)	NO SIG	0.164	3.5 (0.3)	5.2 (0.1)	UNDER	4.04E-07
GO:0005977	glycogen metabolism	65	0.102 (0.032)	0.134 (0.002)	NO SIG	0.161	0.1 (0.4)	5.1 (0.1)	UNDER	0
GO:0007187	G-protein signaling\, coupled to cyclic nucleotide seco	56	0.097 (0.038)	0.134 (0.002)	NO SIG	0.160	1.4 (1.4)	5.1 (0.1)	UNDER	0.004
GO:0019932	second-messenger-mediated signaling	56	0.097 (0.038)	0.134 (0.002)	NO SIG	0.160	1.4 (1.4)	5.1 (0.1)	UNDER	0.004
GO:0019935	cyclic-nucleotide-mediated signaling	56	0.097 (0.038)	0.134 (0.002)	NO SIG	0.160	1.4 (1.4)	5.1 (0.1)	UNDER	0.004
GO:0009112	nucleobase metabolism	44	0.093 (0.041)	0.134 (0.002)	NO SIG	0.157	5.2 (1.7)	5.1 (0.1)	NO SIG	0.483
GO:0006959	humoral immune response	126	0.104 (0.030)	0.134 (0.002)	NO SIG	0.155	3.3 (1.1)	5.1 (0.1)	NO SIG	0.049

GO:0018193	peptidyl-amino acid modification	32	0.058 (0.072)	0.134 (0.002)	NO SIG	0.144	0.9 (1.6)	5.1 (0.1)	UNDER	0.003
GO:0018212	peptidyl-tyrosine modification	32	0.058 (0.072)	0.134 (0.002)	NO SIG	0.144	0.9 (1.6)	5.1 (0.1)	UNDER	0.003
GO:0007268	synaptic transmission	132	0.105 (0.026)	0.134 (0.002)	NO SIG	0.136	5.6 (0.9)	5.1 (0.1)	NO SIG	0.296
GO:0042445	hormone metabolism	49	0.085 (0.043)	0.134 (0.002)	NO SIG	0.129	3.3 (1.1)	5.1 (0.1)	NO SIG	0.048
GO:0007012	actin cytoskeleton reorganization	42	0.086 (0.041)	0.134 (0.002)	NO SIG	0.120	0.6 (1.1)	5.1 (0.1)	UNDER	1.65E-05
GO:0030203	glycosaminoglycan metabolism	39	0.082 (0.044)	0.134 (0.002)	NO SIG	0.118	1.8 (1.9)	5.1 (0.1)	NO SIG	0.036
GO:0006022	aminoglycan metabolism	43	0.085 (0.041)	0.134 (0.002)	NO SIG	0.115	1.6 (1.7)	5.1 (0.1)	NO SIG	0.019
GO:0006665	sphingolipid metabolism	28	0.070 (0.053)	0.134 (0.002)	NO SIG	0.114	0.3 (1.0)	5.1 (0.1)	UNDER	1.58E-06
GO:0006899	non-selective vesicle transport	128	0.099 (0.029)	0.134 (0.002)	NO SIG	0.112	4.9 (0.9)	5.1 (0.1)	NO SIG	0.389
GO:0007262	STAT protein nuclear translocation	33	0.052 (0.067)	0.134 (0.002)	NO SIG	0.108	0.4 (1.1)	5.1 (0.1)	UNDER	7.57E-06
GO:0006112	energy reserve metabolism	69	0.095 (0.031)	0.134 (0.002)	NO SIG	0.108	0.1 (0.4)	5.1 (0.1)	UNDER	0
GO:0009060	aerobic respiration	27	0.083 (0.040)	0.134 (0.002)	NO SIG	0.104	0.0 (0.2)	5.1 (0.1)	UNDER	0
GO:0045333	cellular respiration	27	0.083 (0.040)	0.134 (0.002)	NO SIG	0.104	0.0 (0.2)	5.1 (0.1)	UNDER	0
GO:0019226	transmission of nerve impulse	136	0.101 (0.026)	0.134 (0.002)	NO SIG	0.101	5.5 (0.9)	5.1 (0.1)	NO SIG	0.356
GO:0006220	pyrimidine nucleotide metabolism	65	0.101 (0.026)	0.134 (0.002)	NO SIG	0.099	3.4 (1.1)	5.1 (0.1)	NO SIG	0.057
GO:0009116	nucleoside metabolism	29	0.063 (0.055)	0.134 (0.002)	NO SIG	0.098	0.1 (0.6)	5.1 (0.1)	UNDER	3.55E-15
GO:0007169	transmembrane receptor protein tyrosine kinase signa	120	0.104 (0.023)	0.134 (0.002)	NO SIG	0.096	2.2 (0.8)	5.1 (0.1)	UNDER	2.29E-04
GO:0006695	cholesterol biosynthesis	32	0.082 (0.039)	0.134 (0.002)	NO SIG	0.094	4.1 (2.7)	5.1 (0.1)	NO SIG	0.351
GO:0006334	nucleosome assembly	74	0.095 (0.030)	0.134 (0.002)	NO SIG	0.092	0.6 (0.8)	5.1 (0.1)	UNDER	3.98E-09
GO:0006644	phospholipid metabolism	92	0.091 (0.032)	0.134 (0.002)	NO SIG	0.086	0.9 (0.7)	5.1 (0.1)	UNDER	7.03E-09
GO:0006302	double-strand break repair	54	0.074 (0.043)	0.134 (0.002)	NO SIG	0.079	0.2 (0.6)	5.1 (0.1)	UNDER	6.46E-16
GO:0009069	serine family amino acid metabolism	44	0.060 (0.052)	0.134 (0.002)	NO SIG	0.076	0.3 (0.7)	5.1 (0.1)	UNDER	3.28E-11
GO:0006886	intracellular protein transport	500	0.115 (0.013)	0.134 (0.002)	NO SIG	0.074	3.5 (0.5)	5.2 (0.1)	UNDER	3.02E-04
GO:0009607	response to biotic stimulus	710	0.116 (0.012)	0.135 (0.002)	NO SIG	0.073	4.7 (0.4)	5.1 (0.1)	NO SIG	0.153
GO:0007249	NIK-I-kappaB/NF-kappaB cascade	38	0.066 (0.046)	0.134 (0.002)	NO SIG	0.071	0.1 (0.5)	5.1 (0.1)	UNDER	0
GO:0006865	amino acid transport	30	0.057 (0.051)	0.134 (0.002)	NO SIG	0.064	0.3 (1.0)	5.1 (0.1)	UNDER	6.42E-07
GO:0007259	JAK-STAT cascade	46	0.054 (0.053)	0.134 (0.002)	NO SIG	0.064	2.5 (0.8)	5.1 (0.1)	UNDER	3.65E-04
GO:0009628	response to abiotic stimulus	418	0.113 (0.014)	0.134 (0.002)	NO SIG	0.063	3.6 (0.3)	5.2 (0.1)	UNDER	4.07E-06
GO:0015718	monocarboxylic acid transport	32	0.045 (0.057)	0.134 (0.002)	NO SIG	0.060	1.1 (1.7)	5.1 (0.1)	UNDER	0.007
GO:0007242	intracellular signaling cascade	751	0.116 (0.012)	0.135 (0.002)	NO SIG	0.060	3.2 (0.4)	5.2 (0.1)	UNDER	8.00E-07
GO:0016125	sterol metabolism	108	0.098 (0.023)	0.134 (0.002)	NO SIG	0.060	5.2 (0.9)	5.1 (0.1)	NO SIG	0.453
GO:0009581	perception of external stimulus	426	0.112 (0.014)	0.134 (0.002)	NO SIG	0.055	3.5 (0.3)	5.2 (0.1)	UNDER	1.16E-06
GO:0006338	chromatin modeling	115	0.086 (0.030)	0.134 (0.002)	NO SIG	0.054	1.2 (0.8)	5.2 (0.1)	UNDER	1.34E-06
GO:0016568	chromatin modification	115	0.086 (0.030)	0.134 (0.002)	NO SIG	0.054	1.2 (0.8)	5.2 (0.1)	UNDER	1.34E-06
GO:0016582	non-covalent chromatin modification	115	0.086 (0.030)	0.134 (0.002)	NO SIG	0.054	1.2 (0.8)	5.2 (0.1)	UNDER	1.34E-06
GO:0019538	protein metabolism	216	0.104 (0.019)	0.134 (0.002)	NO SIG	0.052	2.4 (0.4)	5.2 (0.1)	UNDER	5.70E-11
GO:0006413	translational initiation	107	0.093 (0.025)	0.134 (0.002)	NO SIG	0.049	0.3 (0.5)	5.2 (0.1)	UNDER	0
GO:0006749	glutathione metabolism	46	0.079 (0.033)	0.134 (0.002)	NO SIG	0.049	0.2 (0.7)	5.1 (0.1)	UNDER	1.90E-13
GO:0000004	biological_process unknown	39	0.059 (0.045)	0.134 (0.002)	NO SIG	0.048	0.6 (1.2)	5.1 (0.1)	UNDER	8.31E-05
GO:0006310	DNA recombination	90	0.070 (0.039)	0.134 (0.002)	NO SIG	0.048	1.8 (1.1)	5.1 (0.1)	UNDER	9.72E-04
GO:0006376	mRNA splice site selection	26	0.074 (0.035)	0.134 (0.002)	NO SIG	0.046	0.0 (0.3)	5.1 (0.1)	UNDER	0
GO:0006446	regulation of translational initiation	78	0.087 (0.027)	0.134 (0.002)	NO SIG	0.043	0.3 (0.6)	5.1 (0.1)	UNDER	1.37E-15

GO:0030029	actin filament-based process	36	0.079 (0.032)	0.134 (0.002)	NO SIG	0.041	0.0 (0.2)	5.1 (0.1)	UNDER	0
GO:0006497	protein lipidation	100	0.084 (0.029)	0.134 (0.002)	NO SIG	0.041	0.8 (0.8)	5.1 (0.1)	UNDER	8.57E-09
GO:0042157	lipoprotein metabolism	100	0.084 (0.029)	0.134 (0.002)	NO SIG	0.041	0.8 (0.8)	5.1 (0.1)	UNDER	8.57E-09
GO:0042158	lipoprotein biosynthesis	100	0.084 (0.029)	0.134 (0.002)	NO SIG	0.041	0.8 (0.8)	5.1 (0.1)	UNDER	8.57E-09
GO:0008610	lipid biosynthesis	112	0.087 (0.027)	0.134 (0.002)	NO SIG	0.041	0.8 (0.7)	5.2 (0.1)	UNDER	4.12E-10
GO:0006499	N-terminal protein myristoylation	72	0.073 (0.035)	0.134 (0.002)	NO SIG	0.040	0.5 (0.8)	5.1 (0.1)	UNDER	1.66E-09
GO:0018319	protein amino acid myristoylation	72	0.073 (0.035)	0.134 (0.002)	NO SIG	0.040	0.5 (0.8)	5.1 (0.1)	UNDER	1.66E-09
GO:0018377	protein myristoylation	72	0.073 (0.035)	0.134 (0.002)	NO SIG	0.040	0.5 (0.8)	5.1 (0.1)	UNDER	1.66E-09
GO:0006956	complement activation	33	0.042 (0.052)	0.134 (0.002)	NO SIG	0.038	0.2 (0.8)	5.1 (0.1)	UNDER	1.92E-10
GO:0016064	humoral defense mechanism (sensu Vertebrata)	33	0.042 (0.052)	0.134 (0.002)	NO SIG	0.038	0.2 (0.8)	5.1 (0.1)	UNDER	1.92E-10
GO:0006643	membrane lipid metabolism	112	0.082 (0.029)	0.134 (0.002)	NO SIG	0.038	1.2 (0.8)	5.2 (0.1)	UNDER	1.91E-07
GO:0007154	cell communication	3252	0.125 (0.006)	0.136 (0.002)	NO SIG	0.038	4.7 (0.2)	5.2 (0.1)	NO SIG	0.010
GO:0006099	tricarboxylic acid cycle	34	0.067 (0.038)	0.134 (0.002)	NO SIG	0.037	0.0 (0.3)	5.1 (0.1)	UNDER	0
GO:0015711	organic anion transport	40	0.044 (0.050)	0.134 (0.002)	NO SIG	0.036	0.9 (1.3)	5.1 (0.1)	UNDER	7.76E-04
GO:0006445	regulation of translation	124	0.094 (0.022)	0.134 (0.002)	NO SIG	0.035	0.3 (0.5)	5.2 (0.1)	UNDER	0
GO:0007114	budding	27	0.056 (0.043)	0.134 (0.002)	NO SIG	0.034	0.9 (1.7)	5.1 (0.1)	UNDER	0.006
GO:0009410	xenobiotic response	55	0.075 (0.032)	0.134 (0.002)	NO SIG	0.033	0.0 (0.3)	5.1 (0.1)	UNDER	0
GO:0009165	nucleotide biosynthesis	35	0.040 (0.051)	0.134 (0.002)	NO SIG	0.032	0.0 (0.2)	5.1 (0.1)	UNDER	0
GO:0007160	cell-matrix adhesion	85	0.064 (0.038)	0.134 (0.002)	NO SIG	0.032	0.8 (0.9)	5.1 (0.1)	UNDER	6.56E-07
GO:0007229	integrin-mediated signaling pathway	36	0.040 (0.048)	0.134 (0.002)	NO SIG	0.026	0.0 (0.2)	5.1 (0.1)	UNDER	0
GO:0007264	small GTPase mediated signal transduction	242	0.094 (0.021)	0.134 (0.002)	NO SIG	0.026	2.7 (0.7)	5.2 (0.1)	UNDER	2.50E-04
GO:0006820	anion transport	80	0.061 (0.037)	0.134 (0.002)	NO SIG	0.023	0.9 (1.0)	5.1 (0.1)	UNDER	9.18E-06
GO:0008152	metabolism	7800	0.128 (0.003)	0.138 (0.003)	NO SIG	0.021	4.8 (0.1)	5.3 (0.1)	UNDER	5.43E-04
GO:0007005	mitochondrion organization and biogenesis	77	0.079 (0.026)	0.134 (0.002)	NO SIG	0.019	0.2 (0.4)	5.1 (0.1)	UNDER	0
GO:0006807	nitrogen metabolism	46	0.057 (0.037)	0.134 (0.002)	NO SIG	0.017	1.1 (1.1)	5.1 (0.1)	UNDER	1.07E-04
GO:0006944	membrane fusion	51	0.025 (0.050)	0.134 (0.002)	NO SIG	0.015	0.7 (1.0)	5.1 (0.1)	UNDER	2.18E-06
GO:0006979	oxidative stress response	38	0.047 (0.039)	0.134 (0.002)	NO SIG	0.013	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0009147	pyrimidine nucleoside triphosphate metabolism	26	0.058 (0.034)	0.134 (0.002)	NO SIG	0.012	0.0 (0.2)	5.1 (0.1)	UNDER	0
GO:0009148	pyrimidine nucleoside triphosphate biosynthesis	26	0.058 (0.034)	0.134 (0.002)	NO SIG	0.012	0.0 (0.2)	5.1 (0.1)	UNDER	0
GO:0007148	regulation of cell shape and cell size	224	0.090 (0.020)	0.134 (0.002)	NO SIG	0.012	1.9 (0.4)	5.2 (0.1)	UNDER	5.80E-15
GO:0006955	immune response	464	0.101 (0.014)	0.135 (0.002)	NO SIG	0.010	3.8 (0.5)	5.2 (0.1)	UNDER	0.006
GO:0000072	M-phase specific microtubule process	48	0.070 (0.027)	0.134 (0.002)	SLOWER	0.010	2.8 (1.8)	5.1 (0.1)	NO SIG	0.091
GO:0007051	spindle assembly	48	0.070 (0.027)	0.134 (0.002)	SLOWER	0.010	2.8 (1.8)	5.1 (0.1)	NO SIG	0.091
GO:0007052	mitotic spindle assembly	48	0.070 (0.027)	0.134 (0.002)	SLOWER	0.010	2.8 (1.8)	5.1 (0.1)	NO SIG	0.091
GO:0006694	steroid biosynthesis	70	0.070 (0.027)	0.134 (0.002)	SLOWER	0.009	1.9 (1.2)	5.1 (0.1)	UNDER	0.004
GO:0006952	defense response	500	0.102 (0.014)	0.135 (0.002)	SLOWER	0.009	3.8 (0.5)	5.2 (0.1)	UNDER	0.004
GO:0006487	N-linked glycosylation	62	0.062 (0.030)	0.134 (0.002)	SLOWER	0.009	0.0 (0.2)	5.1 (0.1)	UNDER	0
GO:0006457	protein folding	116	0.085 (0.021)	0.134 (0.002)	SLOWER	0.008	0.3 (0.5)	5.2 (0.1)	UNDER	0
GO:0006066	alcohol metabolism	30	0.024 (0.046)	0.134 (0.002)	SLOWER	0.008	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0009605	response to external stimulus	1136	0.111 (0.009)	0.135 (0.002)	SLOWER	0.006	3.9 (0.3)	5.2 (0.1)	UNDER	1.28E-05
GO:0015031	protein transport	689	0.106 (0.011)	0.135 (0.002)	SLOWER	0.005	3.6 (0.4)	5.2 (0.1)	UNDER	4.24E-05
GO:0006119	oxidative phosphorylation	36	0.055 (0.029)	0.134 (0.002)	SLOWER	0.003	0.0 (0.1)	5.1 (0.1)	UNDER	0

GO:0015674	di-\, tri-valent inorganic cation transport	68	0.043 (0.033)	0.134 (0.002)	SLOWER	0.003	0.5 (0.8)	5.1 (0.1)	UNDER	1.39E-09
GO:0006635	fatty acid beta-oxidation	44	0.045 (0.032)	0.134 (0.002)	SLOWER	0.002	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0019395	fatty acid oxidation	44	0.045 (0.032)	0.134 (0.002)	SLOWER	0.002	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0006888	ER to Golgi transport	42	0.028 (0.038)	0.134 (0.002)	SLOWER	0.002	0.4 (0.9)	5.1 (0.1)	UNDER	3.42E-07
GO:0006183	GTP biosynthesis	34	0.031 (0.035)	0.134 (0.002)	SLOWER	0.002	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0009144	purine nucleoside triphosphate metabolism	34	0.031 (0.035)	0.134 (0.002)	SLOWER	0.002	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0009145	purine nucleoside triphosphate biosynthesis	34	0.031 (0.035)	0.134 (0.002)	SLOWER	0.002	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0006164	purine nucleotide biosynthesis	60	0.043 (0.031)	0.134 (0.002)	SLOWER	0.002	0.8 (0.8)	5.1 (0.1)	UNDER	2.13E-07
GO:0006818	hydrogen transport	83	0.059 (0.025)	0.134 (0.002)	SLOWER	0.001	1.2 (0.9)	5.1 (0.1)	UNDER	2.87E-06
GO:0007155	cell adhesion	386	0.088 (0.015)	0.135 (0.002)	SLOWER	0.001	2.6 (0.5)	5.2 (0.1)	UNDER	4.09E-08
GO:0008202	steroid metabolism	177	0.072 (0.020)	0.135 (0.002)	SLOWER	0.001	3.3 (0.6)	5.1 (0.1)	UNDER	0.001
GO:0006163	purine nucleotide metabolism	82	0.049 (0.028)	0.134 (0.002)	SLOWER	0.001	0.7 (0.7)	5.1 (0.1)	UNDER	3.84E-10
GO:0016192	vesicle-mediated transport	442	0.092 (0.014)	0.135 (0.002)	SLOWER	9.83E-04	3.6 (0.5)	5.2 (0.1)	UNDER	4.95E-04
GO:0009108	coenzyme biosynthesis	52	0.038 (0.030)	0.134 (0.002)	SLOWER	7.66E-04	0.2 (0.6)	5.1 (0.1)	UNDER	8.92E-17
GO:0006887	exocytosis	94	0.045 (0.028)	0.134 (0.002)	SLOWER	7.46E-04	1.0 (0.7)	5.1 (0.1)	UNDER	3.08E-09
GO:0009117	nucleotide metabolism	178	0.070 (0.020)	0.135 (0.002)	SLOWER	7.25E-04	1.7 (0.6)	5.2 (0.1)	UNDER	1.51E-09
GO:0009126	purine nucleoside monophosphate metabolism	56	0.038 (0.030)	0.134 (0.002)	SLOWER	7.16E-04	0.2 (0.5)	5.1 (0.1)	UNDER	0
GO:0015931	nucleobase\, nucleoside\, nucleotide and nucleic acid	28	0.043 (0.028)	0.134 (0.002)	SLOWER	6.90E-04	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0006096	glycolysis	94	0.059 (0.023)	0.134 (0.002)	SLOWER	6.65E-04	0.2 (0.5)	5.2 (0.1)	UNDER	0
GO:0009072	aromatic amino acid family metabolism	30	0.015 (0.037)	0.134 (0.002)	SLOWER	6.25E-04	0.0 (0.0)	5.1 (0.1)	UNDER	0
GO:0009074	aromatic amino acid family catabolism	30	0.015 (0.037)	0.134 (0.002)	SLOWER	6.25E-04	0.0 (0.0)	5.1 (0.1)	UNDER	0
GO:0006800	oxygen and radical metabolism	71	0.051 (0.025)	0.134 (0.002)	SLOWER	4.63E-04	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0005975	**carbohydrate metabolism**	**464**	**0.092 (0.012)**	**0.135 (0.002)**	**SLOWER**	**3.18E-04**	**1.1 (0.3)**	**5.2 (0.1)**	**UNDER**	**0**
GO:0030001	metal ion transport	111	0.046 (0.026)	0.134 (0.002)	SLOWER	3.04E-04	0.3 (0.5)	5.2 (0.1)	UNDER	0
GO:0006928	cell motility	381	0.082 (0.015)	0.135 (0.002)	SLOWER	2.69E-04	2.1 (0.4)	5.2 (0.1)	UNDER	1.38E-13
GO:0008652	amino acid biosynthesis	53	0.011 (0.035)	0.134 (0.002)	SLOWER	2.46E-04	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0006731	coenzymes and prosthetic group metabolism	128	0.067 (0.019)	0.134 (0.002)	SLOWER	2.08E-04	0.3 (0.4)	5.2 (0.1)	UNDER	0
GO:0019318	hexose metabolism	162	0.067 (0.019)	0.135 (0.002)	SLOWER	1.69E-04	0.2 (0.3)	5.2 (0.1)	UNDER	0
GO:0005996	monosaccharide metabolism	163	0.067 (0.019)	0.135 (0.002)	SLOWER	1.64E-04	0.2 (0.3)	5.2 (0.1)	UNDER	0
GO:0006092	main pathways of carbohydrate metabolism	166	0.069 (0.018)	0.135 (0.002)	SLOWER	1.61E-04	0.4 (0.4)	5.2 (0.1)	UNDER	0
GO:0006970	osmotic response	41	0.010 (0.034)	0.134 (0.002)	SLOWER	1.54E-04	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0006411	**protein metabolism and modification**	**3244**	**0.117 (0.005)**	**0.137 (0.003)**	**SLOWER**	**1.46E-04**	**3.3 (0.2)**	**5.5 (0.1)**	**UNDER**	**0**
GO:0008150	biological_process	12518	0.128 (0.003)	0.145 (0.004)	SLOWER	1.44E-04	4.8 (0.1)	5.8 (0.2)	UNDER	1.72E-08
GO:0015672	monovalent inorganic cation transport	122	0.058 (0.021)	0.134 (0.002)	SLOWER	1.34E-04	0.1 (0.2)	5.2 (0.1)	UNDER	0
GO:0006937	regulation of muscle contraction	76	0.035 (0.027)	0.134 (0.002)	SLOWER	1.27E-04	0.1 (0.3)	5.1 (0.1)	UNDER	0
GO:0006007	glucose catabolism	106	0.055 (0.021)	0.134 (0.002)	SLOWER	1.20E-04	0.2 (0.4)	5.2 (0.1)	UNDER	0
GO:0019320	hexose catabolism	106	0.055 (0.021)	0.134 (0.002)	SLOWER	1.20E-04	0.2 (0.4)	5.2 (0.1)	UNDER	0
GO:0009056	**catabolism**	**982**	**0.103 (0.008)**	**0.136 (0.002)**	**SLOWER**	**6.72E-05**	**2.7 (0.3)**	**5.3 (0.1)**	**UNDER**	**1.26E-14**
GO:0006095	catabolic carbohydrate metabolism	161	0.064 (0.018)	0.135 (0.002)	SLOWER	6.55E-05	0.2 (0.3)	5.2 (0.1)	UNDER	0
GO:0015985	energy coupled proton transport\, down the electroche	70	0.040 (0.024)	0.134 (0.002)	SLOWER	5.28E-05	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0015986	ATP synthesis coupled proton transport	70	0.040 (0.024)	0.134 (0.002)	SLOWER	5.28E-05	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0015992	proton transport	71	0.040 (0.024)	0.134 (0.002)	SLOWER	4.06E-05	0.0 (0.1)	5.1 (0.1)	UNDER	0

GO:0015980	energy derivation by oxidation of organic compounds	284	0.077 (0.014)	0.135 (0.002)	SLOWER	3.36E-05	0.3 (0.3)	5.2 (0.1)	UNDER	0
GO:0006006	glucose metabolism	130	0.055 (0.020)	0.134 (0.002)	SLOWER	3.33E-05	0.2 (0.3)	5.2 (0.1)	UNDER	0
GO:0009308	amine metabolism	311	0.070 (0.016)	0.135 (0.002)	SLOWER	2.77E-05	4.3 (0.5)	5.1 (0.1)	NO SIG	0.038
GO:0006812	cation transport	213	0.057 (0.019)	0.135 (0.002)	SLOWER	2.66E-05	0.3 (0.4)	5.2 (0.1)	UNDER	0
GO:0009064	glutamine family amino acid metabolism	67	0.006 (0.032)	0.134 (0.002)	SLOWER	2.45E-05	1.5 (0.9)	5.1 (0.1)	UNDER	3.65E-05
GO:0008151	cell growth and/or maintenance	10521	0.125 (0.003)	0.144 (0.003)	SLOWER	2.44E-05	4.5 (0.1)	5.8 (0.1)	UNDER	9.15E-14
GO:0009063	amino acid catabolism	97	0.024 (0.027)	0.135 (0.002)	SLOWER	2.20E-05	0.7 (0.7)	5.1 (0.1)	UNDER	1.10E-10
GO:0009310	amine catabolism	97	0.024 (0.027)	0.135 (0.002)	SLOWER	2.20E-05	0.7 (0.7)	5.1 (0.1)	UNDER	1.10E-10
GO:0009065	glutamine family amino acid catabolism	29	-0.064 (0.049)	0.134 (0.002)	SLOWER	2.16E-05	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0006732	coenzyme metabolism	99	0.050 (0.020)	0.134 (0.002)	SLOWER	1.42E-05	0.1 (0.3)	5.2 (0.1)	UNDER	0
GO:0006811	ion transport	313	0.064 (0.017)	0.135 (0.002)	SLOWER	1.23E-05	0.9 (0.4)	5.2 (0.1)	UNDER	0
GO:0006752	group transfer coenzyme metabolism	37	0.011 (0.029)	0.134 (0.002)	SLOWER	1.21E-05	0.0 (0.2)	5.1 (0.1)	UNDER	0
GO:0000001	mitochondrion inheritance	32	0.015 (0.028)	0.134 (0.002)	SLOWER	1.20E-05	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0000011	vacuole inheritance	32	0.015 (0.028)	0.134 (0.002)	SLOWER	1.20E-05	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0000132	mitotic spindle orientation	32	0.015 (0.028)	0.134 (0.002)	SLOWER	1.20E-05	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0040001	mitotic spindle positioning and orientation	32	0.015 (0.028)	0.134 (0.002)	SLOWER	1.20E-05	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0006414	translational elongation	71	0.039 (0.022)	0.134 (0.002)	SLOWER	8.38E-06	0.4 (0.7)	5.1 (0.1)	UNDER	3.83E-13
GO:0006519	amino acid and derivative metabolism	318	0.067 (0.016)	0.135 (0.002)	SLOWER	8.28E-06	4.2 (0.4)	5.1 (0.1)	NO SIG	0.014
GO:0045045	secretory pathway	179	0.050 (0.020)	0.135 (0.002)	SLOWER	8.21E-06	1.2 (0.6)	5.2 (0.1)	UNDER	1.10E-11
GO:0000041	transition metal transport	43	-0.009 (0.033)	0.134 (0.002)	SLOWER	7.86E-06	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0006823	heavy metal ion transport	43	-0.009 (0.033)	0.134 (0.002)	SLOWER	7.86E-06	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0007013	actin modification	36	0.019 (0.026)	0.134 (0.002)	SLOWER	5.72E-06	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0009306	protein secretion	183	0.048 (0.019)	0.135 (0.002)	SLOWER	3.69E-06	1.2 (0.6)	5.2 (0.1)	UNDER	2.86E-12
GO:0006936	muscle contraction	151	0.044 (0.020)	0.135 (0.002)	SLOWER	3.11E-06	0.1 (0.2)	5.2 (0.1)	UNDER	0
GO:0007033	vacuole organization and biogenesis	36	0.017 (0.026)	0.134 (0.002)	SLOWER	2.85E-06	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0006631	fatty acid metabolism	120	0.032 (0.021)	0.135 (0.002)	SLOWER	5.51E-07	0.0 (0.2)	5.2 (0.1)	UNDER	0
GO:0006810	**transport**	**1459**	**0.098 (0.007)**	**0.137 (0.002)**	**SLOWER**	**3.65E-07**	**2.5 (0.2)**	**5.3 (0.1)**	**UNDER**	**0**
GO:0006520	amino acid metabolism	229	0.040 (0.019)	0.135 (0.002)	SLOWER	2.90E-07	0.9 (0.5)	5.2 (0.1)	UNDER	5.00E-21
GO:0006826	iron transport	28	-0.050 (0.037)	0.134 (0.002)	SLOWER	2.47E-07	0.0 (0.1)	5.1 (0.1)	UNDER	0
GO:0016052	carbohydrate catabolism	141	0.042 (0.018)	0.135 (0.002)	SLOWER	1.61E-07	0.1 (0.3)	5.2 (0.1)	UNDER	0
GO:0006091	energy pathways	392	0.070 (0.012)	0.135 (0.002)	SLOWER	1.04E-07	0.3 (0.2)	5.2 (0.1)	UNDER	0
GO:0006118	electron transport	223	0.054 (0.015)	0.135 (0.002)	SLOWER	8.95E-08	0.2 (0.3)	5.2 (0.1)	UNDER	0
GO:0006629	lipid metabolism	656	0.077 (0.011)	0.136 (0.002)	SLOWER	3.22E-08	1.6 (0.3)	5.3 (0.1)	UNDER	0
GO:0009059	macromolecule biosynthesis	811	0.079 (0.008)	0.136 (0.002)	SLOWER	5.72E-13	0.8 (0.2)	5.3 (0.1)	UNDER	0
GO:0006412	protein biosynthesis	704	0.073 (0.008)	0.136 (0.002)	SLOWER	3.21E-14	0.3 (0.2)	5.3 (0.1)	UNDER	0
GO:0009058	**biosynthesis**	**1302**	**0.085 (0.006)**	**0.137 (0.002)**	**SLOWER**	**1.15E-14**	**1.9 (0.2)**	**5.4 (0.1)**	**UNDER**	**0**
MANUAL	**Swissprot transcription**		**0.254 (0.022)**	**0.129 (0.004)**	**FASTER**	**0**	**16.6 (0.5)**	**4.6 (0.1)**	**OVER**	**0**

Supplementary Table 2: Association of Gene Ontology Categories with Decay Rate for Bud8 Experiments

These tables show details for all Gene Ontology categories. Inferences for manual classification scheme shown at the bottom.

Those categories used in Figure 1B are highlighted in bold. Standard Deviations are shown in parentheses

P-values less than 1E-20 are indicated with a "0".

			Rate *r* (hour^-1)				Half-life < 2hr. Decay %			
GO Number	Category Name	Probe Sets	Group	Non-Group	Distribution	p-value	In-Group	Non-Group	Representation	p-value
GO:0006350	**transcription**	**446**	**0.234 (0.015)**	**0.134 (0.004)**	**FASTER**	**7.45E-11**	**12.2 (0.7)**	**3.7 (0.1)**	**OVER**	**0**
GO:0006351	transcription, DNA-dependent	440	0.234 (0.015)	0.134 (0.004)	FASTER	1.17E-10	12.1 (0.7)	3.7 (0.1)	OVER	0
GO:0006355	regulation of transcription, DNA-dependent	350	0.241 (0.018)	0.135 (0.004)	FASTER	3.26E-09	13.1 (0.8)	3.8 (0.1)	OVER	0
GO:0006139	nucleobase, nucleoside, nucleotide and nucleic a	835	0.185 (0.010)	0.135 (0.004)	FASTER	2.21E-06	7.8 (0.4)	3.8 (0.1)	OVER	2.08E-18
GO:0006366	transcription from Pol II promoter	241	0.227 (0.019)	0.138 (0.004)	FASTER	2.97E-06	11.1 (1.0)	4.1 (0.1)	OVER	1.93E-13
GO:0006357	regulation of transcription from Pol II promoter	138	0.234 (0.025)	0.140 (0.004)	FASTER	7.84E-05	12.2 (1.3)	4.2 (0.1)	OVER	9.87E-11
GO:0008283	cell proliferation	204	0.183 (0.019)	0.141 (0.004)	NO SIG	0.016	8.7 (1.0)	4.2 (0.1)	OVER	1.92E-06
GO:0000079	regulation of CDK activity	34	0.250 (0.051)	0.142 (0.004)	NO SIG	0.018	8.1 (3.3)	4.4 (0.1)	NO SIG	0.130
GO:0000122	negative regulation of transcription from Pol II pro	28	0.261 (0.060)	0.142 (0.004)	NO SIG	0.023	16.2 (3.1)	4.3 (0.1)	OVER	8.57E-05
GO:0016481	negative regulation of transcription	28	0.261 (0.060)	0.142 (0.004)	NO SIG	0.023	16.2 (3.1)	4.3 (0.1)	OVER	8.57E-05
GO:0000086	G2/M transition of mitotic cell cycle	38	0.230 (0.047)	0.142 (0.004)	NO SIG	0.029	7.2 (2.9)	4.4 (0.1)	NO SIG	0.167
GO:0007345	embryogenesis and morphogenesis	45	0.210 (0.036)	0.142 (0.004)	NO SIG	0.031	9.3 (1.9)	4.4 (0.1)	OVER	0.004
GO:0000082	G1/S transition of mitotic cell cycle	50	0.209 (0.042)	0.142 (0.004)	NO SIG	0.056	6.8 (2.5)	4.4 (0.1)	NO SIG	0.174
GO:0000074	**regulation of cell cycle**	**173**	**0.175 (0.021)**	**0.141 (0.004)**	**NO SIG**	**0.057**	**8.0 (1.1)**	**4.3 (0.1)**	**OVER**	**5.31E-04**
GO:0008285	negative regulation of cell proliferation	58	0.207 (0.041)	0.141 (0.004)	NO SIG	0.058	10.9 (2.4)	4.3 (0.1)	OVER	0.003
GO:0009611	response to wounding	43	0.203 (0.043)	0.142 (0.004)	NO SIG	0.076	10.9 (2.1)	4.3 (0.1)	OVER	8.43E-04
GO:0006917	induction of apoptosis	49	0.201 (0.042)	0.142 (0.004)	NO SIG	0.081	5.7 (1.9)	4.4 (0.1)	NO SIG	0.252
GO:0012502	induction of programmed cell death	49	0.201 (0.042)	0.142 (0.004)	NO SIG	0.081	5.7 (1.9)	4.4 (0.1)	NO SIG	0.252
GO:0042127	regulation of cell proliferation	107	0.182 (0.029)	0.141 (0.004)	NO SIG	0.083	8.2 (1.6)	4.3 (0.1)	OVER	0.007
GO:0006954	inflammatory response	26	0.219 (0.062)	0.142 (0.004)	NO SIG	0.107	13.6 (3.2)	4.4 (0.1)	OVER	0.002
GO:0045087	innate immune response	26	0.219 (0.062)	0.142 (0.004)	NO SIG	0.107	13.6 (3.2)	4.4 (0.1)	OVER	0.002
GO:0000278	mitotic cell cycle	202	0.166 (0.019)	0.141 (0.004)	NO SIG	0.108	4.0 (0.9)	4.4 (0.1)	NO SIG	0.304
GO:0007397	histogenesis and organogenesis	39	0.205 (0.052)	0.142 (0.004)	NO SIG	0.113	9.9 (1.9)	4.4 (0.1)	OVER	0.002
GO:0008371	obsolete	138	0.171 (0.024)	0.141 (0.004)	NO SIG	0.118	7.5 (1.0)	4.3 (0.1)	OVER	0.001
GO:0007049	cell cycle	340	0.159 (0.015)	0.141 (0.004)	NO SIG	0.126	5.4 (0.7)	4.3 (0.1)	NO SIG	0.057
GO:0006915	**apoptosis**	**119**	**0.166 (0.023)**	**0.142 (0.004)**	**NO SIG**	**0.147**	**5.0 (0.8)**	**4.4 (0.1)**	**NO SIG**	**0.247**
GO:0012501	programmed cell death	119	0.166 (0.023)	0.142 (0.004)	NO SIG	0.147	5.0 (0.8)	4.4 (0.1)	NO SIG	0.247
GO:0007399	neurogenesis	89	0.172 (0.031)	0.142 (0.004)	NO SIG	0.170	8.7 (1.2)	4.3 (0.1)	OVER	2.05E-04
GO:0016265	death	129	0.162 (0.022)	0.142 (0.004)	NO SIG	0.179	4.6 (0.8)	4.4 (0.1)	NO SIG	0.405
GO:0000084	S phase of mitotic cell cycle	80	0.172 (0.033)	0.142 (0.004)	NO SIG	0.183	4.5 (1.5)	4.4 (0.1)	NO SIG	0.480
GO:0008219	cell death	127	0.161 (0.022)	0.142 (0.004)	NO SIG	0.188	4.7 (0.8)	4.4 (0.1)	NO SIG	0.371
GO:0008202	steroid metabolism	28	0.191 (0.060)	0.142 (0.004)	NO SIG	0.209	8.8 (2.1)	4.4 (0.1)	NO SIG	0.020
GO:0000067	DNA replication and chromosome cycle	90	0.167 (0.034)	0.142 (0.004)	NO SIG	0.232	5.5 (1.5)	4.4 (0.1)	NO SIG	0.235
GO:0006950	stress response	194	0.156 (0.020)	0.142 (0.004)	NO SIG	0.238	6.9 (1.0)	4.3 (0.1)	OVER	0.003

GO:0006260	DNA replication	73	0.167 (0.036)	0.142 (0.004)	NO SIG	0.240	4.9 (1.7)	4.4 (0.1)	NO SIG	0.381
GO:0006468	protein amino acid phosphorylation	205	0.156 (0.020)	0.142 (0.004)	NO SIG	0.241	5.5 (0.7)	4.4 (0.1)	NO SIG	0.064
GO:0006261	DNA dependent DNA replication	45	0.173 (0.046)	0.142 (0.004)	NO SIG	0.250	2.6 (2.1)	4.4 (0.1)	NO SIG	0.201
GO:0007275	**development**	**348**	**0.152 (0.015)**	**0.142 (0.004)**	**NO SIG**	**0.252**	**6.5 (0.6)**	**4.3 (0.1)**	**OVER**	**4.71E-05**
GO:0009613	response to pest/pathogen/parasite	89	0.157 (0.029)	0.142 (0.004)	NO SIG	0.309	7.1 (1.5)	4.4 (0.1)	NO SIG	0.031
GO:0016310	phosphorylation	214	0.151 (0.019)	0.142 (0.004)	NO SIG	0.311	5.3 (0.7)	4.4 (0.1)	NO SIG	0.104
GO:0006913	nucleocytoplasmic transport	43	0.159 (0.036)	0.142 (0.004)	NO SIG	0.316	6.8 (0.8)	4.4 (0.1)	OVER	0.002
GO:0016482	cytoplasmic transport	43	0.159 (0.036)	0.142 (0.004)	NO SIG	0.316	6.8 (0.8)	4.4 (0.1)	OVER	0.002
GO:0009653	morphogenesis	215	0.151 (0.019)	0.142 (0.004)	NO SIG	0.323	6.9 (0.8)	4.3 (0.1)	OVER	2.74E-04
GO:0009887	organogenesis	214	0.151 (0.019)	0.142 (0.004)	NO SIG	0.327	7.0 (0.8)	4.3 (0.1)	OVER	2.47E-04
GO:0006959	humoral immune response	29	0.166 (0.053)	0.142 (0.004)	NO SIG	0.330	8.6 (3.1)	4.4 (0.1)	NO SIG	0.082
GO:0009607	response to biotic stimulus	195	0.149 (0.020)	0.142 (0.004)	NO SIG	0.366	6.1 (0.8)	4.3 (0.1)	NO SIG	0.020
GO:0006952	defense response	132	0.150 (0.023)	0.142 (0.004)	NO SIG	0.374	6.0 (1.0)	4.4 (0.1)	NO SIG	0.050
GO:0007186	G-protein coupled receptor protein signaling pathv	74	0.155 (0.041)	0.142 (0.004)	NO SIG	0.379	6.5 (1.5)	4.4 (0.1)	NO SIG	0.080
GO:0007243	protein kinase cascade	58	0.153 (0.038)	0.142 (0.004)	NO SIG	0.393	6.5 (1.3)	4.4 (0.1)	NO SIG	0.058
GO:0008284	positive regulation of cell proliferation	49	0.153 (0.041)	0.142 (0.004)	NO SIG	0.395	4.9 (1.9)	4.4 (0.1)	NO SIG	0.400
GO:0007267	cell-cell signaling	101	0.149 (0.028)	0.142 (0.004)	NO SIG	0.403	4.9 (0.8)	4.4 (0.1)	NO SIG	0.281
GO:0007276	gametogenesis	40	0.150 (0.037)	0.142 (0.004)	NO SIG	0.416	3.9 (1.5)	4.4 (0.1)	NO SIG	0.372
GO:0006796	phosphate metabolism	290	0.144 (0.017)	0.142 (0.004)	NO SIG	0.452	4.4 (0.6)	4.4 (0.1)	NO SIG	0.490
GO:0007167	enzyme linked receptor protein signaling pathway	50	0.146 (0.034)	0.142 (0.004)	NO SIG	0.456	3.9 (1.0)	4.4 (0.1)	NO SIG	0.298
GO:0007166	cell surface receptor linked signal transduction	175	0.144 (0.022)	0.142 (0.004)	NO SIG	0.462	4.9 (0.8)	4.4 (0.1)	NO SIG	0.245
GO:0006955	immune response	125	0.144 (0.023)	0.142 (0.004)	NO SIG	0.462	5.5 (1.0)	4.4 (0.1)	NO SIG	0.128
GO:0007283	spermatogenesis	26	0.144 (0.046)	0.142 (0.004)	NO SIG	0.482	5.7 (2.0)	4.4 (0.1)	NO SIG	0.261
GO:0000003	reproduction	47	0.144 (0.035)	0.142 (0.004)	NO SIG	0.484	3.4 (1.4)	4.4 (0.1)	NO SIG	0.241
GO:0001501	skeletal development	43	0.142 (0.037)	0.142 (0.004)	NO SIG	0.496	7.8 (2.2)	4.4 (0.1)	NO SIG	0.057
GO:0006281	DNA repair	92	0.141 (0.030)	0.142 (0.004)	NO SIG	0.488	4.1 (1.4)	4.4 (0.1)	NO SIG	0.427
GO:0000087	M phase of mitotic cell cycle	71	0.141 (0.030)	0.142 (0.004)	NO SIG	0.482	3.2 (1.3)	4.4 (0.1)	NO SIG	0.176
GO:0007067	mitosis	70	0.140 (0.030)	0.142 (0.004)	NO SIG	0.471	3.2 (1.3)	4.4 (0.1)	NO SIG	0.188
GO:0000280	nuclear division	84	0.139 (0.028)	0.142 (0.004)	NO SIG	0.459	2.8 (1.2)	4.4 (0.1)	NO SIG	0.093
GO:0007165	**signal transduction**	**570**	**0.141 (0.012)**	**0.142 (0.004)**	**NO SIG**	**0.459**	**3.6 (0.4)**	**4.5 (0.1)**	**NO SIG**	**0.016**
GO:0009100	glycoprotein metabolism	28	0.136 (0.052)	0.142 (0.004)	NO SIG	0.451	3.8 (1.1)	4.4 (0.1)	NO SIG	0.287
GO:0000279	M phase	89	0.139 (0.027)	0.142 (0.004)	NO SIG	0.449	2.7 (1.1)	4.4 (0.1)	NO SIG	0.061
GO:0009309	amine biosynthesis	28	0.136 (0.051)	0.142 (0.004)	NO SIG	0.449	10.3 (1.8)	4.4 (0.1)	OVER	5.96E-04
GO:0006897	endocytosis	34	0.135 (0.049)	0.142 (0.004)	NO SIG	0.444	5.2 (2.3)	4.4 (0.1)	NO SIG	0.367
GO:0006486	protein amino acid glycosylation	26	0.131 (0.055)	0.142 (0.004)	NO SIG	0.417	4.0 (1.2)	4.4 (0.1)	NO SIG	0.387
GO:0009101	glycoprotein biosynthesis	26	0.131 (0.055)	0.142 (0.004)	NO SIG	0.417	4.0 (1.2)	4.4 (0.1)	NO SIG	0.387
GO:0006997	nuclear organization and biogenesis	85	0.135 (0.030)	0.142 (0.004)	NO SIG	0.408	4.3 (0.8)	4.4 (0.1)	NO SIG	0.468
GO:0006259	**DNA metabolism**	**227**	**0.138 (0.018)**	**0.142 (0.004)**	**NO SIG**	**0.406**	**4.0 (0.8)**	**4.4 (0.1)**	**NO SIG**	**0.310**
GO:0006333	chromatin assembly/disassembly	27	0.128 (0.050)	0.142 (0.004)	NO SIG	0.386	7.3 (0.6)	4.4 (0.1)	OVER	2.92E-06
GO:0007001	chromosome organization and biogenesis (sensu	82	0.132 (0.031)	0.142 (0.004)	NO SIG	0.374	4.4 (0.8)	4.4 (0.1)	NO SIG	0.493
GO:0009888	histogenesis	39	0.127 (0.046)	0.142 (0.004)	NO SIG	0.370	1.3 (1.6)	4.4 (0.1)	NO SIG	0.030
GO:0006470	protein amino acid dephosphorylation	97	0.131 (0.028)	0.142 (0.004)	NO SIG	0.349	2.6 (0.9)	4.4 (0.1)	NO SIG	0.022

GO:0016311	dephosphorylation	97	0.131 (0.028)	0.142 (0.004)	NO SIG	0.349	2.6 (0.9)	4.4 (0.1)	NO SIG	0.022
GO:0007600	sensory perception	40	0.123 (0.047)	0.142 (0.004)	NO SIG	0.337	3.9 (1.7)	4.4 (0.1)	NO SIG	0.386
GO:0009582	perception of abiotic stimulus	40	0.124 (0.043)	0.142 (0.004)	NO SIG	0.333	3.8 (1.6)	4.4 (0.1)	NO SIG	0.358
GO:0006397	**mRNA processing**	**81**	**0.128 (0.030)**	**0.142 (0.004)**	**NO SIG**	**0.323**	**1.9 (0.8)**	**4.4 (0.1)**	**UNDER**	**0.001**
GO:0030001	metal ion transport	34	0.115 (0.060)	0.142 (0.004)	NO SIG	0.322	1.0 (1.5)	4.4 (0.1)	UNDER	0.009
GO:0019725	homeostasis	29	0.116 (0.052)	0.142 (0.004)	NO SIG	0.305	6.9 (2.6)	4.4 (0.1)	NO SIG	0.162
GO:0006873	ion homeostasis	27	0.113 (0.056)	0.142 (0.004)	NO SIG	0.299	7.4 (2.8)	4.4 (0.1)	NO SIG	0.135
GO:0030003	cation homeostasis	27	0.113 (0.056)	0.142 (0.004)	NO SIG	0.299	7.4 (2.8)	4.4 (0.1)	NO SIG	0.135
GO:0007169	transmembrane receptor protein tyrosine kinase s	29	0.117 (0.047)	0.142 (0.004)	NO SIG	0.297	0.3 (1.1)	4.4 (0.1)	UNDER	9.15E-05
GO:0006887	exocytosis	29	0.110 (0.059)	0.142 (0.004)	NO SIG	0.296	7.4 (1.2)	4.4 (0.1)	OVER	0.008
GO:0009308	amine metabolism	66	0.121 (0.039)	0.142 (0.004)	NO SIG	0.295	5.5 (1.3)	4.4 (0.1)	NO SIG	0.203
GO:0006519	amino acid and derivative metabolism	66	0.121 (0.039)	0.142 (0.004)	NO SIG	0.294	5.5 (1.3)	4.4 (0.1)	NO SIG	0.203
GO:0006464	**protein modification**	**395**	**0.135 (0.014)**	**0.143 (0.004)**	**NO SIG**	**0.293**	**3.6 (0.5)**	**4.5 (0.1)**	**NO SIG**	**0.030**
GO:0008152	metabolism	2036	0.139 (0.006)	0.144 (0.005)	NO SIG	0.288	4.6 (0.2)	4.3 (0.2)	NO SIG	0.108
GO:0009605	response to external stimulus	316	0.134 (0.015)	0.143 (0.004)	NO SIG	0.276	4.8 (0.6)	4.4 (0.1)	NO SIG	0.248
GO:0000226	microtubule cytoskeleton organization and biogen	32	0.116 (0.042)	0.142 (0.004)	NO SIG	0.263	3.1 (0.2)	4.4 (0.1)	UNDER	2.01E-07
GO:0006461	protein complex assembly	38	0.117 (0.039)	0.142 (0.004)	NO SIG	0.259	1.5 (1.5)	4.4 (0.1)	NO SIG	0.023
GO:0008380	RNA splicing	65	0.119 (0.035)	0.142 (0.004)	NO SIG	0.254	0.9 (1.0)	4.4 (0.1)	UNDER	3.04E-04
GO:0006371	mRNA splicing	59	0.118 (0.037)	0.142 (0.004)	NO SIG	0.252	0.9 (1.1)	4.4 (0.1)	UNDER	5.70E-04
GO:0019538	protein metabolism	54	0.119 (0.035)	0.142 (0.004)	NO SIG	0.249	2.2 (0.8)	4.4 (0.1)	UNDER	0.004
GO:0006396	RNA processing	131	0.126 (0.023)	0.143 (0.004)	NO SIG	0.238	1.8 (0.7)	4.5 (0.1)	UNDER	1.82E-04
GO:0007268	synaptic transmission	41	0.109 (0.047)	0.142 (0.004)	NO SIG	0.238	0.4 (0.9)	4.4 (0.1)	UNDER	4.62E-06
GO:0016070	RNA metabolism	144	0.126 (0.022)	0.143 (0.004)	NO SIG	0.228	1.7 (0.7)	4.5 (0.1)	UNDER	9.52E-05
GO:0008610	lipid biosynthesis	31	0.110 (0.043)	0.142 (0.004)	NO SIG	0.225	0.4 (1.1)	4.4 (0.1)	UNDER	7.30E-05
GO:0006512	ubiquitin cycle	50	0.123 (0.025)	0.142 (0.004)	NO SIG	0.220	0.0 (0.3)	4.4 (0.1)	UNDER	0
GO:0007264	small GTPase mediated signal transduction	75	0.118 (0.031)	0.143 (0.004)	NO SIG	0.220	3.2 (1.2)	4.4 (0.1)	NO SIG	0.150
GO:0007242	intracellular signaling cascade	232	0.128 (0.018)	0.143 (0.004)	NO SIG	0.209	3.2 (0.6)	4.5 (0.1)	NO SIG	0.024
GO:0006323	DNA packaging	70	0.116 (0.032)	0.143 (0.004)	NO SIG	0.209	3.1 (0.7)	4.4 (0.1)	NO SIG	0.029
GO:0006951	response to heat shock	37	0.111 (0.039)	0.142 (0.004)	NO SIG	0.209	4.2 (2.0)	4.4 (0.1)	NO SIG	0.468
GO:0019226	transmission of nerve impulse	44	0.105 (0.046)	0.142 (0.004)	NO SIG	0.206	0.3 (0.8)	4.4 (0.1)	UNDER	1.05E-06
GO:0006497	protein lipidation	27	0.104 (0.045)	0.142 (0.004)	NO SIG	0.198	0.4 (1.2)	4.4 (0.1)	UNDER	4.83E-04
GO:0042157	lipoprotein metabolism	27	0.104 (0.045)	0.142 (0.004)	NO SIG	0.198	0.4 (1.2)	4.4 (0.1)	UNDER	4.83E-04
GO:0042158	lipoprotein biosynthesis	27	0.104 (0.045)	0.142 (0.004)	NO SIG	0.198	0.4 (1.2)	4.4 (0.1)	UNDER	4.83E-04
GO:0009266	temperature response	43	0.112 (0.036)	0.142 (0.004)	NO SIG	0.197	3.7 (1.7)	4.4 (0.1)	NO SIG	0.331
GO:0006643	membrane lipid metabolism	28	0.105 (0.043)	0.142 (0.004)	NO SIG	0.195	2.4 (2.3)	4.4 (0.1)	NO SIG	0.191
GO:0006520	amino acid metabolism	48	0.099 (0.048)	0.143 (0.004)	NO SIG	0.184	1.4 (1.3)	4.4 (0.1)	NO SIG	0.014
GO:0006325	establishment and/or maintenance of chromatin a	62	0.112 (0.033)	0.143 (0.004)	NO SIG	0.180	3.2 (0.4)	4.4 (0.1)	UNDER	8.60E-04
GO:0006629	lipid metabolism	145	0.121 (0.022)	0.143 (0.004)	NO SIG	0.170	3.1 (0.7)	4.4 (0.1)	NO SIG	0.036
GO:0006899	non-selective vesicle transport	43	0.099 (0.044)	0.143 (0.004)	NO SIG	0.164	0.2 (0.7)	4.4 (0.1)	UNDER	3.16E-09
GO:0015672	monovalent inorganic cation transport	38	0.099 (0.044)	0.142 (0.004)	NO SIG	0.163	0.1 (0.6)	4.4 (0.1)	UNDER	3.68E-14
GO:0045045	secretory pathway	51	0.103 (0.039)	0.143 (0.004)	NO SIG	0.155	4.3 (0.8)	4.4 (0.1)	NO SIG	0.438
GO:0007398	ectoderm development	27	0.100 (0.040)	0.142 (0.004)	NO SIG	0.147	0.0 (0.3)	4.4 (0.1)	UNDER	0

GO:0009306	protein secretion	52	0.102 (0.038)	0.143 (0.004)	NO SIG	0.145	4.2 (0.8)	4.4 (0.1)	NO SIG	0.396
GO:0006605	protein targeting	70	0.112 (0.027)	0.143 (0.004)	NO SIG	0.132	2.8 (0.6)	4.4 (0.1)	UNDER	0.002
GO:0006631	fatty acid metabolism	27	0.087 (0.047)	0.142 (0.004)	NO SIG	0.121	0.0 (0.3)	4.4 (0.1)	UNDER	0
GO:0006812	cation transport	64	0.099 (0.037)	0.143 (0.004)	NO SIG	0.120	0.5 (0.8)	4.4 (0.1)	UNDER	7.93E-07
GO:0008544	epidermal differentiation	26	0.098 (0.037)	0.142 (0.004)	NO SIG	0.117	0.0 (0.1)	4.4 (0.1)	UNDER	0
GO:0006612	protein-membrane targeting	26	0.094 (0.040)	0.142 (0.004)	NO SIG	0.115	0.1 (0.6)	4.4 (0.1)	UNDER	1.97E-11
GO:0006928	cell motility	110	0.114 (0.023)	0.143 (0.004)	NO SIG	0.109	3.1 (1.1)	4.4 (0.1)	NO SIG	0.116
GO:0006413	translational initiation	26	0.090 (0.042)	0.142 (0.004)	NO SIG	0.109	0.0 (0.4)	4.4 (0.1)	UNDER	0
GO:0006811	ion transport	92	0.104 (0.030)	0.143 (0.004)	NO SIG	0.104	1.6 (1.0)	4.4 (0.1)	UNDER	0.002
GO:0009117	nucleotide metabolism	48	0.100 (0.033)	0.143 (0.004)	NO SIG	0.102	2.2 (0.4)	4.4 (0.1)	UNDER	2.67E-08
GO:0005975	**carbohydrate metabolism**	**106**	**0.111 (0.024)**	**0.143 (0.004)**	**NO SIG**	**0.095**	**1.9 (0.8)**	**4.4 (0.1)**	**UNDER**	**4.54E-04**
GO:0009581	perception of external stimulus	118	0.111 (0.024)	0.143 (0.004)	NO SIG	0.095	3.6 (1.0)	4.4 (0.1)	NO SIG	0.201
GO:0007517	muscle development	47	0.087 (0.042)	0.143 (0.004)	NO SIG	0.093	4.2 (0.9)	4.4 (0.1)	NO SIG	0.416
GO:0016192	vesicle-mediated transport	130	0.110 (0.025)	0.143 (0.004)	NO SIG	0.092	3.1 (0.7)	4.4 (0.1)	NO SIG	0.038
GO:0016043	cell organization and biogenesis	246	0.122 (0.015)	0.143 (0.004)	NO SIG	0.089	2.7 (0.4)	4.5 (0.1)	UNDER	6.35E-05
GO:0006511	ubiquitin-dependent protein degradation	85	0.116 (0.019)	0.143 (0.004)	NO SIG	0.088	1.4 (0.6)	4.4 (0.1)	UNDER	9.43E-08
GO:0009628	response to abiotic stimulus	115	0.111 (0.023)	0.143 (0.004)	NO SIG	0.086	3.7 (1.0)	4.4 (0.1)	NO SIG	0.222
GO:0006338	chromatin modeling	29	0.078 (0.047)	0.143 (0.004)	NO SIG	0.082	0.0 (0.3)	4.4 (0.1)	UNDER	0
GO:0016568	chromatin modification	29	0.078 (0.047)	0.143 (0.004)	NO SIG	0.082	0.0 (0.3)	4.4 (0.1)	UNDER	0
GO:0016582	non-covalent chromatin modification	29	0.078 (0.047)	0.143 (0.004)	NO SIG	0.082	0.0 (0.3)	4.4 (0.1)	UNDER	0
GO:0006886	intracellular protein transport	136	0.117 (0.018)	0.143 (0.004)	NO SIG	0.081	2.1 (0.6)	4.5 (0.1)	UNDER	4.07E-05
GO:0015031	protein transport	193	0.117 (0.018)	0.143 (0.004)	NO SIG	0.080	3.5 (0.6)	4.4 (0.1)	NO SIG	0.065
GO:0007160	cell-matrix adhesion	33	0.076 (0.047)	0.143 (0.004)	NO SIG	0.079	0.1 (0.5)	4.4 (0.1)	UNDER	2.98E-18
GO:0007154	cell communication	980	0.131 (0.009)	0.145 (0.004)	NO SIG	0.076	3.5 (0.3)	4.6 (0.2)	UNDER	6.03E-04
GO:0007017	microtubule-based process	58	0.103 (0.027)	0.143 (0.004)	NO SIG	0.072	1.8 (0.4)	4.4 (0.1)	UNDER	2.96E-10
GO:0019941	protein-ligand dependent protein degradation	87	0.114 (0.019)	0.143 (0.004)	NO SIG	0.070	1.3 (0.6)	4.4 (0.1)	UNDER	3.65E-08
GO:0008151	cell growth and/or maintenance	2818	0.137 (0.005)	0.148 (0.006)	NO SIG	0.069	4.4 (0.2)	4.4 (0.2)	NO SIG	0.489
GO:0007010	cytoskeleton organization and biogenesis	101	0.112 (0.020)	0.143 (0.004)	NO SIG	0.069	2.2 (0.5)	4.4 (0.1)	UNDER	2.61E-05
GO:0006508	proteolysis and peptidolysis	172	0.117 (0.017)	0.143 (0.004)	NO SIG	0.068	1.6 (0.5)	4.5 (0.1)	UNDER	2.75E-10
GO:0030163	protein degradation	172	0.117 (0.017)	0.143 (0.004)	NO SIG	0.068	1.6 (0.5)	4.5 (0.1)	UNDER	2.75E-10
GO:0006007	glucose catabolism	28	0.080 (0.041)	0.143 (0.004)	NO SIG	0.066	0.9 (1.6)	4.4 (0.1)	NO SIG	0.014
GO:0019320	hexose catabolism	28	0.080 (0.041)	0.143 (0.004)	NO SIG	0.066	0.9 (1.6)	4.4 (0.1)	NO SIG	0.014
GO:0007028	cytoplasm organization and biogenesis	158	0.116 (0.017)	0.143 (0.004)	NO SIG	0.065	1.9 (0.5)	4.5 (0.1)	UNDER	1.24E-06
GO:0009057	macromolecule catabolism	181	0.117 (0.017)	0.143 (0.004)	NO SIG	0.062	1.5 (0.4)	4.5 (0.1)	UNDER	1.91E-11
GO:0006445	regulation of translation	31	0.081 (0.039)	0.143 (0.004)	NO SIG	0.056	0.1 (0.4)	4.4 (0.1)	UNDER	0
GO:0005996	monosaccharide metabolism	39	0.085 (0.036)	0.143 (0.004)	NO SIG	0.054	0.7 (1.1)	4.4 (0.1)	UNDER	4.83E-04
GO:0019318	hexose metabolism	39	0.085 (0.036)	0.143 (0.004)	NO SIG	0.054	0.7 (1.1)	4.4 (0.1)	UNDER	4.83E-04
GO:0006006	glucose metabolism	32	0.080 (0.038)	0.143 (0.004)	NO SIG	0.051	0.8 (1.4)	4.4 (0.1)	UNDER	0.005
GO:0016052	carbohydrate catabolism	34	0.078 (0.037)	0.143 (0.004)	NO SIG	0.039	0.8 (1.3)	4.4 (0.1)	UNDER	0.003
GO:0006996	organelle organization and biogenesis	140	0.110 (0.018)	0.143 (0.004)	NO SIG	0.038	1.6 (0.4)	4.5 (0.1)	UNDER	5.91E-11
GO:0008150	biological_process	3460	0.137 (0.005)	0.151 (0.006)	NO SIG	0.036	4.3 (0.2)	4.5 (0.2)	NO SIG	0.304
GO:0006095	catabolic carbohydrate metabolism	42	0.080 (0.032)	0.143 (0.004)	NO SIG	0.025	0.6 (1.0)	4.4 (0.1)	UNDER	1.60E-04

GO:0006731	coenzymes and prosthetic group metabolism	32	0.075 (0.034)	0.143 (0.004)	NO SIG	0.023		0.0 (0.3)	4.4 (0.1)	UNDER	0
GO:0006092	main pathways of carbohydrate metabolism	43	0.080 (0.031)	0.143 (0.004)	NO SIG	0.023		0.6 (1.0)	4.4 (0.1)	UNDER	1.09E-04
GO:0007148	regulation of cell shape and cell size	57	0.085 (0.028)	0.143 (0.004)	NO SIG	0.022		1.9 (0.6)	4.4 (0.1)	UNDER	1.12E-05
GO:0006810	**transport**	**387**	**0.115 (0.013)**	**0.144 (0.004)**	**NO SIG**	**0.017**		**2.4 (0.4)**	**4.5 (0.1)**	**UNDER**	**1.68E-07**
GO:0015980	energy derivation by oxidation of organic compour	70	0.087 (0.025)	0.143 (0.004)	NO SIG	0.013		0.4 (0.6)	4.4 (0.1)	UNDER	3.67E-10
GO:0006936	muscle contraction	46	0.062 (0.036)	0.143 (0.004)	NO SIG	0.013		1.2 (1.2)	4.4 (0.1)	UNDER	0.005
GO:0009056	**catabolism**	**246**	**0.106 (0.015)**	**0.144 (0.004)**	**SLOWER**	**0.007**		**1.3 (0.4)**	**4.5 (0.1)**	**UNDER**	**1.92E-15**
GO:0006118	electron transport	53	0.067 (0.029)	0.143 (0.004)	SLOWER	0.005		0.1 (0.4)	4.4 (0.1)	UNDER	1.50E-20
GO:0006091	energy pathways	96	0.080 (0.024)	0.143 (0.004)	SLOWER	0.004		0.3 (0.5)	4.5 (0.1)	UNDER	1.84E-14
GO:0006457	protein folding	27	0.048 (0.033)	0.143 (0.004)	SLOWER	0.002		0.0 (0.0)	4.4 (0.1)	UNDER	0
GO:0007155	cell adhesion	132	0.082 (0.021)	0.144 (0.004)	SLOWER	0.002		1.9 (0.7)	4.5 (0.1)	UNDER	2.25E-04
GO:0009058	**biosynthesis**	**318**	**0.101 (0.012)**	**0.145 (0.004)**	**SLOWER**	**2.10E-04**		**1.8 (0.3)**	**4.6 (0.1)**	**UNDER**	**8.33E-18**
GO:0006411	**protein metabolism and modification**	**862**	**0.115 (0.008)**	**0.147 (0.004)**	**SLOWER**	**2.09E-04**		**2.5 (0.2)**	**4.8 (0.2)**	**UNDER**	**3.12E-15**
GO:0009059	macromolecule biosynthesis	201	0.084 (0.013)	0.144 (0.004)	SLOWER	7.86E-06		0.6 (0.2)	4.5 (0.1)	UNDER	0
GO:0006412	protein biosynthesis	175	0.077 (0.013)	0.144 (0.004)	SLOWER	5.18E-07		0.1 (0.2)	4.5 (0.1)	UNDER	0
MANUAL	**Swissprot transcription**		**0.252 (0.039)**	**0.137 (0.007)**	**FASTER**	**1.64E-07**		**15.5 (1.1)**	**3.9 (0.1)**	**OVER**	**0**

Supplementary Table 3: Functional Analysis of Decaying Transcripts in Yeast

This table shows results of a previously published yeast decay experiment analyzed with the same statistical inference procedures used in Supplementary Tables 1 and 2. Gene Ontology categories corresponding to those shown in Figure 1 are highlighted in bold. Manual classifications of transcription-related genes are attached to the end of this list. For the inference of percentage of fast decying transcripts, we defined "fast" as a half-life of ten minutes or less. Standard Deviations are shown in parentheses. P-values less than 1E-20 are indicated with a "0". We note that most of GO categories on this list appear to be decaying faster than the set of genes on the chip as a whole. This trend may be related to our observation that the 1811 genes of "unknown" function constitute one of the slowest decaying categories. Many of the genes in this category are predicted open reading frames with little characterization. Therefore, it is conceivable that transcripts of this type decay "slowly" because the microarray is detecting noise rather than actual transcripts for these genes. If true, this would lead to underestimates of decay rate and percent of fast decayers for the "non" group (g') genes. However, using the set of all GO-associated genes (i.e. the set of genes biased to move faster) rather than all genes as Sg yields similar results (i.e. transcription fast, metabolism slow): p-values are larger but still highly significant.

			Rate *r* (minutes^-1)				Half Life <10 min. %			
GO Number	Category Name	Probes	Group	Non-Group	Distribution	p-value	In-Group	Non-Group	Representation	p-value
GO:0006350	**transcription**	**223**	**0.082 (0.001)**	**0.055 (0.000)**	**FASTER**	**0**	**27.9 (1.4)**	**9.4 (0.2)**	**OVER**	**0**
GO:0006351	transcription, DNA-dependent	182	0.085 (0.001)	0.055 (0.000)	FASTER	0	30.4 (1.6)	9.5 (0.2)	OVER	0
GO:0006139	nucleobase, nucleoside, nucleotide and nucleic	443	0.073 (0.000)	0.054 (0.000)	FASTER	0	21.8 (0.9)	9.1 (0.2)	OVER	0
GO:0006360	transcription from Pol I promoter	78	0.105 (0.002)	0.055 (0.000)	FASTER	0	47.7 (2.5)	9.6 (0.2)	OVER	0
GO:0008151	cell growth and/or maintenance	1668	0.062 (0.000)	0.053 (0.000)	FASTER	0	13.7 (0.4)	8.4 (0.2)	OVER	0
GO:0016043	cell organization and biogenesis	416	0.068 (0.000)	0.055 (0.000)	FASTER	0	18.0 (0.8)	9.5 (0.2)	OVER	0
GO:0006364	rRNA processing	54	0.104 (0.002)	0.055 (0.000)	FASTER	0	46.2 (3.3)	9.9 (0.2)	OVER	0
GO:0006365	35S primary transcript processing	32	0.117 (0.003)	0.055 (0.000)	FASTER	0	54.8 (4.0)	10.0 (0.2)	OVER	0
GO:0007046	ribosome biogenesis	62	0.099 (0.002)	0.055 (0.000)	FASTER	0	42.3 (3.0)	9.8 (0.2)	OVER	0
GO:0016070	RNA metabolism	137	0.078 (0.001)	0.055 (0.000)	FASTER	0	27.8 (1.7)	9.7 (0.2)	OVER	0
GO:0006396	RNA processing	132	0.079 (0.001)	0.055 (0.000)	FASTER	0	28.2 (1.7)	9.8 (0.2)	OVER	0
GO:0008152	metabolism	1228	0.062 (0.000)	0.054 (0.000)	FASTER	0	14.0 (0.5)	8.9 (0.2)	OVER	0
GO:0042254	ribosome biogenesis and assembly	89	0.085 (0.001)	0.055 (0.000)	FASTER	0	32.7 (2.2)	9.8 (0.2)	OVER	0
GO:0007028	cytoplasm organization and biogenesis	319	0.068 (0.001)	0.055 (0.000)	FASTER	0	18.7 (0.9)	9.7 (0.2)	OVER	0
GO:0006366	transcription from Pol II promoter	76	0.075 (0.001)	0.056 (0.000)	FASTER	0	21.5 (2.3)	10.1 (0.2)	OVER	4.40E-07
GO:0006357	regulation of transcription from Pol II promoter	40	0.078 (0.002)	0.056 (0.000)	FASTER	0	28.7 (3.4)	10.1 (0.2)	OVER	1.57E-08
GO:0006355	regulation of transcription, DNA-dependent	67	0.073 (0.001)	0.056 (0.000)	FASTER	0	21.7 (2.5)	10.1 (0.2)	OVER	2.09E-06
GO:0006464	**protein modification**	**96**	**0.069 (0.001)**	**0.056 (0.000)**	**FASTER**	**0**	**14.0 (1.8)**	**10.2 (0.2)**	**NO SIG**	**0.020**
GO:0007600	sensory perception	30	0.078 (0.002)	0.056 (0.000)	FASTER	0	20.0 (2.3)	10.2 (0.2)	OVER	8.32E-06
GO:0007606	chemosensory perception	30	0.078 (0.002)	0.056 (0.000)	FASTER	0	20.0 (2.3)	10.2 (0.2)	OVER	8.32E-06
GO:0009582	perception of abiotic stimulus	30	0.078 (0.002)	0.056 (0.000)	FASTER	0	20.0 (2.3)	10.2 (0.2)	OVER	8.32E-06
GO:0009593	perception of chemical substance	30	0.078 (0.002)	0.056 (0.000)	FASTER	0	20.0 (2.3)	10.2 (0.2)	OVER	8.32E-06
GO:0019236	pheromone response	30	0.078 (0.002)	0.056 (0.000)	FASTER	0	20.0 (2.3)	10.2 (0.2)	OVER	8.32E-06
GO:0006259	**DNA metabolism**	**128**	**0.067 (0.001)**	**0.056 (0.000)**	**FASTER**	**0**	**18.2 (1.7)**	**10.1 (0.2)**	**OVER**	**6.64E-07**
GO:0006323	DNA packaging	38	0.081 (0.002)	0.056 (0.000)	FASTER	0	27.5 (3.7)	10.1 (0.2)	OVER	1.31E-06
GO:0006325	establishment and/or maintenance of chromatir	38	0.081 (0.002)	0.056 (0.000)	FASTER	0	27.5 (3.7)	10.1 (0.2)	OVER	1.31E-06
GO:0007049	cell cycle	245	0.064 (0.001)	0.055 (0.000)	FASTER	0	14.4 (1.1)	10.0 (0.2)	OVER	2.94E-05
GO:0000278	mitotic cell cycle	146	0.067 (0.001)	0.056 (0.000)	FASTER	0	16.6 (1.5)	10.1 (0.2)	OVER	8.36E-06
GO:0007001	chromosome organization and biogenesis (sens	41	0.079 (0.002)	0.056 (0.000)	FASTER	0	25.8 (3.5)	10.1 (0.2)	OVER	4.54E-06

GO:0007010	cytoskeleton organization and biogenesis	138	0.066 (0.001)	0.056 (0.000)	FASTER	0	15.0 (1.3)	10.1 (0.2)	OVER	9.57E-05
GO:0006997	nuclear organization and biogenesis	61	0.072 (0.001)	0.056 (0.000)	FASTER	0	19.4 (2.5)	10.2 (0.2)	OVER	1.33E-04
GO:0006810	**transport**	**291**	**0.062 (0.001)**	**0.055 (0.000)**	**FASTER**	**0**	**13.1 (0.8)**	**10.1 (0.2)**	**OVER**	**1.76E-04**
GO:0000084	S phase of mitotic cell cycle	48	0.071 (0.001)	0.056 (0.000)	FASTER	0	20.1 (2.3)	10.2 (0.2)	OVER	1.33E-05
GO:0006260	DNA replication	48	0.071 (0.001)	0.056 (0.000)	FASTER	0	20.1 (2.3)	10.2 (0.2)	OVER	1.33E-05
GO:0007005	mitochondrion organization and biogenesis	43	0.070 (0.001)	0.056 (0.000)	FASTER	0	20.4 (2.9)	10.2 (0.2)	OVER	2.43E-04
GO:0006261	DNA dependent DNA replication	41	0.072 (0.002)	0.056 (0.000)	FASTER	0	20.6 (2.6)	10.2 (0.2)	OVER	2.71E-05
GO:0006996	organelle organization and biogenesis	231	0.062 (0.001)	0.056 (0.000)	FASTER	0	13.2 (0.9)	10.1 (0.2)	OVER	7.72E-04
GO:0007275	**development**	**88**	**0.066 (0.001)**	**0.056 (0.000)**	**FASTER**	**0**	**14.7 (1.6)**	**10.2 (0.2)**	**OVER**	**0.003**
GO:0006411	**protein metabolism and modification**	**591**	**0.059 (0.000)**	**0.055 (0.000)**	**FASTER**	**0**	**11.5 (0.6)**	**10.1 (0.2)**	**NO SIG**	**0.021**
GO:0007124	pseudohyphal growth	27	0.072 (0.002)	0.056 (0.000)	FASTER	0	14.6 (3.1)	10.2 (0.2)	NO SIG	0.082
GO:0007150	growth pattern	27	0.072 (0.002)	0.056 (0.000)	FASTER	0	14.6 (3.1)	10.2 (0.2)	NO SIG	0.082
GO:0040007	growth	27	0.072 (0.002)	0.056 (0.000)	FASTER	0	14.6 (3.1)	10.2 (0.2)	NO SIG	0.082
GO:0007148	regulation of cell shape and cell size	39	0.075 (0.002)	0.056 (0.000)	FASTER	0	15.4 (2.4)	10.2 (0.2)	NO SIG	0.014
GO:0006352	transcription initiation	28	0.073 (0.002)	0.056 (0.000)	FASTER	1.15E-19	20.7 (4.6)	10.2 (0.2)	NO SIG	0.012
GO:0000067	DNA replication and chromosome cycle	89	0.064 (0.001)	0.056 (0.000)	FASTER	2.10E-19	15.7 (1.7)	10.2 (0.2)	OVER	7.87E-04
GO:0042255	ribosome assembly	34	0.069 (0.001)	0.056 (0.000)	FASTER	3.65E-19	24.6 (2.6)	10.2 (0.2)	OVER	1.55E-08
GO:0042257	ribosomal subunit assembly	34	0.069 (0.001)	0.056 (0.000)	FASTER	3.65E-19	24.6 (2.6)	10.2 (0.2)	OVER	1.55E-08
GO:0006796	phosphate metabolism	51	0.069 (0.001)	0.056 (0.000)	FASTER	2.76E-18	15.3 (2.0)	10.2 (0.2)	OVER	0.005
GO:0006468	protein amino acid phosphorylation	37	0.070 (0.002)	0.056 (0.000)	FASTER	1.10E-17	15.2 (2.1)	10.2 (0.2)	OVER	0.009
GO:0006486	protein amino acid glycosylation	37	0.069 (0.002)	0.056 (0.000)	FASTER	1.68E-17	13.3 (3.6)	10.2 (0.2)	NO SIG	0.198
GO:0009101	glycoprotein biosynthesis	37	0.069 (0.002)	0.056 (0.000)	FASTER	1.68E-17	13.3 (3.6)	10.2 (0.2)	NO SIG	0.198
GO:0007165	**signal transduction**	**70**	**0.066 (0.001)**	**0.056 (0.000)**	**FASTER**	**2.22E-17**	**14.6 (1.9)**	**10.2 (0.2)**	**NO SIG**	**0.011**
GO:0009100	glycoprotein metabolism	38	0.068 (0.001)	0.056 (0.000)	FASTER	2.16E-16	13.0 (3.5)	10.3 (0.2)	NO SIG	0.220
GO:0019538	protein metabolism	41	0.067 (0.001)	0.056 (0.000)	FASTER	1.72E-15	12.1 (3.3)	10.3 (0.2)	NO SIG	0.288
GO:0006886	intracellular protein transport	124	0.061 (0.001)	0.056 (0.000)	FASTER	9.28E-15	12.6 (1.3)	10.2 (0.2)	NO SIG	0.038
GO:0016310	phosphorylation	45	0.065 (0.001)	0.056 (0.000)	FASTER	3.65E-12	12.5 (1.7)	10.3 (0.2)	NO SIG	0.094
GO:0015031	protein transport	159	0.060 (0.001)	0.056 (0.000)	FASTER	5.02E-12	11.3 (1.1)	10.2 (0.2)	NO SIG	0.175
GO:0000283	establishment of cell polarity (sensu Saccharon	51	0.067 (0.002)	0.056 (0.000)	FASTER	1.38E-11	15.9 (2.0)	10.2 (0.2)	OVER	0.002
GO:0030010	establishment of cell polarity	51	0.067 (0.002)	0.056 (0.000)	FASTER	1.38E-11	15.9 (2.0)	10.2 (0.2)	OVER	0.002
GO:0030468	establishment of cell polarity (sensu Fungi)	51	0.067 (0.002)	0.056 (0.000)	FASTER	1.38E-11	15.9 (2.0)	10.2 (0.2)	OVER	0.002
GO:0006811	ion transport	34	0.064 (0.001)	0.056 (0.000)	FASTER	1.45E-11	14.0 (1.7)	10.2 (0.2)	NO SIG	0.016
GO:0009581	perception of external stimulus	57	0.064 (0.001)	0.056 (0.000)	FASTER	1.61E-11	13.8 (1.5)	10.2 (0.2)	NO SIG	0.011
GO:0009628	response to abiotic stimulus	57	0.064 (0.001)	0.056 (0.000)	FASTER	1.61E-11	13.8 (1.5)	10.2 (0.2)	NO SIG	0.011
GO:0007163	establishment and/or maintenance of cell polari	52	0.066 (0.002)	0.056 (0.000)	FASTER	4.03E-11	15.6 (2.0)	10.2 (0.2)	OVER	0.003
GO:0030012	establishment and/or maintenance of cell polari	52	0.066 (0.002)	0.056 (0.000)	FASTER	4.03E-11	15.6 (2.0)	10.2 (0.2)	OVER	0.003
GO:0030467	establishment and/or maintenance of cell polari	52	0.066 (0.002)	0.056 (0.000)	FASTER	4.03E-11	15.6 (2.0)	10.2 (0.2)	OVER	0.003
GO:0008380	RNA splicing	54	0.064 (0.001)	0.056 (0.000)	FASTER	1.01E-10	17.5 (2.2)	10.2 (0.2)	OVER	4.27E-04
GO:0006887	exocytosis	32	0.066 (0.002)	0.056 (0.000)	FASTER	7.25E-10	20.6 (3.0)	10.2 (0.2)	OVER	2.44E-04
GO:0009057	macromolecule catabolism	116	0.060 (0.001)	0.056 (0.000)	FASTER	2.25E-09	12.7 (1.4)	10.2 (0.2)	NO SIG	0.044
GO:0000226	microtubule cytoskeleton organization and biog	42	0.064 (0.001)	0.056 (0.000)	FASTER	2.81E-09	15.1 (2.5)	10.2 (0.2)	NO SIG	0.028
GO:0006605	protein targeting	77	0.061 (0.001)	0.056 (0.000)	FASTER	3.34E-09	13.1 (1.8)	10.2 (0.2)	NO SIG	0.060

GO:0006371	mRNA splicing	40	0.063 (0.001)	0.056 (0.000)	FASTER	1.21E-08	18.9 (2.4)	10.2 (0.2)	OVER	1.71E-04
GO:0006397	**mRNA processing**	**54**	**0.063 (0.001)**	**0.056 (0.000)**	**FASTER**	**2.60E-08**	**18.8 (2.2)**	**10.2 (0.2)**	**OVER**	**5.18E-05**
GO:0007154	cell communication	139	0.060 (0.001)	0.056 (0.000)	FASTER	5.13E-08	11.7 (1.2)	10.2 (0.2)	NO SIG	0.107
GO:0006947	cell-cell fusion	57	0.062 (0.001)	0.056 (0.000)	FASTER	2.84E-07	11.7 (1.4)	10.3 (0.2)	NO SIG	0.151
GO:0007322	mating (sensu Saccharomyces)	57	0.062 (0.001)	0.056 (0.000)	FASTER	2.84E-07	11.7 (1.4)	10.3 (0.2)	NO SIG	0.151
GO:0009292	genetic exchange	57	0.062 (0.001)	0.056 (0.000)	FASTER	2.84E-07	11.7 (1.4)	10.3 (0.2)	NO SIG	0.151
GO:0030461	mating (sensu Fungi)	57	0.062 (0.001)	0.056 (0.000)	FASTER	2.84E-07	11.7 (1.4)	10.3 (0.2)	NO SIG	0.151
GO:0007017	microtubule-based process	48	0.062 (0.001)	0.056 (0.000)	FASTER	3.75E-07	13.5 (2.3)	10.2 (0.2)	NO SIG	0.080
GO:0016192	vesicle-mediated transport	84	0.060 (0.001)	0.056 (0.000)	FASTER	1.20E-06	9.5 (1.3)	10.3 (0.2)	NO SIG	0.272
GO:0007242	intracellular signaling cascade	43	0.063 (0.001)	0.056 (0.000)	FASTER	1.36E-06	11.4 (2.8)	10.3 (0.2)	NO SIG	0.334
GO:0000074	**regulation of cell cycle**	**46**	**0.067 (0.002)**	**0.056 (0.000)**	**FASTER**	**1.43E-06**	**15.9 (2.7)**	**10.2 (0.2)**	**NO SIG**	**0.017**
GO:0009605	response to external stimulus	82	0.061 (0.001)	0.056 (0.000)	FASTER	1.72E-06	12.3 (1.5)	10.2 (0.2)	NO SIG	0.082
GO:0009306	protein secretion	59	0.061 (0.001)	0.056 (0.000)	FASTER	1.91E-06	11.7 (1.8)	10.3 (0.2)	NO SIG	0.218
GO:0045045	secretory pathway	59	0.061 (0.001)	0.056 (0.000)	FASTER	1.91E-06	11.7 (1.8)	10.3 (0.2)	NO SIG	0.218
GO:0000279	M phase	127	0.060 (0.001)	0.056 (0.000)	FASTER	2.31E-06	12.9 (1.5)	10.2 (0.2)	NO SIG	0.033
GO:0007151	sporulation (sensu Saccharomyces)	26	0.063 (0.002)	0.056 (0.000)	FASTER	2.49E-06	7.6 (1.8)	10.3 (0.2)	NO SIG	0.068
GO:0030437	sporulation (sensu Fungi)	26	0.063 (0.002)	0.056 (0.000)	FASTER	2.49E-06	7.6 (1.8)	10.3 (0.2)	NO SIG	0.068
GO:0006812	cation transport	28	0.062 (0.001)	0.056 (0.000)	FASTER	7.91E-06	13.4 (2.1)	10.3 (0.2)	NO SIG	0.068
GO:0016288	cytokinesis	34	0.064 (0.002)	0.056 (0.000)	FASTER	4.30E-05	14.9 (1.3)	10.2 (0.2)	OVER	2.66E-04
GO:0000087	M phase of mitotic cell cycle	66	0.060 (0.001)	0.056 (0.000)	FASTER	5.63E-05	12.3 (2.4)	10.2 (0.2)	NO SIG	0.193
GO:0007114	budding	61	0.060 (0.001)	0.056 (0.000)	FASTER	7.22E-05	12.5 (1.6)	10.2 (0.2)	NO SIG	0.085
GO:0007264	small GTPase mediated signal transduction	37	0.062 (0.002)	0.056 (0.000)	FASTER	7.75E-05	13.0 (3.1)	10.3 (0.2)	NO SIG	0.189
GO:0030435	sporulation	27	0.062 (0.002)	0.056 (0.000)	FASTER	1.62E-04	7.3 (1.7)	10.3 (0.2)	NO SIG	0.044
GO:0007067	mitosis	65	0.060 (0.001)	0.056 (0.000)	FASTER	1.70E-04	12.4 (2.4)	10.2 (0.2)	NO SIG	0.185
GO:0006281	DNA repair	53	0.060 (0.001)	0.056 (0.000)	FASTER	2.49E-04	12.4 (2.7)	10.2 (0.2)	NO SIG	0.212
GO:0009059	macromolecule biosynthesis	274	0.057 (0.000)	0.056 (0.000)	FASTER	0.001	9.7 (0.9)	10.3 (0.2)	NO SIG	0.263
GO:0006511	ubiquitin-dependent protein degradation	88	0.057 (0.001)	0.056 (0.000)	NO SIG	0.059	10.0 (1.7)	10.3 (0.2)	NO SIG	0.441
GO:0019941	protein-ligand dependent protein degradation	88	0.057 (0.001)	0.056 (0.000)	NO SIG	0.059	10.0 (1.7)	10.3 (0.2)	NO SIG	0.441
GO:0000280	nuclear division	88	0.057 (0.001)	0.056 (0.000)	NO SIG	0.121	11.0 (2.0)	10.3 (0.2)	NO SIG	0.361
GO:0006508	proteolysis and peptidolysis	98	0.056 (0.001)	0.056 (0.000)	NO SIG	0.225	9.9 (1.6)	10.3 (0.2)	NO SIG	0.407
GO:0000072	M-phase specific microtubule process	26	0.056 (0.001)	0.056 (0.000)	NO SIG	0.471	10.6 (3.6)	10.3 (0.2)	NO SIG	0.468
GO:0006412	protein biosynthesis	234	0.056 (0.000)	0.056 (0.000)	NO SIG	0.478	9.3 (0.9)	10.3 (0.2)	NO SIG	0.124
GO:0007121	polar budding	28	0.056 (0.002)	0.056 (0.000)	NO SIG	0.409	11.2 (2.5)	10.3 (0.2)	NO SIG	0.360
GO:0009058	**biosynthesis**	**377**	**0.055 (0.000)**	**0.056 (0.000)**	**NO SIG**	**0.168**	**8.6 (0.7)**	**10.4 (0.2)**	**UNDER**	**0.006**
GO:0007047	cell wall organization and biogenesis	55	0.055 (0.001)	0.056 (0.000)	NO SIG	0.149	8.0 (1.6)	10.3 (0.2)	NO SIG	0.067
GO:0045229	external protective structure organization and b	55	0.055 (0.001)	0.056 (0.000)	NO SIG	0.149	8.0 (1.6)	10.3 (0.2)	NO SIG	0.067
GO:0009607	response to biotic stimulus	26	0.054 (0.002)	0.056 (0.000)	NO SIG	0.122	8.5 (3.2)	10.3 (0.2)	NO SIG	0.293
GO:0008150	biological_process	3496	0.056 (0.000)	0.056 (0.000)	NO SIG	0.120	10.3 (0.2)	10.2 (0.4)	NO SIG	0.448
GO:0030029	actin filament-based process	37	0.054 (0.001)	0.056 (0.000)	NO SIG	0.092	5.5 (0.5)	10.3 (0.2)	UNDER	1.33E-18
GO:0030163	protein degradation	102	0.055 (0.001)	0.056 (0.000)	NO SIG	0.088	9.5 (1.5)	10.3 (0.2)	NO SIG	0.308
GO:0009056	**catabolism**	**149**	**0.055 (0.001)**	**0.056 (0.000)**	**NO SIG**	**0.046**	**11.1 (1.3)**	**10.2 (0.2)**	**NO SIG**	**0.250**
GO:0009308	amine metabolism	58	0.054 (0.001)	0.056 (0.000)	NO SIG	0.039	7.0 (1.7)	10.3 (0.2)	NO SIG	0.023

GO:0006520	amino acid metabolism	57	0.053 (0.001)	0.056 (0.000)	NO SIG	0.017		7.1 (1.7)	10.3 (0.2)	NO SIG	0.029
GO:0008652	amino acid biosynthesis	37	0.053 (0.002)	0.056 (0.000)	NO SIG	0.015		6.3 (1.7)	10.3 (0.2)	NO SIG	0.010
GO:0009309	amine biosynthesis	37	0.053 (0.002)	0.056 (0.000)	NO SIG	0.015		6.3 (1.7)	10.3 (0.2)	NO SIG	0.010
GO:0006519	amino acid and derivative metabolism	59	0.053 (0.001)	0.056 (0.000)	NO SIG	0.012		6.9 (1.6)	10.3 (0.2)	NO SIG	0.018
GO:0006873	ion homeostasis	38	0.054 (0.001)	0.056 (0.000)	NO SIG	0.011		3.5 (1.4)	10.3 (0.2)	UNDER	3.32E-07
GO:0019725	homeostasis	38	0.054 (0.001)	0.056 (0.000)	NO SIG	0.011		3.5 (1.4)	10.3 (0.2)	UNDER	3.32E-07
GO:0006629	lipid metabolism	78	0.054 (0.001)	0.056 (0.000)	SLOWER	0.008		6.3 (1.4)	10.3 (0.2)	UNDER	0.002
GO:0007015	actin filament organization	32	0.053 (0.001)	0.056 (0.000)	SLOWER	0.006		3.2 (0.6)	10.3 (0.2)	UNDER	0
GO:0006512	ubiquitin cycle	33	0.051 (0.002)	0.056 (0.000)	SLOWER	0.001		11.1 (2.9)	10.3 (0.2)	NO SIG	0.382
GO:0030036	actin cytoskeleton organization and biogenesis	34	0.052 (0.001)	0.056 (0.000)	SLOWER	0.001		3.0 (0.6)	10.3 (0.2)	UNDER	0
GO:0007126	meiosis	29	0.051 (0.002)	0.056 (0.000)	SLOWER	7.59E-04		8.6 (3.0)	10.3 (0.2)	NO SIG	0.291
GO:0006897	endocytosis	29	0.052 (0.001)	0.056 (0.000)	SLOWER	4.91E-05		3.5 (0.2)	10.3 (0.2)	UNDER	0
GO:0006950	stress response	53	0.048 (0.001)	0.056 (0.000)	SLOWER	8.84E-11		6.7 (2.1)	10.3 (0.2)	NO SIG	0.043
GO:0030003	cation homeostasis	31	0.048 (0.001)	0.056 (0.000)	SLOWER	3.23E-16		1.1 (1.7)	10.3 (0.2)	UNDER	1.27E-08
GO:0005996	monosaccharide metabolism	34	0.039 (0.001)	0.056 (0.000)	SLOWER	0		4.9 (2.6)	10.3 (0.2)	NO SIG	0.019
GO:0019318	hexose metabolism	34	0.039 (0.001)	0.056 (0.000)	SLOWER	0		4.9 (2.6)	10.3 (0.2)	NO SIG	0.019
GO:0005975	**carbohydrate metabolism**	**82**	**0.042 (0.001)**	**0.056 (0.000)**	**SLOWER**	**0**		**6.3 (1.3)**	**10.3 (0.2)**	**UNDER**	**0.001**
GO:0006006	glucose metabolism	28	0.036 (0.001)	0.056 (0.000)	SLOWER	0		4.0 (2.6)	10.3 (0.2)	UNDER	0.008
GO:0006095	catabolic carbohydrate metabolism	45	0.036 (0.001)	0.056 (0.000)	SLOWER	0		4.5 (1.7)	10.3 (0.2)	UNDER	4.39E-04
GO:0016052	carbohydrate catabolism	26	0.033 (0.001)	0.056 (0.000)	SLOWER	0		4.3 (2.8)	10.3 (0.2)	NO SIG	0.017
GO:0006092	main pathways of carbohydrate metabolism	49	0.035 (0.001)	0.056 (0.000)	SLOWER	0		4.2 (1.6)	10.3 (0.2)	UNDER	5.77E-05
GO:0006091	energy pathways	62	0.037 (0.001)	0.056 (0.000)	SLOWER	0		3.3 (1.3)	10.4 (0.2)	UNDER	1.30E-08
GO:0015980	energy derivation by oxidation of organic compo	62	0.037 (0.001)	0.056 (0.000)	SLOWER	0		3.3 (1.3)	10.4 (0.2)	UNDER	1.30E-08
GO:0000004	biological_process unknown	1811	0.050 (0.000)	0.059 (0.000)	SLOWER	0		7.2 (0.3)	12.2 (0.3)	UNDER	0
GO:0007033	vacuole organization and biogenesis	44	0.039 (0.001)	0.056 (0.000)	SLOWER	0		0.0 (0.1)	10.4 (0.2)	UNDER	0
MANUAL	**Description Line Transcription**		**0.069 (0.001)**	**0.055 (0.000)**	**FASTER**	**0**		**17.9 (1.2)**	**9.8 (0.2)**	**OVER**	**1.00E-05**
MANUAL	**Swissprot Transcription**		**0.069 (0.001)**	**0.055 (0.000)**	**FASTER**	**0**		**16.1 (1.4)**	**10.0 (0.2)**	**OVER**	**3.06E-11**

Supplementary Table 4: Number of Probe Sets Matched (Motif vs. Sequence Set)

The number of probe sets contributing to the analysis in Supplementary Tables 5 and 6 are indicated. Where the number of probe sets was less than 25, statistical testing was not performed ("n.a.").

Four HepG2 Experiments (Supplementary Table 5)

Name	Sequence	5'UTR	ORF	3'UTR	WHOLESEQ
1	[AT]ATTTA[AT]	1894	4148	7214	10217
2A	ATTTATTTATTTATTTATTTA	117	na	54	153
2B	ATTTATTTATTTATTTA	144	na	88	208
2C	[AT]ATTTATTTATTTA[AT]	186	na	162	302
2D	[AT]{2}ATTTATTTA[AT]{2}	271	na	431	574
2E	[AT]{4}ATTTA[AT]{4}	563	139	1625	1990
B1	TTTTAAAT	687	430	2865	3601
B-1	GGGCCTGG	593	1719	1119	2981
B2	TTTTAATTT	426	81	1266	1537
B-2	CCCAGCCCC	449	846	855	1843
B3	AAATATTTT	373	267	1175	1632
B4	AATATTTTT	315	118	1030	1304
H1	TTTTTTT	2049	1035	6386	8027
H-1	CCGCCTC	1460	2075	1226	3965
H2	TTTTTAAA	1029	640	4411	5350
H-2	CCAGCCTC	664	1549	1137	2910
H3	TTGTAAATA	172	107	641	884
MEG	TTATTTATT	456	118	946	1286
MEGSHORT	TATTTAT	946	1277	3538	4982

Single Bud8 Experiment (Supplementary Table 6)

Name	Sequence	5'UTR	ORF	3'UTR	WHOLESEQ
1	[AT]ATTTA[AT]	542	1179	2101	2952
2A	ATTTATTTATTTATTTATTTA	32	na	na	43
2B	ATTTATTTATTTATTTA	38	na	na	55
2C	[AT]ATTTATTTATTTA[AT]	50	na	43	79
2D	[AT]{2}ATTTATTTA[AT]{2}	70	na	117	155
2E	[AT]{4}ATTTA[AT]{4}	159	42	493	601
B1	TTTTAAAT	190	114	821	1021
B-1	GGGCCTGG	170	460	295	807
B2	TTTTAATTT	120	26	381	465
B-2	CCCAGCCCC	128	237	226	509
B3	AAATATTTT	99	71	332	457
B4	AATATTTTT	92	31	310	389
H1	TTTTTTT	561	299	1837	2295
H-1	CCGCCTC	407	560	331	1091
H2	TTTTTAAA	293	171	1280	1542
H-2	CCAGCCTC	178	426	309	794
H3	TTGTAAATA	52	31	178	250
MEG	TTATTTATT	118	33	279	367
MEGSHORT	TATTTAT	265	354	1041	1441

Supplementary Table 5: Motif Tabulations for 4x HepG2 Experiments

Several motifs were analyzed for their correlation with increased/decreased mRNA decay activity. Two basic inferences were used: 1) Changes in average decay rate (A-C = DRI) and 2) Changes in Percentage of "Fast" Decayers (D-E = PFDI). For both comparisons, the values for probe sets in genes with a motif in a particular position (A,D) were compared with the values for probe sets without the motif (B,D) to determine p-values (C,E). For the p-values, bold denotes increased rate or over-representation with 99% or greater probability and italics denotes decreased rate or under-representation. As described in the methods, motifs 1-2E and MEG/MEGSHORT were obtained from published studies. The other motifs were determined from analysis of multimers enriched in the decaying pool (see Experimental Procedures): "B" and "H" series motifs came from analysis of Bud8 and HepG2 decay respectively. "-" denotes underrepresentation in decaying sequence. "na" indicates where insufficient number of sequence/probe set matches (<= 25) precluded statistical inference. Standard deviations are shown in parentheses. P-values less than 1E-20 are indicated by "0".

A) In-Group Decay Rate (hour^-1)

	Decay Rate for Sequences with motif in...			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	0.149 (0.008)	0.167 (0.005)	0.177 (0.004)	0.164 (0.003)
2A	0.120 (0.026)	na	0.179 (0.039)	0.129 (0.024)
2B	0.111 (0.024)	na	0.144 (0.036)	0.118 (0.021)
2C	0.113 (0.021)	na	0.135 (0.025)	0.117 (0.017)
2D	0.118 (0.018)	na	0.154 (0.016)	0.144 (0.014)
2E	0.124 (0.014)	0.235 (0.030)	0.196 (0.009)	0.191 (0.008)
B1	0.138 (0.012)	0.219 (0.018)	0.173 (0.006)	0.171 (0.006)
B-1	0.125 (0.012)	0.126 (0.007)	0.120 (0.009)	0.126 (0.006)
B2	0.129 (0.015)	0.182 (0.039)	0.156 (0.010)	0.156 (0.009)
B-2	0.148 (0.014)	0.167 (0.011)	0.125 (0.010)	0.143 (0.007)
B3	0.124 (0.017)	0.192 (0.022)	0.189 (0.010)	0.184 (0.009)
B4	0.140 (0.018)	0.203 (0.036)	0.194 (0.011)	0.189 (0.010)
H1	0.139 (0.007)	0.153 (0.010)	0.166 (0.004)	0.159 (0.004)
H-1	0.142 (0.008)	0.150 (0.007)	0.119 (0.009)	0.141 (0.005)
H2	0.166 (0.010)	0.181 (0.014)	0.178 (0.005)	0.177 (0.005)
H-2	0.125 (0.012)	0.177 (0.008)	0.120 (0.009)	0.152 (0.006)
H3	0.145 (0.023)	0.195 (0.035)	0.221 (0.014)	0.202 (0.012)
MEG	0.133 (0.014)	0.159 (0.035)	0.190 (0.011)	0.182 (0.010)
MEGSHORT	0.146 (0.010)	0.165 (0.010)	0.189 (0.006)	0.177 (0.005)

B) Non-Group Decay Rate (hour^-1)

	Decay Rate for Sequences without motif in...			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	0.132 (0.002)	0.125 (0.002)	0.109 (0.003)	0.102 (0.003)
2A	0.134 (0.002)	na	0.134 (0.002)	0.134 (0.002)
2B	0.134 (0.002)	na	0.134 (0.002)	0.134 (0.002)
2C	0.134 (0.002)	na	0.134 (0.002)	0.134 (0.002)
2D	0.134 (0.002)	na	0.134 (0.002)	0.134 (0.002)
2E	0.134 (0.002)	0.133 (0.002)	0.128 (0.002)	0.128 (0.002)
B1	0.134 (0.002)	0.132 (0.002)	0.127 (0.002)	0.126 (0.002)
B-1	0.134 (0.002)	0.135 (0.002)	0.135 (0.002)	0.135 (0.002)
B2	0.134 (0.002)	0.134 (0.002)	0.132 (0.002)	0.132 (0.002)
B-2	0.134 (0.002)	0.133 (0.002)	0.134 (0.002)	0.133 (0.002)
B3	0.134 (0.002)	0.133 (0.002)	0.130 (0.002)	0.129 (0.002)
B4	0.134 (0.002)	0.134 (0.002)	0.131 (0.002)	0.130 (0.002)
H1	0.133 (0.002)	0.133 (0.002)	0.118 (0.003)	0.117 (0.003)
H-1	0.133 (0.002)	0.132 (0.002)	0.135 (0.002)	0.132 (0.003)
H2	0.132 (0.002)	0.132 (0.002)	0.121 (0.002)	0.118 (0.003)
H-2	0.134 (0.002)	0.130 (0.002)	0.135 (0.002)	0.131 (0.002)
H3	0.134 (0.002)	0.134 (0.002)	0.131 (0.002)	0.131 (0.002)
MEG	0.134 (0.002)	0.134 (0.002)	0.131 (0.002)	0.131 (0.002)
MEGSHORT	0.133 (0.002)	0.132 (0.002)	0.122 (0.002)	0.120 (0.002)

C) p-value for Decay-Rate Shift

	p-value for In-Group rate change			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	0.016	**2.02E-12**	**0**	**0**
2A	0.295	na	0.120	0.422
2B	0.169	na	0.394	0.211
2C	0.159	na	0.485	0.153
2D	0.185	na	0.096	0.231
2E	0.223	**3.50E-04**	**1.13E-13**	**1.38E-14**
B1	0.384	**1.48E-06**	**1.52E-11**	**1.57E-13**
B-1	0.233	0.118	0.053	0.056
B2	0.384	0.108	0.011	**0.005**
B-2	0.156	**7.72E-04**	0.174	0.080
B3	0.275	**0.004**	**6.05E-09**	**5.11E-10**
B4	0.364	0.027	**3.43E-09**	**6.77E-10**
H1	0.210	0.029	**0**	**3.50E-20**
H-1	0.168	**0.005**	0.036	0.060
H2	**3.83E-04**	**4.84E-04**	**0**	**0**
H-2	0.211	**1.93E-08**	0.059	**4.18E-04**
H3	0.310	0.040	**4.77E-11**	**6.96E-10**
MEG	0.479	0.235	**2.57E-07**	**1.00E-07**
MEGSHORT	0.112	**5.03E-04**	**0**	**0**

D) Percentage In-Group w/ Half-Life < 2 hr.

	% Transcripts with motif having Half-life < 2hr.			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	6.3 (0.3)	6.9 (0.2)	7.9 (0.2)	7.1 (0.1)
2A	5.1 (1.1)	na	9.5 (1.9)	5.2 (1.0)
2B	4.2 (0.9)	na	8.0 (1.5)	4.7 (0.8)
2C	3.8 (0.8)	na	6.7 (1.0)	4.5 (0.6)
2D	4.5 (0.7)	na	6.7 (0.7)	6.1 (0.6)
2E	4.2 (0.5)	14.5 (1.2)	9.8 (0.4)	9.6 (0.3)
B1	5.1 (0.5)	13.2 (0.8)	7.6 (0.3)	7.7 (0.2)
B-1	4.1 (0.4)	4.5 (0.3)	4.6 (0.3)	4.5 (0.2)
B2	3.8 (0.5)	9.1 (1.6)	6.4 (0.4)	6.3 (0.4)
B-2	4.9 (0.5)	7.0 (0.4)	3.5 (0.3)	5.0 (0.3)
B3	4.8 (0.6)	8.4 (0.9)	9.8 (0.4)	8.9 (0.4)
B4	5.3 (0.7)	11.7 (1.8)	9.5 (0.5)	9.3 (0.4)
H1	6.0 (0.3)	6.4 (0.4)	7.7 (0.2)	7.2 (0.2)
H-1	5.0 (0.3)	5.4 (0.3)	3.9 (0.3)	5.1 (0.2)
H2	7.7 (0.4)	8.7 (0.6)	8.4 (0.2)	8.2 (0.2)
H-2	3.8 (0.4)	8.6 (0.3)	3.1 (0.3)	6.1 (0.2)
H3	5.2 (1.1)	11.1 (1.5)	11.5 (0.7)	10.2 (0.5)
MEG	5.2 (0.5)	8.1 (1.5)	9.4 (0.5)	8.7 (0.4)
MEGSHORT	6.3 (0.4)	6.3 (0.4)	8.5 (0.3)	7.6 (0.2)

D) Percentage Non-Group w/ Half-Life < 2 hr.

	% Transcripts w/out motif having Half-life < 2hr.			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	5.0 (0.1)	4.6 (0.1)	3.5 (0.1)	3.0 (0.1)
2A	5.1 (0.1)	na	5.1 (0.1)	5.1 (0.1)
2B	5.1 (0.1)	na	5.1 (0.1)	5.1 (0.1)
2C	5.1 (0.1)	na	5.1 (0.1)	5.1 (0.1)
2D	5.1 (0.1)	na	5.1 (0.1)	5.1 (0.1)
2E	5.2 (0.1)	5.1 (0.1)	4.7 (0.1)	4.6 (0.1)
B1	5.1 (0.1)	4.9 (0.1)	4.7 (0.1)	4.6 (0.1)
B-1	5.2 (0.1)	5.2 (0.1)	5.2 (0.1)	5.2 (0.1)
B2	5.2 (0.1)	5.1 (0.1)	5.0 (0.1)	5.0 (0.1)
B-2	5.1 (0.1)	5.0 (0.1)	5.2 (0.1)	5.1 (0.1)
B3	5.1 (0.1)	5.1 (0.1)	4.8 (0.1)	4.8 (0.1)
B4	5.1 (0.1)	5.1 (0.1)	4.9 (0.1)	4.8 (0.1)
H1	5.0 (0.1)	5.1 (0.1)	3.9 (0.1)	3.7 (0.1)
H-1	5.1 (0.1)	5.1 (0.1)	5.2 (0.1)	5.1 (0.1)
H2	5.0 (0.1)	5.0 (0.1)	4.2 (0.1)	4.0 (0.1)
H-2	5.2 (0.1)	4.8 (0.1)	5.2 (0.1)	5.0 (0.1)
H3	5.1 (0.1)	5.1 (0.1)	4.9 (0.1)	4.9 (0.1)
MEG	5.1 (0.1)	5.1 (0.1)	4.9 (0.1)	4.9 (0.1)
MEGSHORT	5.1 (0.1)	5.0 (0.1)	4.4 (0.1)	4.3 (0.1)

D) p-value for Enrichment/Depletion of Motif

	p-value for In-Group Representation change			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	**2.95E-06**	**0**	**0**	**0**
2A	0.492	na	0.011	0.477
2B	0.142	na	0.033	0.282
2C	0.042	na	0.064	0.158
2D	0.172	na	**0.009**	0.042
2E	0.023	**1.70E-14**	**0**	**0**
B1	0.443	**0**	**0**	**0**
B-1	*0.004*	*0.007*	0.036	*5.65E-04*
B2	*0.007*	**0.008**	**7.24E-04**	**1.94E-04**
B-2	0.348	**2.96E-06**	*1.78E-07*	0.249
B3	0.298	**6.33E-05**	**0**	**0**
B4	0.404	**7.94E-05**	**2.00E-20**	**0**
H1	**2.33E-04**	**0.001**	**0**	**0**
H-1	0.384	0.151	*2.86E-05*	0.362
H2	**1.93E-10**	**2.03E-10**	**0**	**0**
H-2	*2.89E-04*	**0**	*8.56E-11*	**9.39E-07**
H3	0.460	**4.43E-05**	**0**	**0**
MEG	0.450	0.024	**7.50E-20**	**1.00E-20**
MEGSHORT	**8.75E-04**	**0.001**	**0**	**0**

Supplementary Table 6: Motif Tabulations for Bud8 Experiments

Several motifs were analyzed for their correlation with increased/decreased mRNA decay activity. Two basic inferences were used: 1) Changes in average decay rate (A-C = DRI) and 2) Changes in Percentage of "Fast" Decayers (D-E = PFDI). For both comparisons, the values for probe sets in genes with a motif in a particular position (A,D) were compared with the values for probe sets without the motif (B,D) to determine p-values (C,E). For the p-values, bold denotes increased rate or over-representation with 99% or greater probability and italics denotes decreased rate or under-representation. As described in the methods, motifs 1-2E and MEG/MEGSHORT were obtained from published studies. The other motifs were determined from analysis of multimers enriched in the decaying pool (see Experimental Procedures): "B" and "H" series motifs came from analysis of Bud8 and HepG2 decay respectively. "-" denotes underrepresentation in decaying sequence. "na" indicates where insufficient number of sequence/probe set matches (<= 25) precluded statistical inference. Standard deviations are shown in parentheses. P-values less than 1E-20 are indicated by "0".

A) In-Group Decay Rate (hour^-1)

	Decay Rate for Sequences with motif in...			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	0.154 (0.013)	0.161 (0.009)	0.178 (0.006)	0.166 (0.005)
2A	0.106 (0.056)	na	na	0.114 (0.049)
2B	0.101 (0.049)	na	na	0.099 (0.042)
2C	0.111 (0.042)	na	0.105 (0.041)	0.099 (0.033)
2D	0.115 (0.039)	na	0.160 (0.027)	0.148 (0.024)
2E	0.142 (0.026)	0.164 (0.045)	0.199 (0.014)	0.191 (0.013)
B1	0.144 (0.023)	0.224 (0.031)	0.171 (0.010)	0.173 (0.009)
B-1	0.119 (0.021)	0.147 (0.012)	0.127 (0.013)	0.138 (0.009)
B2	0.136 (0.028)	0.162 (0.066)	0.180 (0.015)	0.179 (0.014)
B-2	0.144 (0.024)	0.157 (0.015)	0.102 (0.015)	0.137 (0.011)
B3	0.139 (0.034)	0.170 (0.044)	0.179 (0.017)	0.173 (0.015)
B4	0.135 (0.036)	0.176 (0.059)	0.188 (0.017)	0.179 (0.016)
H1	0.149 (0.012)	0.146 (0.018)	0.169 (0.007)	0.163 (0.006)
H-1	0.148 (0.013)	0.150 (0.011)	0.131 (0.015)	0.146 (0.008)
H2	0.167 (0.017)	0.172 (0.025)	0.179 (0.008)	0.177 (0.007)
H-2	0.132 (0.022)	0.163 (0.013)	0.132 (0.014)	0.147 (0.010)
H3	0.183 (0.052)	0.231 (0.053)	0.242 (0.023)	0.228 (0.020)
MEG	0.124 (0.028)	0.136 (0.046)	0.197 (0.019)	0.183 (0.016)
MEGSHORT	0.134 (0.019)	0.152 (0.016)	0.191 (0.009)	0.176 (0.008)

D) Percentage In-Group w/ Half-Life < 2 hr.

	% Transcripts with motif having Half-life < 2hr.			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	5.1 (0.4)	5.2 (0.3)	6.9 (0.2)	5.8 (0.2)
2A	2.5 (1.4)	na	na	2.4 (1.5)
2B	2.1 (1.2)	na	na	1.9 (1.2)
2C	2.0 (1.2)	na	0.9 (1.4)	1.5 (0.9)
2D	2.0 (1.2)	na	6.4 (1.2)	5.1 (1.0)
2E	3.8 (0.9)	5.9 (1.5)	9.0 (0.6)	8.1 (0.5)
B1	4.4 (0.8)	10.7 (1.2)	6.5 (0.4)	6.4 (0.4)
B-1	2.4 (0.6)	5.6 (0.5)	3.8 (0.5)	4.6 (0.3)
B2	3.4 (1.0)	9.9 (2.3)	7.0 (0.6)	6.9 (0.6)
B-2	4.4 (0.9)	5.3 (0.7)	2.6 (0.5)	3.8 (0.4)
B3	4.7 (1.0)	6.6 (1.6)	7.1 (0.6)	6.8 (0.5)
B4	3.7 (1.0)	7.3 (2.5)	7.5 (0.7)	6.9 (0.6)
H1	5.3 (0.4)	2.8 (0.5)	6.9 (0.3)	6.2 (0.2)
H-1	5.3 (0.4)	5.4 (0.5)	4.5 (0.5)	5.1 (0.3)
H2	5.8 (0.6)	5.8 (1.0)	7.5 (0.3)	7.1 (0.3)
H-2	3.7 (0.8)	6.0 (0.5)	3.0 (0.5)	4.7 (0.4)
H3	5.9 (2.0)	14.3 (2.6)	12.3 (1.0)	11.0 (0.9)
MEG	3.3 (0.9)	0.3 (0.9)	9.1 (0.8)	7.7 (0.6)
MEGSHORT	4.8 (0.5)	3.9 (0.6)	7.6 (0.4)	6.3 (0.3)

B) Non-Group Decay Rate (hour^-1)

	Decay Rate for Sequences without motif in...			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	0.141 (0.004)	0.137 (0.004)	0.120 (0.005)	0.115 (0.005)
2A	0.142 (0.004)	na	na	0.142 (0.004)
2B	0.143 (0.004)	na	na	0.143 (0.004)
2C	0.142 (0.004)	na	0.142 (0.004)	0.143 (0.004)
2D	0.143 (0.004)	na	0.142 (0.004)	0.142 (0.004)
2E	0.142 (0.004)	0.142 (0.004)	0.137 (0.004)	0.136 (0.004)
B1	0.142 (0.004)	0.140 (0.004)	0.137 (0.004)	0.135 (0.004)
B-1	0.143 (0.004)	0.142 (0.004)	0.143 (0.004)	0.143 (0.004)
B2	0.142 (0.004)	0.142 (0.004)	0.139 (0.004)	0.139 (0.004)
B-2	0.142 (0.004)	0.142 (0.004)	0.144 (0.004)	0.143 (0.004)
B3	0.142 (0.004)	0.142 (0.004)	0.140 (0.004)	0.139 (0.004)
B4	0.142 (0.004)	0.142 (0.004)	0.139 (0.004)	0.139 (0.004)
H1	0.141 (0.004)	0.142 (0.004)	0.129 (0.004)	0.127 (0.005)
H-1	0.142 (0.004)	0.141 (0.004)	0.143 (0.004)	0.141 (0.004)
H2	0.141 (0.004)	0.141 (0.004)	0.131 (0.004)	0.129 (0.004)
H-2	0.143 (0.004)	0.140 (0.004)	0.143 (0.004)	0.141 (0.004)
H3	0.142 (0.004)	0.142 (0.004)	0.139 (0.004)	0.138 (0.004)
MEG	0.143 (0.004)	0.142 (0.004)	0.139 (0.004)	0.139 (0.004)
MEGSHORT	0.143 (0.004)	0.141 (0.004)	0.131 (0.004)	0.130 (0.004)

D) Percentage Non-Group w/ Half-Life < 2 hr.

	% Transcripts w/out motif having Half-life < 2hr.			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	4.3 (0.1)	4.2 (0.1)	2.9 (0.2)	2.8 (0.2)
2A	4.4 (0.1)	na	na	4.4 (0.1)
2B	4.4 (0.1)	na	na	4.4 (0.1)
2C	4.4 (0.1)	na	4.4 (0.1)	4.4 (0.1)
2D	4.4 (0.1)	na	4.4 (0.1)	4.4 (0.1)
2E	4.4 (0.1)	4.4 (0.1)	3.9 (0.1)	3.9 (0.1)
B1	4.4 (0.1)	4.3 (0.1)	4.0 (0.1)	3.9 (0.1)
B-1	4.5 (0.1)	4.3 (0.1)	4.4 (0.1)	4.4 (0.1)
B2	4.4 (0.1)	4.4 (0.1)	4.2 (0.1)	4.2 (0.1)
B-2	4.4 (0.1)	4.4 (0.1)	4.5 (0.1)	4.5 (0.1)
B3	4.4 (0.1)	4.4 (0.1)	4.2 (0.1)	4.2 (0.1)
B4	4.4 (0.1)	4.4 (0.1)	4.2 (0.1)	4.2 (0.1)
H1	4.3 (0.1)	4.5 (0.1)	3.1 (0.2)	3.1 (0.2)
H-1	4.3 (0.1)	4.3 (0.1)	4.4 (0.1)	4.2 (0.1)
H2	4.3 (0.1)	4.4 (0.1)	3.5 (0.1)	3.3 (0.1)
H-2	4.4 (0.1)	4.3 (0.1)	4.5 (0.1)	4.3 (0.1)
H3	4.4 (0.1)	4.3 (0.1)	4.1 (0.1)	4.1 (0.1)
MEG	4.4 (0.1)	4.4 (0.1)	4.1 (0.1)	4.2 (0.1)
MEGSHORT	4.4 (0.1)	4.4 (0.1)	3.6 (0.1)	3.7 (0.2)

C) p-value for Decay-Rate Shift

	p-value for In-Group rate change			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	0.164	**0.010**	**1.92E-13**	**6.14E-12**
2A	0.261	na	na	0.282
2B	0.202	na	na	0.150
2C	0.230	na	0.178	0.092
2D	0.240	na	0.250	0.396
2E	0.489	0.312	**5.50E-06**	**2.18E-05**
B1	0.469	**0.004**	**0.001**	**1.50E-04**
B-1	0.130	0.352	0.126	0.310
B2	0.408	0.384	**0.005**	**0.003**
B-2	0.467	0.166	*0.003*	0.319
B3	0.461	0.265	0.011	0.014
B4	0.424	0.284	**0.003**	**0.007**
H1	0.270	0.412	**3.32E-07**	**1.22E-06**
H-1	0.323	0.220	0.224	0.280
H2	0.071	0.108	**1.39E-07**	**9.70E-09**
H-2	0.317	0.055	0.226	0.277
H3	0.217	0.047	**3.45E-06**	**7.95E-06**
MEG	0.257	0.450	**0.001**	**0.004**
MEGSHORT	0.325	0.253	**8.05E-10**	**1.40E-07**

D) p-value for Enrichment/Depletion of Motif

	p-value for In-Group Representation change			
Motif	5'UTR	ORF	3'UTR	WHOLESEQ
1	0.038	**0.002**	**0**	**0**
2A	0.088	na	na	0.091
2B	0.026	na	na	0.015
2C	0.021	na	*0.005*	*9.96E-04*
2D	0.022	na	0.052	0.239
2E	0.246	0.156	**4.22E-16**	**1.44E-13**
B1	0.490	**3.78E-08**	**7.61E-09**	**2.83E-10**
B-1	*1.84E-04*	**0.007**	0.104	0.275
B2	0.151	**0.009**	**1.06E-05**	**1.37E-06**
B-2	0.488	0.092	*8.51E-05*	0.081
B3	0.376	0.073	**7.31E-06**	**2.78E-06**
B4	0.246	0.126	**7.70E-07**	**3.96E-06**
H1	0.016	*0.001*	**0**	**0**
H-1	0.017	**0.009**	0.454	**0.008**
H2	**0.009**	0.075	**0**	**0**
H-2	0.161	**0.001**	*0.001*	0.173
H3	0.232	**6.20E-05**	**3.44E-16**	**2.70E-15**
MEG	0.099	*3.38E-06*	**4.44E-11**	**1.17E-08**
MEGSHORT	0.223	0.163	**0**	**2.22E-15**

Supplementary Table 7: Affymetrix U95Av2 Accessions with "Transcription" Gene Ontology Codes

These genes were classified as belonging to the process "transcription" according to the Compugen Gene Ontology resource.

Accession	Gene Description		Accession	Gene Description
U59302	HSU59302 Human steroid receptor coactivator-1		U07132	HSU07132 Human steroid hormone receptor Ner-l
U68233	Human farnesol receptor HRR-1 (HRR-1) mRNA, complete c		X62534	H.sapiens HMG-2 mRNA /cds=(214,843) /gb=X62534 /gi=323
L37368	Human (clone E5.1) RNA-binding protein mRNA, complete		U50079	HSU50079 Human histone deacetylase HD1 mRNA,
AF039081	Homo sapiens Cre binding protein-like 2 mRNA, complete		U39840	Human hepatocyte nuclear factor-3 alpha (HNF-3 alpha)
L07758	Human IEF SSP 9502 mRNA, complete cds /cds=(87,1592) /		AJ007989	Homo sapiens mRNA for transcription factor TBX6 /cds=(
X63564	H.sapiens mRNA for RNA polymerase II largest subunit /		X74142	H.sapiens HBF-1 mRNA for transcription factor /cds=(21
J05448	HUMRPOLAA Human RNA polymerase subunit hRPB 3		X74143	H.sapiens HBF-2 mRNA for transcription factor /cds=(20
L37127	HUMRPIA Homo sapiens RNA polymerase II mRNA,		M29039	HUMJUNCAA Human transactivator (jun-B) gene,
U75370	Human mitochondrial RNA polymerase mRNA, nuclear gene		M24748	HUMTHRA1A Human thyroid hormone receptor alph
U37690	HSU37690 Human RNA polymerase II subunit (hsR		M95925	Human leucine zipper on the D14S46E locus mRNA, comple
X82125	H.sapiens HOK-2 mRNA for zinc finger protein /cds=(653		M95929	Human homeobox protein (PHOX1) mRNA, 3 end /cds=(0,59
X84373	H.sapiens mRNA for nuclear factor RIP140 /cds=(287,376		M97388	HUMDR1TATA Human TATA binding protein-associa
X89416	HSRNAPPP5 H.sapiens mRNA for protein phosphat		AF009801	Homo sapiens homeodomain protein (BAPX1) mRNA, complet
AF008442	Homo sapiens RNA polymerase I subunit hRPA39 mRNA, com		L29277	HUMAPRF Homo sapiens DNA-binding protein (APR
D38251	Homo sapiens mRNA for RPB5 (XAP4), complete cds /cds=(		AB006684	Homo sapiens APECED gene for AIRE-1, AIRE-2, AIRE-3, c
L40407	Homo sapiens thyroid receptor interactor (TRIP9) gene,		AF028706	Homo sapiens zinc-finger protein of the cerebellum 3 (
AB002388	Human mRNA for KIAA0390 gene, complete cds /cds=(138,4		Y00664	Human germ line n-myc gene /cds=(141,1511) /gb=Y00664
AF041259	Homo sapiens breast cancer putative transcription fact		Z36715	H.sapiens mRNA for Net transcription factor /cds=(279,
D86975	Human mRNA for KIAA0222 gene, complete cds /cds=(318,3		M98833	Human ERGB transcription factor (FLI-1 homolog) mRNA,
M26747	HUMCERBA Human c-erbA mRNA, complete cds"		U09366	Human zinc finger protein ZNF133 /cds=(445,2409) /gb=U
M69238	Human aryl hydrocarbon receptor nuclear translocator (		U23946	HSU23946 Human putative tumor suppressor (LUC
X63563	H.sapiens mRNA for RNA polymerase II 140 kDa subunit /		X76342	H.sapiens ADH7 mRNA /cds=UNKNOWN /gb=X76342 /gi=541674
U29332	Homo sapiens heart protein (FHL-2) mRNA, complete cds		U09367	Human zinc finger protein ZNF136 /cds=(71,1693) /gb=U0
M29550	Human calcineurin A1 mRNA, complete cds /cds=(107,1651		U09368	Human zinc finger protein ZNF140 /cds=(272,1645) /gb=U
M29551	Human calcineurin A2 mRNA, complete cds /cds=(116,1690		D12765	HUME1AF Human mRNA for E1A-F
D17532	Human mRNA for RCK, complete cds /cds=(339,1757) /gb=D		L19871	HUMATF3X Human activating transcription facto
M10901	HUMGCRA Human glucocorticoid receptor alpha m		L19872	Human AH-receptor mRNA, complete cds /cds=(375,2921) /
U37689	HSU37689 Human RNA polymerase II subunit (hsR		U28251	Human Krueppel-type zinc finger protein (ZNF169) gene,
U22431	HSU22431 Human hypoxia-inducible factor 1 alp		U31986	Human cartilage-specific homeodomain protein Cart-1 mR
U89387	Human RNA polymerase II subunit hsRPB4 gene, complete		M27691	HUMCREB Human transactivator protein (CREB) m
U52427	Human RNA polymerase II seventh subunit (rpb-7) gene,		X79200	HSSYTSSX Homo spaiens mRNA for SYT-SSX protei
AF011573	Homo sapiens zinc finger protein (ZnF20) mRNA, complet		X12492	HSCTF1 Human mRNA for CAAT-box binding transc
AF040253	Homo sapiens transcription factor Tat-CT1 mRNA, comple		X79201	H.sapiens mRNA for SYT /cds=(3,1178) /gb=X79201 /gi=53
AF008203	Homo sapiens homeobox protein (Alx3) mRNA, complete cd		S66431	S66431 RBP2=retinoblastoma binding protein 2
AB020700	Homo sapiens mRNA for KIAA0893 protein, complete cds /		AB002332	Human mRNA for KIAA0334 gene, complete cds /cds=(251,2
AB023232	Homo sapiens mRNA for KIAA1015 protein, complete cds /		U73328	Human DLX7 (Dlx7) mRNA, complete cds /cds=(246,749) /g
M69296	Human estrogen receptor-related protein (variant ER fr		AJ005814	Homo sapiens mRNA for hoxA7 protein /cds=(106,798) /gb
AB014528	Homo sapiens mRNA for KIAA0628 protein, complete cds /		M80783	Human B12 protein mRNA, complete cds /cds=(153,1103) /
U22377	Human Zn-15 related zinc finger protein (rlf) mRNA, co		X06614	HSRRA Human mRNA for receptor of retinoic aci
AB014569	Homo sapiens mRNA for KIAA0669 protein, complete cds /		M30607	Human zinc finger protein Y-linked (ZFY) mRNA, complet
AF044288	Homo sapiens basic-helix-loop-helix-PAS orphan MOP3 (M		AJ000041	Homo sapiens mRNA for HOXC11 /cds=(44,958) /gb=AJ00004
AF025770	Homo sapiens C2H2 zinc finger protein (ZNF189) mRNA, c		U13044	Human nuclear respiratory factor-2 subunit alpha mRNA,
AB019246	Homo sapiens mRNA for FTZ-F1 related protein, complete		L76569	Homo sapiens (clones cYG3, B5P6C4) fragile X E mental
U52191	Human SMCY (H-Y) mRNA, complete cds /cds=(275,4894) /g		M24069	Human DNA-binding protein A (dbpA) gene, 3 end /cds=(
U24576	U24576 Homo sapiens breast tumor autoantigen		L75847	Human zinc finger protein 45 (ZNF45) mRNA, complete cd
AF038451	Homo sapiens secreted cement gland protein XAG-2 homol		X07384	Human mRNA for GLI protein /cds=(78,3398) /gb=X07384 /
D50419	Homo sapiens mRNA for OTK18, complete cds /cds=(345,24		U80669	HSU80669 Homo sapiens androgen regulated home
L22342	Human nuclear phosphoprotein mRNA, complete cds /cds=(		M83667	HUMNFIL6BA Human NF-IL6-beta protein mRNA, co
S68134	CREM=cyclic AMP-responsive element modulator beta isof		X58964	H.sapiens gene for MHC class II regulatory factor RFX
U62392	Homo sapiens zinc finger protein mRNA, complete cds /c		X58965	HSNM23H2G H.sapiens RNA for nm23-H2 gene
AF059194	Homo sapiens basic-leucine zipper transcription factor		Z46967	H.sapiens mRNA for calicin (partial) /cds=(0,1757) /gb
AB002294	Human mRNA for KIAA0296 gene, complete cds /cds=(423,5		L14812	HUMP107B Human retinoblastoma related protein
M82827	Human fusion protein mRNA, complete cds /cds=(324,857)		U38480	HSU38480 Human retinoid X receptor-gamma mRNA
U62767	Human PP35 mRNA, complete cds /cds=(191,1114) /gb=U627		V00568	HSMYC1 Human mRNA encoding the c-myc oncogene
AL050276	Homo sapiens mRNA; cDNA DKFZp566F123 (from clone DKFZp		S69369	S69369 PAX3A=transcription factor [human, adu
AF019214	Homo sapiens HMG box containing protein 1 mRNA, comple		S69370	S69370 PAX3B=transcription factor {alternativ
D43945	Homo sapiens mRNA for TFEC isoform (or TFECL), complet		Z49825	H.sapiens mRNA for hepatocyte nuclear factor 4 alpha /
U80987	HSU80987 Human transcription factor TBX5 mRNA		Z49826	H.sapiens mRNA for hepatocyte nuclear factor 4 gamma /
U69274	Human zinc finger protein mRNA, complete cds /cds=(161		U31248	Human zinc finger protein (ZNF174) mRNA, complete cds
AB028021	Homo sapiens HNF-3beta mRNA for hepatocyte nuclear fac		X51730	HSPREC Human mRNA and promoter DNA for proges
U43189	HSU43189 Human Ets transcription factors NERF		X96744	H.sapiens PAX7 gene, exon 1 (and joined CDS) /cds=(599
M99587	Homo sapiens chromosome 4p homeobox mRNA sequence /cds		M86546	H.sapiens PBX1a and PBX1b mRNA, complete cds /cds=(123
AB002333	Human mRNA for KIAA0335 gene, complete cds /cds=(226,4		U90840	Homo sapiens SSX3 (SSX3) mRNA, complete cds /cds=(53,6
AB002350	Human mRNA for KIAA0352 gene, complete cds /cds=(86,22		U90841	Homo sapiens SSX4 (SSX4) mRNA, complete cds /cds=(0,56
U57317	HSU57317 Homo sapiens p300/CBP-associated fac		U90842	Homo sapiens SSX5 (SSX5) mRNA, complete cds /cds=(0,56
AF038960	Homo sapiens SKD1 homolog mRNA, complete cds /cds=(189		X71346	Homo sapiens HNF1-B mRNA /cds=UNKNOWN /gb=X71346 /gi=4
U70862	HSU70862 Human nuclear factor I B3 mRNA, comp		X71348	Homo sapiens vHNF1-C mRNA /cds=UNKNOWN /gb=X71348 /gi=
AF022654	Homo sapiens homeodomain protein (OG12) mRNA, complete		M13929	HUMMYCPOA Human c-myc-P64 mRNA, initiating fr
AF068744	Homo sapiens double homeodomain protein (DUX2) mRNA, c		S62138	TLS/CHOP=hybrid gene {translocation breakpoint} [human
U38864	Human zinc-finger protein C2H2-150 mRNA, complete cds		L01042	Human HIV1 tata element modulatory factor mRNA sequenc
U57796	Human zinc finger protein (LD5-1) mRNA, complete cds /		S62140	TLS=translocated in liposarcoma [human, mRNA, 1824 nt]
AB018326	Homo sapiens mRNA for KIAA0783 protein, complete cds /		M77810	HUMGATA2A Human transcription factor GATA-2 (
AF055376	Homo sapiens short form transcription factor C-MAF (c-		L07592	Human peroxisome proliferator activated receptor mRNA,
AB011107	Homo sapiens mRNA for KIAA0535 protein, complete cds /		Y08639	H.sapiens mRNA for nuclear orphan receptor ROR-beta /c
U43292	Human MDS1B (MDS1) mRNA, complete cds /cds=(307,816) /		U11690	Human faciogenital dysplasia (FGD1) mRNA, complete cds
AF072810	Homo sapiens transcription factor WSTF mRNA, complete		AJ010277	Homo sapiens mRNA for TBX19 protein /cds=(51,1397) /gb
AF028840	Homo sapiens Kruppel-associated box protein mRNA, comp		S67970	ZNF75=KRAB zinc finger [human, lung fibroblast, mRNA,
AF064804	Homo sapiens transcription factor SUPT3H (SUPT3H) mRNA		M21985	HUMSRTR2A Human steroid receptor TR2 mRNA, co

AF074606	Homo sapiens histone acetyltransferase (HBO1) mRNA, co		U85707	Human leukemogenic homolog protein (MEIS1) mRNA, compl
AB011141	Homo sapiens mRNA for KIAA0569 protein, complete cds /		M16801	Human mineralocorticoid receptor mRNA (hMR), complete
AB007886	Homo sapiens KIAA0426 mRNA, complete cds /cds=(145,195		L26494	Homo sapiens (oct-6) mRNA, complete cds /cds=(41,1387)
D87673	D87673 Homo sapiens mRNA for heat shock trans		L06895	HUMMAD Homo sapiens antagonizer of myc transc
M29960	HUMTR211 Human steroid receptor (TR2-11) mRNA		X63380	Homo sapiens mRNA for serum response factor-related pr
AB020639	Homo sapiens mRNA for KIAA0832 protein, complete cds /		X70683	H.sapiens mRNA for SOX-4 protein /cds=(350,1774) /gb=X
U12431	Human ELAV-like neuronal protein 1 (hel-N1) mRNA, comp		U25435	Human transcriptional repressor (CTCF) mRNA, complete
D86966	Human mRNA for KIAA0211 gene, complete cds /cds=(570,4		AF060865	Homo sapiens chromosome 16 zinc finger protein ZNF210
D86969	Human mRNA for KIAA0215 gene, complete cds /cds=(298,2		AF012130	Homo sapiens brachyury variant A (TBX1) mRNA, complete
AB028964	Homo sapiens mRNA for KIAA1041 protein, complete cds /		AF012131	Homo sapiens brachyury variant B (TBX1) mRNA, complete
AB023162	Homo sapiens mRNA for KIAA0945 protein, complete cds /		U09410	Human zinc finger protein ZNF131 mRNA, partial cds /cd
AF060503	Homo sapiens zinc finger protein (ZF5128) mRNA, comple		U09411	Human zinc finger protein ZNF132 mRNA, complete cds /c
U16028	HSU16028 Human CRE-BP1 transcription factor m		U09412	Human zinc finger protein ZNF134 mRNA, complete cds /c
AB023189	Homo sapiens mRNA for KIAA0972 protein, complete cds /		U09413	Human zinc finger protein ZNF135 mRNA, complete cds /c
U08015	Human NF-ATc mRNA, complete cds /cds=(239,2389) /gb=U0		U09414	Human zinc finger protein ZNF137 mRNA, complete cds /c
U85430	Human transcription factor NFATx4 mRNA, complete cds /		U89331	Human pseudoautosomal homeodomain-containing protein (
U51334	Human putative RNA binding protein (RBP56) mRNA, compl		M24899	Human triiodothyronine (ear7) mRNA, complete cds /cds=
U22680	Human X2 box repressor mRNA, complete cds /cds=(165,34		X69115	H.sapiens ZNF37A mRNA for zinc finger protein /cds=(0,
AF096870	Homo sapiens estrogen-responsive B box protein (EBBP)		U88629	HSU88629 Human RNA polymerase II elongation f
M55422	Human Krueppel-related zinc finger protein (H-plk) mRN		M19720	HUMMYC3L Human L-myc protein gene, complete c
AF074382	Homo sapiens IkB kinase gamma subunit (IKK-gamma) mRNA		M93119	Human zinc-finger DNA-binding motifs (IA-1) mRNA, comp
X91868	H.sapiens mRNA for SIX1 protein /cds=(275,1129) /gb=X9		U35612	Homo sapiens SOX22 protein (SOX22) mRNA, complete cds
M57609	Human DNA-binding protein (GLI3) mRNA, complete cds /c		AB008913	Homo sapiens mRNA for Pax-4, complete cds /cds=(0,1052
D28118	HUMDB1 Human mRNA for DB1, complete cds"		X12556	HSDBLPRO Human mRNA for dbl proto-oncogene
U70663	Human zinc finger transcription factor hEZF (EZF) mRNA		D31716	Human mRNA for GC box bindig protein, complete cds /cd
Z30425	HSONHORE H.sapiens mRNA for orphan nuclear ho		AF071771	Homo sapiens SPH-binding factor mRNA, partial cds /cds
X68560	H.sapiens SPR-2 mRNA for GT box binding protein /cds=(		L12701	HUMEN2A02 Human engrailed protein (EN2) gene,
X68561	H.sapiens SPR-1 mRNA for GT box binding protein /cds=(		M13228	HUMNMCY1A Human N-myc oncogene protein mRNA
X95463	H.sapiens mRNA for ox19 protein /cds=(88,1371) /gb=X95		X16155	Human mRNA for chicken ovalbumin upstream promoter tra
L17131	Human high mobility group protein (HMG-I(Y)) gene exon		M74297	HUMHOX14 Human homeobox 1.4 protein mRNA, com
X16281	Human mRNA for zinc finger protein (clone 431) /cds=(0		U26914	HSU26914 Human ras-responsive element binding
X16282	Human mRNA for zinc finger protein (clone 647) /cds=(0		M85164	Homo sapiens SRF accessory protein 1B (SAP-1) mRNA, co
AF041339	Homo sapiens homeodomain protein (PITX3) mRNA, complet		M85165	Homo sapiens SRF accessory protein 1A (SAP-1) mRNA, co
U13219	Human forkhead protein FREAC-1 mRNA, complete cds /cds		U65404	HSU65404 Human erythroid-specific transcripti
X59841	Human PBX3 mRNA /cds=UNKNOWN /gb=X59841 /gi=35314 /		S82592	Evi-1=Evi-1 protein {3 region, deletion region} [human
X59842	Human PBX2 mRNA /cds=UNKNOWN /gb=X59842 /gi=35312 /		X59739	Human ZFX mRNA for put. transcription activator, isofo
U13220	Homo sapiens forkhead protein FREAC-2 mRNA, complete c		U34962	Human transcription factor HCSX (hCsx) mRNA, complete
U19765	Human nucleic acid binding protein gene, complete cds		M58297	Human zinc finger protein 42 (MZF-1) mRNA, complete cd
X59871	Human TCF-1 mRNA for T cell factor 1 (splice form C) /		M15400	HUMRBS Human retinoblastoma susceptibility mR
U05237	Human fetal Alz-50-reactive clone 1 (FAC1) mRNA, compl		U11701	Human LIM-homeobox domain protein (hLH-2) mRNA, comple
U51003	Human DLX-2 (Dlx2) mRNA, complete cds /cds=UNKNOWN /gb		X07495	Human mRNA for cp19 homeobox from HOX-3 locus /cds=(60
L16464	HUMETSONC Human ETS oncogene (PEP1) mRNA, com		X14798	HSCETS1 Human DNA for c-ets-1 proto-oncogene
J03069	HUMMYCL2A Human MYCL2 gene, complete cds"		X86691	H.sapiens mRNA for 218kD Mi-2 protein /cds=(89,5827) /
X60655	H.sapiens EVX1 mRNA /cds=(228,1451) /gb=X60655 /gi=312		X97548	H.sapiens mRNA for TIF1beta zinc finger protein /cds=(
X14894	Human mRNA for myogenic factor Myf-5 /cds=(42,809) /gb		X52541	HSEGR1 Human mRNA for early growth response p
D89928	Homo sapiens HKL1 mRNA, complete cds /cds=(152,1969) /		U11732	HSU11732 Human ets-like gene (tel) mRNA, comp
L16499	Human orphan homeobox protein (PRH) mRNA, complete cds		L35269	Homo sapiens zinc finger protein 35 (ZNF35) gene /cds=
AF009353	Homo sapiens transcription intermediary factor 1 (TIF1		U16031	HSU16031 Human transcription factor IL-4 Stat
X53390	Human mRNA for upstream binding factor (hUBF) /cds=(14		U15306	Human cysteine-rich sequence-specific DNA-binding prot
M86737	Human high mobility group box (SSRP1) mRNA, complete c		X52560	Human gene for nuclear factor NF-IL6 /cds=(0,1037) /gb
U87309	Human hVps41p (HVPS41) mRNA, complete cds /cds=(29,259		U48730	HSU48730 Homo sapiens transcription factor St
M60618	Human nuclear autoantigen (SP-100) mRNA, complete cds		X17651	Human Myf-4 mRNA for myogenic determination factor /cd
M79462	HUMPML1 Human PML-1 mRNA, complete CDS"		Y09445	HSTBX5 H.sapiens mRNA for transcription facto
M79463	HUMPML2 Human PML-2 mRNA, complete CDS"		U76388	Human steroidogenic factor 1 mRNA, complete cds /cds=(
U31468	Homo sapiens homeobox protein (GBX2) gene, complete cd		L41870	HUMRB1MRNA Homo sapiens retinoblastoma suscep
AF042832	Homo sapiens forkhead-related transcription factor FRE		X82324	H.sapiens Brain 4 mRNA /cds=(32,1117) /gb=X82324 /gi=7
M22898	HUMP53A11 Human phosphoprotein p53 gene, exon		AF019415	untitled /cds=(105,926) /gb=AF019415 /gi=3955194 /
U51095	HSU51095 Human homeobox protein Cdx1 mRNA, co		M24900	Human triiodothyronine recptor (THRA1, ear1), and ear2
U51096	HSU51096 Human homeobox protein Cdx2 mRNA, co		U68385	Human Meis1-related protein 2 (MRG2), mRNA, partial cd
X55544	H.sapiens cDNA for TREB protein /cds=(0,815) /gb=X5554		K00650	HUMFOS Human fos proto-oncogene (c-fos), comp
AF001433	Human requiem (HREQ) mRNA, complete cds /cds=(41,1216)		U22662	HSU22662 Human nuclear orphan receptor LXR-al
U14755	Human LIM domain transcription factor LIM-1 (hLIM-1) m		D15050	Human mRNA for transcription factor AREB6, complete cd
J03827	Y box binding protein-1 (YB-1) mRNA /cds=(126,1079) /g		U00115	Human zinc-finger protein (bcl-6) mRNA, complete cds /
X13403	Human mRNA for octamer-binding protein Oct-1 /cds=(59,		M16937	Human homeo box c1 protein, mRNA, complete cds /cds=(9
S57212	hMEF2C=myocyte enhancer-binding factor 2 [human, skele		M16938	Human homeo box c8 protein, mRNA, complete cds /cds=(4
U34360	Human lymphoid nuclear protein (LAF-4) mRNA, complete		X02469	HSP53 Human mRNA for p53 cellular tumor antig
M97676	HUMHOX7 Homo sapiens (region 7) homeobox prot		U13948	Human zinc finger/leucine zipper protein (AF10) mRNA,
L03427	Human zinc finger protein basonuclin mRNA, complete cd		Z50781	H.sapiens mRNA for leucine zipper protein /cds=(135,36
AF001461	Homo sapiens Kruppel-like zinc finger protein Zf9 mRNA		M36542	Human lymphoid-specific transcription factor mRNA, com
L49169	Human G0S3 mRNA, complete cds /cds=(593,1609) /gb=L491		AF081280	Homo sapiens nucleoplasmin-3 (NPM3) mRNA, complete cds
X58431	HSHOX22 Human Hox2.2 gene for a homeobox prot		X92518	H.sapiens mRNA for HMGI-C protein /cds=UNKNOWN /gb=X92
D87073	Human mRNA for KIAA0236 gene, complete cds /cds=(436,5		S83308	SOX5=Sry-related HMG box gene {alternatively spliced}
M62810	Human mitochondrial transcription factor 1 mRNA, compl		AB006867	Homo sapiens mRNA for hSOX20 protein, complete cds /cd
X99894	HSGSFGENE H.sapiens mRNA coding for insulin p		AJ001699	Homo sapiens mRNA for Brachyury (T) protein /cds=(159,
U36500	Human lymphoid-specific SP100 homolog (LYSP100-B) mRNA		AF003540	Homo sapiens Krueppel family zinc finger protein (znfp
U36501	Human SP100-B (SP100-B) mRNA, complete cds /cds=(31,20		M27878	Human DNA binding protein (HPF2) mRNA, complete cds /c
AF076215	Homo sapiens prophet of Pit1 (PROP1) mRNA, complete cd		AF084199	Homo sapiens beta-interferon gene positive-regulatory
M96980	Homo sapiens myelin transcription factor 1 (MTF1) mRNA		U37251	Human KRAB zinc finger protein (ZNF177) mRNA, splicing
U77968	Human neuronal PAS1 (NPAS1) mRNA, complete cds /cds=(2		L12579	Human alternatively spliced CUTL1 mRNA, complete cds /
U90304	Human iroquois-class homeodomain protein IRX-2a mRNA,		U63809	Homo sapiens prostate apoptosis response protein par-4
M91196	Homo sapiens DNA-binding protein mRNA, complete cds /c		AF054180	Homo sapiens hematopoietic cell derived zinc finger pr

U77970	Human neuronal PAS2 (NPAS2) mRNA, complete cds /cds=(2		AB018287	Homo sapiens mRNA for KIAA0744 protein, complete cds /
M94046	Human zinc finger protein (MAZ) mRNA /cds=UNKNOWN /gb=		L04282	HUMTB Human CACCC box-binding protein mRNA, c
U43843	Human h-neuro-d4 protein mRNA, complete cds /cds=(6,10		AF038951	Homo sapiens DNA-binding protein mRNA, complete cds /c
X75918	H.sapiens mRNA for NOT /cds=(317,2113) /gb=X75918 /gi=		AF028008	Homo sapiens SP1-like zinc finger transcription factor
S74720	DAX-1=DSS-AHC critical region on X chromosome, gene 1		AF004849	Homo sapiens PKY protein kinase mRNA, complete cds /cd
X99101	HSRNAERB H.sapiens mRNA for estrogen receptor		U38896	Human zinc finger protein C2H2-171 mRNA, complete cds
X12794	HSEAR2 Human v-erbA related ear-2 gene		U41843	Human Dr1-associated corepressor (DRAP1) mRNA, complet
AF076292	Homo sapiens TGF-beta/activin signal transducer FAST-1		U53831	Homo sapiens interferon regulatory factor 7B mRNA, com
X69391	H.sapiens mRNA for ribosomal protein L6 /cds=(26,892)		D89859	Homo sapiens mRNA for zinc finger 5 protein, complete
U16997	Human orphan receptor ROR gamma mRNA, complete cds /cd		AB007947	Homo sapiens mRNA for KIAA0478 protein, complete cds /
M57732	Human hepatic nuclear factor 1 (TCF1) mRNA, complete c		AF041381	Homo sapiens putative transcriptional repressor E2F-6
V01512	HSCFOS Human cellular oncogene c-fos (complet		U93237	HSU93237 Human menin (MEN1) gene, complete cd
L25270	Human XE169 mRNA, complete cds /cds=(531,5213) /gb=L25		D43642	Human YL-1 mRNA for YL-1 protein (nuclear protein with
U10492	Human Mox1 protein (MOX1) mRNA, complete cds /cds=(29,		X99268	H.sapiens mRNA for B-HLH DNA binding protein /cds=(110
AF059575	untitled /cds=(0,1139) /gb=AF059575 /gi=3335523 /		U47677	HSE2F1S03 Human transcription factor E2F1 (E2
J03133	HUMTFSP1 Human transcription factor SP1 mRNA,		M96577	HUME2F Homo sapiens (E2F-1) pRB-binding prote
M87503	Human IFN-responsive transcription factor subunit mRNA		U27459	Human origin recognition complex protein 2 homolog hOR
AF017418	Homo sapiens homeobox protein MEIS2 (MEIS2) mRNA, part		L15409	HUMHIPLIND Homo sapiens (clone g7) von Hippel
AF104902	Homo sapiens ZIC2 protein (ZIC2) mRNA, complete cds /c		AB006626	Homo sapiens mRNA for KIAA0288 gene, complete cds /cds
U20648	Human zinc finger protein (ZNF154) mRNA, partial cds /		AF068754	Homo sapiens heat shock factor binding protein 1 HSBP1
AB005535	AB005535 Homo sapiens mRNA for Clock, partial		X52221	HSERCC25 H.sapiens ERCC2 gene, exons 1 & 2 (p
J03161	HUMSRF Human serum response factor (SRF) mRNA		AF054284	Homo sapiens spliceosomal protein SAP 155 mRNA, comple
L31881	Human nuclear factor I-X mRNA, complete cds /cds=(36,1		U61837	Homo sapiens putative cyclin G1 interacting protein mR
U51127	HSU51127 Human interferon regulatory factor 5		D80008	Human mRNA for KIAA0186 gene, complete cds /cds=(94,68
AF005887	AF005887 Homo sapiens ATF family member ATF6		D32143	Human mRNA for biliverdin-IXbeta reductase I /cds=(109
M95585	HUMHLF Human hepatic leukemia factor (HLF) mR		X12433	Human pHS1-2 mRNA with ORF homologous to membrane rece
X89750	H.sapiens mRNA for TGIF protein /cds=(311,1129) /gb=X8		AB011111	Homo sapiens mRNA for KIAA0539 protein, complete cds /
U35113	HSU35113 Human metastasis-associated mta1 mRN		U18919	Human chromosome 17q12-21 mRNA, clone pOV-2, partial c
X52773	HSRARLP Human mRNA for retinoic acid receptor		AF084523	Homo sapiens cellular repressor of E1A-stimulated gene
S74017	S74017 Nrf2=NF-E2-like basic leucine zipper t		AF074723	Homo sapiens RNA polymerase transcriptional regulation
M60721	HUMHB24 Human homeobox gene, complete cds"		AF016270	Homo sapiens thyroid hormone receptor coactivating pro
L46353	HUMHMGIC03 Homo sapiens high-mobility group p		U66615	HSU66615 Human SWI/SNF complex 155 KDa subuni
Y00291	Human hap mRNA encoding a DNA-binding hormone receptor		U66616	HSU66616 Human SWI/SNF complex 170 KDa subuni
U59831	Human transcription factor, forkhead related activator		U66617	HSU66617 Human SWI/SNF complex 60 KDa subunit
U43185	HSU43185 Human signal transducer and activato		U46691	Human putative chromatin structure regulator (SUPT6H)
U78722	Homo sapiens zinc finger protein 165 (Zpf165) mRNA, co		U66618	HSU66618 Human SWI/SNF complex 60 KDa subunit
U07559	Human ISL-1 (Islet-1) mRNA, complete cds /cds=(248,128		U66619	HSU66619 Human SWI/SNF complex 60 KDa subunit
X59244	Human ZNF43 mRNA /cds=(337,2748) /gb=X59244 /gi=38031		U03494	Human transcription factor LSF mRNA, complete cds /cds
D14520	Human mRNA for GC-Box binding protein BTEB2, complete		U03495	Human transcription factor LSF-ID mRNA, complete cds /
M83221	HUMIRELA Homo sapiens I-Rel mRNA, complete cd		AB006572	Homo sapiens RMP mRNA for RPB5 meidating protein, comp
X59268	Human mRNA for general transcription factor IIB /cds=(		L47345	HUMELONA Homo sapiens elongin A mRNA, complet
X52011	H.sapiens MYF6 gene encoding a muscle determination fa		U79751	Human basic-leucine zipper nuclear factor (JEM-1) mRNA
X86174	H.sapiens mRNA for SSX1 protein /cds=(91,657) /gb=X861		AB000468	Homo sapiens mRNA for zinc finger protein, complete cd
X74594	HSRB2P130 H.sapiens mRNA for Rb2/p130 protein		D50495	HUMTEF Homo sapiens mRNA for transcription el
M55654	Human TATA-binding protein mRNA, complete cds /cds=(24		U63810	Homo sapiens WD40 protein Ciao 1 mRNA, complete cds /c
U48213	Human D-site binding protein gene, promoter region and		L40904	H. sapiens peroxisome proliferator activated receptor
U43923	HSU43923 Human transcription factor SUPT4H mR		AF069733	Homo sapiens ADA3-like protein mRNA, complete cds /cds
AF048722	Homo sapiens ALL1 responsive protein ARP1c (ARP1) mRNA		AF069735	Homo sapiens PCAF associated factor 65 alpha mRNA, com
X69438	H.sapiens zinc finger gene pAT133 /cds=(231,1691) /gb=		U69609	Human transcriptional repressor (GCF2) mRNA, complete
X52056	HSSPI1 Human mRNA for spi-1 proto-oncogene		AB006651	Homo sapiens EXLM1 mRNA, complete cds /cds=(119,4483)
U52682	Human lymphocyte specific interferon regulatory factor		U85658	Human transcription factor ERF-1 mRNA, complete cds /c
M21535	HUMERG11 Human erg protein (ets-related gene)		U60666	Human testis specific leucine rich repeat protein (TSL
AF016052	Homo sapiens zinc finger protein ZNF191 (ZNF191) gene,		AB011076	Homo sapiens mRNA for UTF1, complete cds /cds=(15,1040
X51345	Human jun-B mRNA for JUN-B protein /cds=(253,1296) /gb		U13045	Human nuclear respiratory factor-2 subunit beta 1 mRNA
U48250	HSU48250 Human protein kinase C-binding prote		AF031383	Homo sapiens hMed7 (MED7) mRNA, complete cds /cds=(66,
L13740	HUMTR3A Human TR3 orphan receptor mRNA, compl		U91543	Homo sapiens zinc-finger helicase (hZFH) mRNA, complet
Y08223	H.sapiens MFH-1 gene /cds=(0,1505) /gb=Y08223 /gi=1869		U38847	Human TAR RNA loop binding protein (TRP-185) mRNA, com
D43638	Human mRNA for MTG8a protein, complete cds /cds=(411,2		U03644	Human recepin mRNA, complete cds /cds=(32,1387) /gb=U0
D89377	Homo sapiens mRNA for MSX-2, complete cds /cds=(61,864		AB021663	Homo sapiens mRNA for leucine-zipper protein, complete
D10216	Human mRNA for Pit-1/GHF-1, complete cds /cds=(125,100		U78082	Human RNA polymerase transcriptional regulation mediat
X96381	H.sapiens erm gene, exon 2,3,4,5 (and joined CDS) /cds		AF012108	Homo sapiens Amplified in Breast Cancer (AIB1) mRNA, c
U28687	Human zinc finger containing protein ZNF157 (ZNF157) m		U08336	Human basic helix-loop-helix transcription factor mRNA
L13773	Human AF-4 mRNA, complete cds /cds=(420,4052) /gb=L137		U58198	Human interleukin enhancer binding factor 3 mRNA, comp
M84739	Human autoantigen calreticulin mRNA, complete cds /cds		AB010882	Homo sapiens mRNA for hSNF2H, complete cds /cds=(202,3
L10102	Homo sapiens sex-determining region Y (SRY) gene, comp		AB021742	Homo sapiens NDRF gene for neuroD-related factor, comp
AJ001183	Homo sapiens mRNA for Sox10 protein /cds=(120,1520) /g		U02683	Human alpha palindromic binding protein mRNA, complete
M76766	HUMTFIIB Human transcription factor (TFIIB) m		L34587	HUMRPIE Homo sapiens RNA polymerase II elonga
J03258	HUMVDR Human vitamin D receptor mRNA, complet		U04847	Human Ini1 mRNA, complete cds /cds=(69,1226) /gb=U0484
L07919	Human homeodomain protein DLX-2 mRNA, 3 end /cds=(0,7		M81601	HUMTEFSII Human transcription elongation fact
S77763	nuclear factor erythroid 2 isoform f=basic leucine zip		M31523	HUMTFAA Human transcription factor (E2A) mRNA
Z21966	HSMPOUHOX H.sapiens mPOU homeobox protein mRN		L08424	Homo sapiens achaete scute homologous protein (ASH1) m
U43203	Human thyroid transcription factor 1 (TTF-1) mRNA, com		X52611	HSAP2 Human mRNA for transcription factor AP-
D79998	Human mRNA for KIAA0176 gene, partial cds /cds=(0,797)		U00968	Human SREBP-1 mRNA, complete cds /cds=(166,3609) /gb=U
U12767	HSU12767 Human mitogen induced nuclear orphan		X64229	H.sapiens dek mRNA /cds=(33,1160) /gb=X64229 /gi=30502
AJ002607	Homo sapiens HOX11L1 gene, exon 1 and joined CDS /cds=		U11861	Human G10 homolog (edg-2) mRNA, complete cds /cds=(379
Y09723	H.sapiens mRNA for Miz-1 protein /cds=(126,2537) /gb=Y		M96954	Homo sapiens nucleolysin TIAR mRNA, complete cds /cds=
AF032885	Homo sapiens forkhead protein (FKHR) mRNA, complete cd		AF006513	Homo sapiens CHD1 mRNA, complete cds /cds=(163,5292) /
AF032886	Homo sapiens forkhead protein (FKHRL1) mRNA, complete		AF006514	Homo sapiens CHD2 mRNA, complete cds /cds=(707,5926) /
X68011	H.sapiens ZNF81 gene /cds=(0,1013) /gb=X68011 /gi=4543		U08998	Human TAR RNA binding protein 2 (TRBP2) mRNA, complete
L27586	Human TR4 orphan receptor mRNA, complete cds /cds=(222		U84570	Human A2 mRNA, complete cds /cds=(239,883) /gb=U84570
X63741	H.sapiens pilot mRNA /cds=(353,1516) /gb=X63741 /gi=35		M36711	Human sequence-specific DNA-binding protein (AP-2) mRN

X76061	H.sapiens p130 mRNA for 130K protein /cds=(69,3488) /g		U96915	Homo sapiens sin3 associated polypeptide p18 (SAP18) m
X82629	H.sapiens mRNA for Mox-2 /cds=(265,1176) /gb=X82629 /g		D26155	Human mRNA for transcriptional activator hSNF2a, compl
U41813	Human class I homeoprotein (HOXA9) mRNA, partial cds /		L23959	HUMDP1A Homo sapiens E2F-related transcriptio
U15655	HSU15655 Human ets domain protein ERF mRNA, c		AF045184	Homo sapiens nuclear receptor coactivator NCoA-62 mRNA
M38258	HUMRARGA Human retinoic acid receptor gamma 1		AF010312	Homo sapiens Pig7 (PIG7) mRNA, complete cds /cds=(79,7
M26679	HUMHOX13G Homo sapiens homeobox protein (HOX-		X89887	Homo sapiens mRNA for WD repeat protein (HIRA) /cds=(2
X68060	H.sapiens topIIb mRNA for topoisomerase IIb /cds=(0,48		X07024	HSCCG1 Human X chromsome mRNA for CCG1 protei
X76091	H.sapiens HRFX2 mRNA /cds=(159,2330) /gb=X76091 /gi=45		U65092	Human melanocyte-specific gene 1 (msg1) mRNA, complete
X76092	H.sapiens HRFX3 mRNA /cds=(8,2131) /gb=X76092 /gi=4524		U65093	Human msg1-related gene 1 (mrg1) mRNA, complete cds /c
L38487	HUMHERRA1 Human estrogen receptor-related pro		Y11392	Homo sapiens mRNA; candidate gene for APECED /cds=(68,
U07664	HSHB9HB2 Human HB9 homeobox gene, exons 2 and		M74719	Human SEF2-1B protein (SEF2-1B) mRNA, complete cds /cd
L05072	HUMIFNRF1A Homo sapiens interferon regulatory		M63896	Homo sapiens transcriptional enhancer factor (TEF1) DN
Z46629	Homo sapiens SOX9 mRNA /cds=(359,1888) /gb=Z46629 /gi=		M33336	HUMCAMPPK Human cAMP-dependent protein kinase
AF005220	Homo sapiens transcription factor HOXD13 (Hoxd13) gene		X87838	H.sapiens mRNA for beta-catenin /cds=(214,2559) /gb=X8
U18543	Human zinc-finger protein mRNA, complete cds /cds=(110		X87843	HSCYCHASS H.sapiens mRNA for cyclin H assembl
U57029	Human T-cell leukemia virus enhancer factor (HTLF) mRN		D90359	Human CCG1 mRNA /cds=(51,5669) /gb=D90359 /gi=559319 /
X59372	HSHOX4C Human HOX4C mRNA for a homeobox prote		D13317	Human mRNA for transcription factor, E4TF1-53, complet
X59373	HSHOX4D Human HOX4D mRNA for a homeobox prote		M80627	Human HEB helix-loop-helix protein (HEB) mRNA, complet
U37431	HSU37431 Human HOXA1 mRNA, long transcript an		U52960	Human RNA polymerase II complex component SRB7 mRNA, c
Y13436	Homo sapiens sox1 gene /cds=(0,1163) /gb=Y13436 /gi=41		U63824	Human transcription factor RTEF-1 (RTEF1) mRNA, comple
M15024	HUMCMYBLA Human c-myb mRNA, complete cds"		S78825	Id1 (Id1-b)=transcription regulator helix-loop-helix p
L11672	HUMKRUPZN Human Kruppel related zinc finger p		X72889	H.sapiens hbrm mRNA /cds=(222,4982) /gb=X72889 /gi=414
U49020	Human myocyte-specific enhancer factor 2A (MEF2A) gene		S73885	AP-4=basic helix-loop-helix DNA-binding protein [human
M29580	Human zinc-finger protein 7 (ZFP7) mRNA, complete cds		X77956	H.sapiens Id1 mRNA /cds=(35,499) /gb=X77956 /gi=457784
M29581	Human zinc-finger protein 8 (ZFP8) mRNA, 3 end /cds=(		U14680	HSU14680 Human breast and ovarian cancer susc
M64497	Human apolipoprotein AI regulatory protein (ARP-1) mRN		U29175	Human transcriptional activator (BRG1) mRNA, complete
L02932	HUMPPAR Human peroxisome proliferator activat		X83928	HSTAFII28 H.sapiens mRNA for transcription fa
U57052	Human Hoxb-13 mRNA, complete cds /cds=(54,908) /gb=U57		U75308	HSU75308 Human TBP-associated factor (hTAFII1
X51417	HSSTHOR2 Human mRNA for steroid hormone recep		Y09321	H.sapiens TAFII105 mRNA, partial /cds=(0,2405) /gb=Y09
U09848	Human zinc finger protein (ZNF139) mRNA, partial cds /		U13991	HSU13991 Human TATA-binding protein associate
D78579	Homo sapiens mRNA for neuron derived orphan receptor,		U72209	HSU72209 Human YY1-associated factor 2 (YAF2)
U09850	Human zinc finger protein (ZNF143) mRNA, complete cds		AF013956	Homo sapiens Polycomb 2 homolog (hPc2) mRNA, complete
AF000993	Homo sapiens ubiquitous TPR motif, X isoform (UTX) mRN		Z14000	H.sapiens RING1 gene /cds=(75,1208) /gb=Z14000 /gi=296
AF000994	Homo sapiens ubiquitous TPR motif, Y isoform (UTY) mRN		AB014591	Homo sapiens mRNA for KIAA0691 protein, complete cds /
X95701	H.sapiens mRNA for GATA-6 DNA-binding protein /cds=(71		U71267	Human potential transcriptional repressor NOT4Hp (NOT4
X51435	Human PRDII-BF1 gene for a DNA-binding protein /cds=(3		U07563	HSABLGR3 Human proto-oncogene tyrosine-protei
X17254	Human mRNA for the transcription factor Eryf1 /cds=(11		X16416	Human c-abl mRNA encoding p150 protein /cds=(147,3539)
D16815	Homo sapiens mRNA for EAR-1r, complete cds /cds=(305,2		M14752	HUMABLA Human c-abl gene, complete cds"
X55005	HSCERBAR Homo sapiens mRNA for thyroid hormon		U90919	Human clones 23667 and 23775 zinc finger protein mRNA,
AB002803	Homo sapiens BACH1 mRNA, complete cds /cds=(118,2328)		U93867	HSU93867 Human RNA polymerase III subunit (RP
L78440	HUMSTAT4R Homo sapiens STAT4 mRNA, complete c		U93868	Human RNA polymerase III subunit (RPC32) mRNA, complet
AB015132	Homo sapiens UKLF mRNA for ubiquitous Kruppel like fac		U93869	Human RNA polymerase III subunit (RPC39) mRNA, complet
M84820	HUMRXRB Human retinoid X receptor beta (RXR-b		D88152	Homo sapiens mRNA for acetyl-coenzyme A transporter, c
U38904	Human zinc finger protein C2H2-25 mRNA, complete cds /		AL050274	Homo sapiens mRNA; cDNA DKFZp566C243 (from clone DKFZp
X16576	Human KUP mRNA for protein with two zinc fingers /cds=		AF038961	Homo sapiens SL15 protein mRNA, complete cds /cds=(16,
X68879	H.sapiens EMX1 mRNA /cds=(0,461) /gb=X68879 /gi=31139		AF004876	Homo sapiens 54TMp (54tm) mRNA, complete cds /cds=(115
M13666	Human c-myb mRNA, 3 end /cds=(0,833) /gb=M13666 /gi=1		AL080122	Homo sapiens mRNA; cDNA DKFZp564O123 (from clone DKFZp
X68880	H.sapiens EMX2 mRNA /cds=(0,476) /gb=X68880 /gi=31141		AF043724	Homo sapiens hepatitis A virus cellular receptor 1 (hH
U06233	Human POU domain protein (Brn-3b) mRNA, complete cds /		AL080155	Homo sapiens mRNA; cDNA DKFZp434J154 (from clone DKFZp
M25269	HUMELK1A Homo sapiens tyrosine kinase (ELK1)		U41816	Human C-1 mRNA, complete cds /cds=(11,403) /gb=U41816
X99350	H.sapiens HFH4 gene, exon 1 and joined CDS /cds=(216,1		AF042384	Homo sapiens BC-2 protein mRNA, complete cds /cds=(129
X15875	Human mRNA for cAMP response element (CRE-BP1) binding		U03886	Human GS2 mRNA, complete cds /cds=(130,891) /gb=U03886
Z11898	H.sapiens OTF3 mRNA encoding octamer binding protein 3		Y12670	Homo sapiens mRNA for leptin receptor gene-related pro
M17254	Human erg2 gene encoding erg2 protein, complete cds /c		D14659	Human mRNA for KIAA0103 gene, complete cds /cds=(6,899
U66561	Human kruppel-related zinc finger protein (ZNF184) mRN		D63477	Human mRNA for KIAA0143 gene, partial cds /cds=(0,2658
X70394	H.sapiens OZF mRNA /cds=(856,1734) /gb=X70394 /gi=4687		AB018257	Homo sapiens mRNA for KIAA0714 protein, partial cds /c
Y07661	H.sapiens USF2 gene /cds=(0,1040) /gb=Y07661 /gi=18060		AF063605	Homo sapiens brain my047 protein mRNA, complete cds /c
J04076	Human early growth response 2 protein (EGR2) mRNA, com		D31815	HUMSMP30 Human mRNA for SMP-30 (senescence ma
J04088	HUMTOPII Human DNA topoisomerase II (top2) mR		U73524	Human putative ATP/GTP-binding protein (HEAB) mRNA, co
X65233	H.sapiens mRNA for Zinc-finger protein (ZNFpT17) /cds=		AL050369	Homo sapiens mRNA; cDNA DKFZp566J153 (from clone DKFZp
AF040963	Homo sapiens Mad4 homolog (Mad4) mRNA, complete cds /c		U59151	Human Cbf5p homolog (CBF5) mRNA, complete cds /cds=(80
U40462	HSU40462 Human Ikaros/LyF-1 homolog (hIk-1) m		D80007	Human mRNA for KIAA0185 gene, partial cds /cds=(0,5655
M68864	Human ORF mRNA, complete cds /cds=(135,1031) /gb=M6886		Y12065	Homo sapiens mRNA for nucleolar protein hNop56 /cds=(2
D85131	D85131 Homo sapiens mRNA for Myc-associated z		X56597	Human humFib mRNA for fibrillarin /cds=(59,1024) /gb=X
X52943	Human mRNA for ATF-a transcription factor /cds=(103,15		M58460	Human 75-kD autoantigen (PM-Sc1) mRNA, complete cds /c
L16794	Human transcription factor (MEF2) mRNA, complete cds /		D29641	Human mRNA for KIAA0052 gene, partial cds /cds=(0,2510
U68723	HSU68723 Human checkpoint suppressor 1 mRNA,		X98494	H.sapiens mRNA for M phase phosphoprotein 10 /cds=(0,2
U68727	HSU68727 Human homeobox-containing protein mR		U28042	HSU28042 Human DEAD box RNA helicase-like pro
U70212	Human SIM1 mRNA, complete cds /cds=(216,2516) /gb=U702		U02619	Human TFIIIC Box B-binding subunit mRNA, complete cds
U17163	Human transcription factor ETV1 mRNA, complete cds /cd		L05515	HUMCREBPA Homo sapiens cAMP response element-
U44059	Human thyrotroph embryonic factor (TEF) mRNA, complete		M85234	Human nuclease sensitive element binding protein-1 mRN
M68891	HUMGATA Human GATA-binding protein (GATA2) mR		U68019	HSU68019 Homo sapiens mad protein homolog (hM
AF000297	Homo sapiens Human nkx-2.8 homeobox gene~~Exon 1 and i		AF017307	Homo sapiens Ets-related transcription factor (ERT) mR
D76435	Homo sapiens mRNA for Zic protein, complete cds /cds=(		AF044209	Homo sapiens nuclear receptor co-repressor N-CoR mRNA,
M65217	HUMHSF2 Human heat shock factor 2 (HSF2) mRNA		L32832	Homo sapiens zinc finger homeodomain protein (ATBF1-A)
AB007901	Homo sapiens KIAA0441 mRNA, complete cds /cds=(168,226		U65928	HSU65928 Human Jun activation domain binding
U23752	Human SOX-11 mRNA, complete cds /cds=(75,1400) /gb=U23		D14041	Homo sapiens mRNA for H-2K binding factor-2, complete
U50648	HSIIPKR17 Human interferon-inducible RNA-depe		AF072242	Homo sapiens methyl-CpG binding protein MBD2 (MBD2) mR
U28049	Human TBX2 (TXB2) mRNA, complete cds /cds=(47,2155) /g		AB007296	Homo sapiens mRNA for hGLI2, complete cds, clone-hGli2
U04897	Human orphan hormone nuclear receptor RORalpha1 mRNA,		U59736	Human transcription factor (NFATc.b) mRNA, complete cd
L20433	Human octamer binding transcription factor 1 (OTF1) mR		AB006909	Homo sapiens mRNA for A-type microphthalmia associated

S54641	HZF-16=Kruppel-related zinc finger gene homolog {alter		U08191	Human R kappa B mRNA, complete cds /cds=(2220,5216) /g
M97935	Homo sapiens transcription factor ISGF-3 mRNA, complet		AF017789	Homo sapiens putative transcription factor CA150 mRNA,
M97936	Human transcription factor ISGF-3 mRNA sequence /cds=U		AF084260	Homo sapiens signalosome subunit 2 (SGN2) mRNA, comple
D87328	Homo sapiens mRNA for HCS, complete cds /cds=(1231,341		AF049703	Homo sapiens E74-like factor 5 (ELF5) mRNA, complete c
U35376	Human repressor transcriptional factor (ZNF85) mRNA, c		AF032387	Homo sapiens snRNA activating protein complex 190kD su
D83778	Human mRNA for KIAA0194 gene, partial cds /cds=(0,4309		AL050265	Homo sapiens mRNA; cDNA DKFZp564O1716 (from clone DKFZ
L15309	Human zinc finger protein (ZNF141) mRNA, complete cds		U96131	Homo sapiens HPV16 E1 protein binding protein mRNA, co
U80456	Human transcription factor SIM2 long form mRNA, comple		AB022663	Homo sapiens HFB30 mRNA, complete cds /cds=(236,1660)
U80457	Human transcription factor SIM2 short form mRNA, compl		L10403	Homo sapiens DNA binding protein for surfactant protei
U18671	HSU18671 Human Stat2 gene, complete cds"		U32645	Human myeloid elf-1 like factor (MEF) mRNA, complete c
L07335	Homo sapiens (clone 6AR33) HMG box mRNA, 3 end cds /c		U88528	Human transcription factor LZIP mRNA, complete cds /cd
AJ222700	Homo sapiens mRNA for TSC-22 protein /cds=(191,625) /g		D88208	Homo sapiens hSGT1 mRNA for hSgt1p, complete cds /cds=
X71135	H.sapiens sox3 gene /cds=(0,1331) /gb=X71135 /gi=46879		L24203	HUMDK Homo sapiens ataxia-telangiectasia grou
U48436	Homo sapiens fragile X mental retardation protein FMR2		AB012124	Homo sapiens TCFL5 mRNA for transcription factor-like
X60287	HSMAXM H.sapiens max mRNA		AF048730	Homo sapiens cyclin T1 mRNA, complete cds /cds=(44,222
X17360	HSHOX51 Human HOX 5.1 gene for HOX 5.1 protei		AF048731	Homo sapiens cyclin T2a mRNA, complete cds /cds=(32,20
D88827	Homo sapiens mRNA for zinc finger protein FPM315, comp		AF049460	Homo sapiens nuclear DEAF-1 related transcriptional re
X63131	H.sapiens My1 (PML) mRNA /cds=(141,2042) /gb=X63131 /g		U05040	Human FUSE binding protein mRNA, complete cds /cds=(26
U74612	Human hepatocyte nuclear factor-3/fork head homolog 11		AF035309	Homo sapiens clone 23598 mRNA, complete cds /cds=(880,
Z11933	H.sapiens mRNA for N-Oct 3, N-Oct5a, and N-Oct 5b prot		U91935	Human retina-derived POU-domain factor-1 mRNA, alterna
S79325	SYT...SSX1 {translocation breakpoint} [human, synovial		AF064094	Homo sapiens KL04P mRNA, complete cds /cds=(108,1439)
M93650	Human paired box gene (PAX6) homologue, complete cds /		U17838	Human zinc finger protein RIZ mRNA, complete cds /cds=
J04101	HUMETS1A Human erythroblastosis virus oncogen		D45132	HUMHOXY1 Homo sapiens mRNA for zinc-finger DN
J04102	HUMETS2A Human erythroblastosis virus oncogen		AB010812	Homo sapiens Nrf3 mRNA for NF-E2-related factor 3, com
AF046001	Homo sapiens zinc finger transcription factor (ZNF207)		AB020634	Homo sapiens mRNA for KIAA0827 protein, complete cds /
U47741	Human CREB-binding protein (CBP) mRNA, complete cds /c		AB028979	Homo sapiens mRNA for KIAA1056 protein, complete cds /
U47742	HSU47742 Human monocytic leukaemia zinc finge		AF010403	Homo sapiens ALR mRNA, complete cds /cds=(0,15788) /gb
Y07707	H.sapiens mRNA for ITBA4 gene /cds=(0,716) /gb=Y07707		AF081287	Homo sapiens serine phosphatase FCP1a (FCP1) mRNA, com
X16663	Human HS1 gene for heamatopoietic lineage cell specifi		U32849	Homo sapiens Nmi mRNA, complete cds /cds=(280,1203) /g
X16665	Human HOX2H mRNA from the Hox2 locus /cds=(78,1148) /g		U37146	Human silencing mediator of retinoid and thyroid hormo
J04111	HUMJUNA Human c-jun proto oncogene (JUN), com		M80261	HUMAPE Human apurinic endonuclease (APE) mRNA
X16666	Human HOX2I mRNA from the Hox2 locus /cds=(6,911) /gb=		M96944	Human B-cell specific transcription factor (BSAP) mRNA
X16667	Human HOX2G mRNA from the Hox2 locus /cds=(363,1658) /		AJ238381	Homo sapiens pax9 gene, exons 1-2 and joined CDS /cds=
X15949	HSIRF2 Human mRNA for interferon regulatory f		L20298	Homo sapiens transcription factor (CBFB) mRNA, 3 end
U67369	Human growth factor independence-1 (Gfi-1) mRNA, compl		L13744	Human AF-9 mRNA, complete cds /cds=(195,1901) /gb=L137
X68985	HSHEPLF H.sapiens mRNA for hepatic leukemia f		U01877	HSU01877 Human p300 protein mRNA, complete cd
D82344	Homo sapiens mRNA for NBPhox, complete cds /cds=(360,1		X96401	H.sapiens mRNA for ROX protein /cds=(212,1960) /gb=X96
U22376	HSU22376 Human (c-myb) gene, complete primary		U44754	Human PSE-binding factor PTF gamma subunit mRNA, compl
X87870	H.sapiens mRNA for hepatocyte nuclear factor 4a /cds=(		U44755	Human PSE-binding factor PTF delta subunit mRNA, compl
X87871	H.sapiens mRNA for hepatocyte nuclear factor 4b /cds=(		AB007131	Homo sapiens mRNA for HSF2BP, complete cds /cds=(332,1
D28588	Human mRNA for KIAA0048 gene, complete cds /cds=(334,2		Y12812	H.sapiens RFXAP mRNA /cds=(0,818) /gb=Y12812 /gi=20734
AJ000644	Homo sapiens mRNA for SPOP /cds=(157,1281) /gb=AJ00064		AF094760	Homo sapiens RFXANK (RFXANK) mRNA, complete cds /cds=(
M97287	Human MAR/SAR DNA binding protein (SATB1) mRNA, comple		U74667	HSU74667 Human tat interactive protein (TIP60
X61755	HSHOX3D Human HOX3D gene for homeoprotein HOX		M95809	Human basic transcription factor 62kD subunit (BTF2),
U81984	Human endothelial PAS domain protein 1 (EPAS1) mRNA, c		J03473	HUMRISDAD Human poly(ADP-ribose) synthetase m
M77698	HUMKRP Homo sapiens GLI-Krupple related prote		X15217	HSSNOAR Human sno oncogene mRNA for snoA prot
U31814	Human transcriptional regulator homolog RPD3 mRNA, com		X15218	HSSKIR Human ski oncogene mRNA
X72631	H.sapiens mRNA encoding Rev-ErbAalpha /cds=UNKNOWN /gb		D14887	Human mRNA for TFIIA-42, complete cds /cds=(404,1534)
X66079	H.sapiens Spi-B mRNA /cds=(5,793) /gb=X66079 /gi=36562		S66427	S66427 RBP1=retinoblastoma binding protein 1
X64624	H.sapiens mRNA for RDC-1 POU domain containing protein		X96717	H.sapiens mRNA for transcription factor TFE3 /cds=(234
X66087	HSAMYB2 H.sapiens a-myb mRNA /NOTE=replacemen		L22846	HUME2F2A Homo sapiens transcription factor E2
L16896	Human zinc finger protein mRNA, complete cds /cds=(155		X64037	H.sapiens mRNA for RNA polymerase II associated protei
U13695	HSU13695 Human homolog of yeast mutL (hPMS1)		U21858	HSU21858 Human transcriptional activation fac
X58072	Human hGATA3 mRNA for trans-acting T-cell specific tra		M37197	HUMCBF Human CCAAT-box-binding factor (CBF) m
U05681	HSBCL3S2 Human proto-oncogene (BCL3) gene, ex		X13293	HSBMYB Human mRNA for B-myb gene
L13203	Human HNF-3/fork-head homolog-3 HFH-3 mRNA, complete c		Z49194	H.sapiens mRNA for oct-binding factor /cds=(523,1293)
L08895	Homo sapiens MADS/MEF2-family transcription factor (ME		L40371	Homo sapiens thyroid receptor interactor (TRIP4) mRNA,
X90828	H.sapiens mRNA for transcription factor, Lbx1 /cds=(14		L40388	HUMTRIP15M Homo sapiens thyroid receptor inte
D32257	Human GTF3A mRNA for Xenopus transcription factor IIIA		L04791	HUMERCC6A Human excision repair protein ERCC6
D50405	Human mRNA for RPD3 protein, complete cds /cds=(63,151		X75042	HSRNAREL H.sapiens rel proto-oncogene mRNA
X03635	HSERR Human mRNA for oestrogen receptor		U71300	Human snRNA activating protein complex 50kD subunit (S
U70370	Human hindlimb expressed homeobox protein backfoot (Bf		M90354	Human BTF3 protein homologue gene, complete cds /cds=(
M74099	Human displacement protein (CCAAT) mRNA /cds=UNKNOWN /		M90355	Human BTF3 protein homologue gene, complete cds /cds=(
X56681	HSJUNDR Human junD mRNA		M90356	Human BTF3 protein homologue gene, complete cds /cds=(
Y09980	H.sapiens HOXD3 gene /cds=(378,1676) /gb=Y09980 /gi=18		M90357	Human basic transcription factor 3a (BTF3a) gene /cds=
L04510	Human nucleotide binding protein mRNA, complete cds /c		X80230	HSSTPKC2K H.sapiens mRNA (clone C-2k) mRNA fo
X56687	H.sapiens mRNA for autoantigen NOR-90 /cds=(179,2362)		U16282	Human ELL mRNA, complete cds /cds=(12,1877) /gb=U16282
X82850	HSTTF1 H.sapiens mRNA for thyroid transcript		X16901	Human mRNA for RAP30 subunit of transcription initiati
Z56281	HSIRF3MR H.sapiens mRNA for interferon regula		AF015553	Homo sapiens TFII-I protein (TFII-I) mRNA, complete cd
S38742	S38742 HOX11=HOX11 homeodomain {homeobox} [hu		AF026445	Homo sapiens cofactor of initiator function (CIF150) m
D14822	Human chimeric mRNA derived from AML1 gene and MTG8(ET		U77948	HSU77948 Human Bruton s tyrosine kinase-assoc
D14823	Human chimeric mRNA derived from AML1 gene and MTG8(ET		X95525	H.sapiens mRNA for TAFII100 protein /cds=(23,2422) /gb
D14827	HUMTHP1 Human mRNA for Tax helper protein 1,		U57693	Human TFIID subunits TAF20 and TAF15 mRNA, complete cd
M55914	HUMCMYCQ Human c-myc binding protein (MBP-1)		X84002	HSTAFII20 H.sapiens TAFII20 mRNA for transcri
M64673	HUMHSF1 Human heat shock factor 1 (TCF5) mRNA		X84003	HSTAFII18 H.sapiens TAFII18 mRNA for transcri
M92299	Human homeobox 2.1 protein (HOX2A) mRNA, complete cds		U14193	HSU14193 Human TFIIA gamma subunit mRNA, comp
X58840	Human mRNA for variant hepatic nuclear factor 1 (vHNF1		D38550	Human mRNA for KIAA0075 gene, partial cds /cds=(0,524)
X66867	HSMAXG H.sapiens max gene		L20320	Human protein serine/threonine kinase stk1 mRNA, compl
U20240	Human C/EBP gamma mRNA, complete cds /cds=(250,702) /g		U75309	HSU75309 Human TBP-associated factor (hTAFII1
AB002296	Human mRNA for KIAA0298 gene, complete cds /cds=(2865,		X77743	HSCDKAK H.sapiens CDK activating kinase mRNA
L35035	Homo sapiens ribose 5-phosphate isomerase (RPI) mRNA /		U18062	HSU18062 Human TFIID subunit TAFII55 (TAFII55

M91585	Human Br140 mRNA, complete cds /cds=(28,3672) /gb=M915		X63468	H.sapiens mRNA for transcription factor TFIIE alpha /c
X07290	Human HF.12 gene mRNA /cds=(0,589) /gb=X07290 /gi=3207		X63469	H.sapiens mRNA for transcription factor TFIIE beta /cd
AF084645	Homo sapiens orphan nuclear receptor (PAR1) mRNA, comp		AB020661	Homo sapiens mRNA for KIAA0854 protein, complete cds /
M91592	Human zinc-finger protein (ZNF76) gene, partial cds /c		U90547	Human Ro/SSA ribonucleoprotein homolog (RoRet) mRNA, c
X52332	Homo sapiens mRNA for zinc finger protein 10 /cds=(70,		D21205	HUMERFP Human mRNA for estrogen responsive fi
AF078096	Homo sapiens forkhead/winged helix-like transcription		D13636	Human mRNA for KIAA0011 gene, complete cds /cds=(39,27
M23263	HUMARB Human androgen receptor mRNA, complete		M25077	Human SS-A/Ro ribonucleoprotein autoantigen 60 kd subu
X66899	HSEWS H.sapiens EWS mRNA		J04137	Human 60-kdal ribonucleoprotein (Ro) mRNA, complete cd
X16706	Human fra-2 mRNA /cds=(3,983) /gb=X16706 /gi=31464 /		U75276	HSU75276 Human TFIIB related factor hBRF (HBR
X16707	Human fra-1 mRNA /cds=(34,849) /gb=X16707 /gi=31462 /		AB018281	Homo sapiens mRNA for KIAA0738 protein, complete cds /
U49250	Human putative cerebral cortex transcriptional regulat		L25444	Homo sapiens (TAFII70-alpha) mRNA, complete cds /cds=(
X51630	HSWT1 Human Wilms tumor WT1 mRNA for zinc fin		AF073771	Homo sapiens RNA polymerase II termination factor mRNA
L34357	Homo sapiens GATA-4 mRNA, complete cds /cds=(240,1568)			
M82882	Human cis-acting sequence /cds=UNKNOWN /gb=M82882 /gi=			
U89995	Human DNA binding protein FKHL15 (FKHL15) mRNA, comple			
U04270	Human putative potassium channel subunit (h-erg) mRNA,			
AF027219	Homo sapiens ZNF202 beta (ZNF202) mRNA, complete cds /			
D89667	D89667 Homo sapiens mRNA for c-myc binding pr			
U82759	Human homeodomain protein HoxA9 mRNA, complete cds /cd			

Supplementary Table 8: Affymetrix U95Av2 Accessions with "Biosynthesis" Gene Ontology Codes

These genes were classified as belonging to the process "biosynthesis" according to the Computen Gene Ontology resource.

Accession	Gene Description		Accession	Gene Description
D78335	Human mRNA for 5-terminal region of UMK, complete cds		D50929	Human mRNA for KIAA0139 gene, complete cds /cds=(128,4
X54199	Human mRNA for GARS-AIRS-GART /cds=UNKNOWN /gb=X54199		M84711	HUMFTE1A Human v-fos transformation effector
L28957	Homo sapiens CTP-phosphocholine cytidyltransferase mRN		L06498	Homo sapiens ribosomal protein S20 (RPS20) mRNA, compl
Y09022	H.sapiens mRNA for Not56-like protein /cds=(31,1347) /		L06499	Homo sapiens ribosomal protein L37a (RPL37A) mRNA, com
M11058	Human 3-hydroxy-3-methylglutaryl coenzyme A reductase		D32050	Human mRNA for alanyl-tRNA synthetase, complete cds /c
X69141	H.sapiens mRNA for squalene synthase /cds=(91,1344) /g		Z28407	H.sapiens mRNA for ribosomal protein L8 /cds=(43,816)
D11466	Homo sapiens mRNA for PIG-A protein, complete cds /cds		X55715	Human Hums3 mRNA for 40S ribosomal protein s3 /cds=(22
X78283	H.sapiens mRNA for aryl sulfotransferase (ST1A3) /cds=		D32053	Homo sapiens mRNA for Lysyl tRNA Synthetase, complete
M31642	Human hypoxanthine phosphoribosyltransferase (HPRT) mR		U94855	Homo sapiens translation initiation factor 3 47 kDa su
M27396	Human asparagine synthetase mRNA, complete cds /cds=(1		U14966	Human ribosomal protein L5 mRNA, complete cds /cds=(30
U61263	Human acetolactate synthase homolog mRNA, complete cds		U14968	Human ribosomal protein L27a mRNA, complete cds /cds=(
U68418	Human branched chain aminotransferase precursor (BCATm		U14969	Human ribosomal protein L28 mRNA, complete cds /cds=(2
U21551	Human ECA39 mRNA, complete cds /cds=(0,1154) /gb=U2155		X06323	Human MRL3 mRNA for ribosomal protein L3 homologue (M
D78586	HUMMUPCAD Human CAD mRNA for multifunctional		U14970	Human ribosomal protein S5 mRNA, complete cds /cds=(37
X01630	Human mRNA for argininosuccinate synthetase /cds=(75,1		U14971	Human ribosomal protein S9 mRNA, complete cds /cds=(35
X59834	Human rearranged mRNA for glutamine synthase /cds=(109		U14972	Human ribosomal protein S10 mRNA, complete cds /cds=(1
M77836	Human pyrroline 5-carboxylate reductase mRNA, complete		X65923	H.sapiens fau mRNA /cds=(56,457) /gb=X65923 /gi=31302
J00139	HUMFOL5 Human dihydrofolate reductase gene, e		L05095	Homo sapiens ribosomal protein L30 mRNA, complete cds
J00140	Human dihydrofolate reductase gene /cds=(42,605) /gb=J		X17206	Human mRNA for LLRep3 /cds=(240,905) /gb=X17206 /gi=34
AF006043	Homo sapiens 3-phosphoglycerate dehydrogenase mRNA, co		D14660	Human mRNA for KIAA0104 gene, complete cds /cds=(34,87
Y10275	H.sapiens mRNA for L-3-phosphoserine phosphatase /cds=		U62962	Human Int-6 mRNA, complete cds /cds=(22,1359) /gb=U629
D88674	D88674 Homo sapiens mRNA for antizyme inhibit		X54326	H.sapiens mRNA for glutaminyl-tRNA synthetase /cds=(58
X16277	Human gene for ornithine decarboxylase ODC (EC 4.1.1.1		J02645	HUMEIF2A Human translational initiation facto
M33764	HUMSODB Human ornithine decarboxylase gene, c		L18960	HUMEIF4C Human protein synthesis factor (eIF-
D78361	HUMODAZ Human mRNA for ornithine decarboxylas		M94314	Homo sapiens ribosomal protein L30 mRNA, complete cds
AF057297	Homo sapiens ornithine decarboxylase antizyme 2 (OAZ2)		U09953	Human ribosomal protein L9 mRNA, complete cds /cds=(29
M21154	HUMAMD Human S-adenosylmethionine decarboxyla		X15940	Human mRNA for ribosomal protein L31 /cds=(7,384) /gb=
D79986	Human mRNA for KIAA0164 gene, complete cds /cds=(253,3		X79865	H.sapiens Mrp17 mRNA /cds=(137,733) /gb=X79865 /gi=131
U67171	Human selenoprotein W (selW) mRNA, complete cds /cds=(		X53777	Human L23 mRNA for putative ribosomal protein /cds=(13
L40410	Homo sapiens thyroid receptor interactor (TRIP3) mRNA,		Z25749	H.sapiens gene for ribosomal protein S7 /cds=(81,665)
L40411	Homo sapiens thyroid receptor interactor (TRIP8) mRNA,		M63180	Human threonyl-tRNA synthetase mRNA, complete cds /cds
U31525	Human glycogenin mRNA, complete cds /cds=(127,1128) /g		U04953	Human isoleucyl-tRNA synthetase mRNA, complete cds /cd
M55621	Human N-acetylglucosaminyltransferase I (GlcNAc-TI) mR		X55954	Human mRNA for HL23 ribosomal protein homologue /cds=(
AF061016	Homo sapiens UDP-glucose dehydrogenase (UGDH) mRNA, co		AF035280	Homo sapiens clone 23689 mRNA, complete cds /cds=(46,1
U67368	Human multiple exostosis 2 (EXT2) gene /cds=(30,2186)		X64707	H.sapiens BBC1 mRNA /cds=(51,686) /gb=X64707 /gi=29382
S79639	S79639 EXT1=putative tumour suppressor/heredi		AB018284	Homo sapiens mRNA for KIAA0741 protein, complete cds /
X92720	H.sapiens mRNA for phosphoenolpyruvate carboxykinase /		AF022229	Homo sapiens translation initiation factor 6 (eIF6) mR
V00572	Human mRNA encoding phosphoglycerate kinase /cds=(79,1		M64716	Human ribosomal protein S25 mRNA, complete cds /cds=(7
K03515	Human neuroleukin mRNA, complete cds /cds=(15,1691) /g		AB011086	Homo sapiens mRNA for KIAA0514 protein, complete cds /
U85773	Human phosphomannomutase (PMM2) mRNA, complete cds /cd		X79234	H.sapiens mRNA for ribosomal protein L11 /cds=(0,536)
D29805	Human mRNA for beta-1,4-galactosyltransferase, complet		D23660	Human mRNA for ribosomal protein, complete cds /cds=(5
AF038660	Homo sapiens chromosome 1p33-p34 beta-1,4-galactosyltr		D23661	Human mRNA for ribosomal protein L37, complete cds /cd
AF038661	Homo sapiens chromosome 1q21-1q23 beta-1,4-galactosylt		Z21507	H.sapiens EF-1delta gene encoding human elongation fac
Z34975	H.sapiens LDLC mRNA /cds=(95,2311) /gb=Z34975 /gi=5756		X06617	Human mRNA for ribosomal protein S11 /cds=(15,491) /gb
D79206	Homo sapiens gene for ryudocan core protein, exon1-5,		M15353	Homo sapiens cap-binding protein mRNA, complete cds /c
Z48199	H.sapiens syndecan-1 gene (exons 2-5) /cds=(0,866) /gb		X60489	Human mRNA for elongation factor-1-beta /cds=(235,912)
L09235	Human vacuolar ATPase (isoform VA68) mRNA, complete cd		L06845	Human cysteinyl-tRNA synthetase mRNA, partial cds /cds
D14710	Human mRNA for ATP synthase alpha subunit, complete cd		L19161	HUMIEF2G Human translation initiation factor
D89052	Homo sapiens mRNA for proton-ATPase-like protein, comp		M13932	Human ribosomal protein S17 mRNA, complete cds /cds=(2
M62762	Human vacuolar H+ ATPase proton channel subunit mRNA,		M13934	Human ribosomal protein S14 gene, complete cds /cds=(0
X16396	Human mRNA for NAD-dependent methylene tetrahydrofolat		J05032	Human aspartyl-tRNA synthetase alpha-2 subunit mRNA, c
J04031	HUMMDMCSF Human methylenetetrahydrofolate deh		S80343	S80343 ArgRS=arginyl-tRNA synthetase [human,
L35546	Homo sapiens gamma-glutamylcysteine synthetase light s		X80822	H.sapiens mRNA for ORF /cds=(133,555) /gb=X80822 /gi=5
M90656	Human gamma-glutamylcysteine synthetase (GCS) mRNA, co		J02984	Human insulinoma rig-analog mRNA encoding DNA-binding
U55206	Homo sapiens human gamma-glutamyl hydrolase (hGH) mRNA		Z49148	H.sapiens mRNA for ribosomal protein L29 /cds=(29,508)
D78177	Homo sapiens mRNA for quinolinate phosphoribosyl trans		Y11681	Homo sapiens mRNA for mitochondrial ribosomal protein
U89606	Human pyridoxal kinase mRNA, complete cds /cds=(6,944)		U54558	Homo sapiens translation initiation factor eIF3 p66 su
U51240	Human lysosomal-associated multitransmembrane protein		U54559	Homo sapiens translation initiation factor eIF3 p40 su
L76259	Homo sapiens PTS gene, complete cds /cds=(68,505) /gb=		U18937	Human histidyl-tRNA synthetase homolog (HO3) mRNA, com
M76231	Human sepiapterin reductase mRNA, complete cds /cds=(2		U49436	HSU49436 Human translation initiation factor
AF042386	Homo sapiens cyclophilin-33B (CYP-33) mRNA, complete c		L01124	Human ribosomal protein S13 (RPS13) mRNA, complete cds
X52851	Human cyclophilin gene for cyclophilin (EC 5.2.1.8) /c		Z12962	H.sapiens mRNA for homologue to yeast ribosomal protei
U29344	Human breast carcinoma fatty acid synthase mRNA, compl		U78525	Homo sapiens eukaryotic translation initiation factor
X17025	Human homolog of yeast IPP isomerase /cds=(50,736) /gb		U79262	Human deoxyhypusine synthase mRNA, complete cds /cds=(
M88468	Homo sapiens mevalonate kinase mRNA, complete cds /cds		M55409	HUMPANCAN Homo sapiens pancreatic tumor-relat
AJ002190	Homo sapiens cDNA for dihydroxyacetone phosphate acylt		L25899	Human ribosomal protein L10 mRNA, complete cds /cds=(1
L19783	Human GPI-H mRNA, complete cds /cds=(60,626) /gb=L1978		U26266	Human deoxyhypusine synthase mRNA, complete cds /cds=(
AF040707	Homo sapiens candidate tumor suppressor gene 21 protei		U09510	Human glycyl-tRNA synthetase mRNA, complete cds /cds=(
D42138	D42138 Homo sapiens mRNA for PIG-B, complete		X56932	H.sapiens mRNA for 23 kD highly basic protein /cds=(17
D85418	D85418 Homo sapiens mRNA for phosphatidylinos		U10117	Human endothelial-monocyte activating polypeptide II m
AB002135	Homo sapiens mRNA for glycosylphosphatidylinositol anc		U24169	Human JTV-1 (JTV-1) mRNA, complete cds /cds=(113,1051)
D13435	HUMPIGF Human mRNA for PIG-F (phosphatidyl-in		D21851	Human mRNA for KIAA0028 gene, partial cds /cds=(185,28
AL049933	Homo sapiens mRNA; cDNA DKFZp564K1216 (from clone DKFZ		X59303	Human G7a mRNA for valyl-tRNA synthetase /cds=(219,401
M87434	Human 71 kDa 25 oligoadenylate synthetase (p69 2-5A s		AF097441	Homo sapiens phenylalanine-tRNA synthetase (FARS1) mRN
M86707	Homo sapiens myristoyl CoA-protein N-myristoyltransfer		D13630	Human mRNA for KIAA0005 gene, complete cds /cds=(80,13
M74491	Human ADP-ribosylation factor 3 mRNA, complete cds /cd		AF064607	Homo sapiens GC20 protein mRNA, complete cds /cds=(70,
X04828	Human mRNA for G(i) protein alpha-subunit (adenylate c		AF104913	Homo sapiens eukaryotic protein synthesis initiation f
X07767	HSPKA Human mRNA for cAMP-dependent protein k		AF038957	Homo sapiens translation initiation factor 4e mRNA, co

M14333	HUMCSYNA Homo sapiens c-syn protooncogene mRN		U73824	Human p97 mRNA, complete cds /cds=(306,3029) /gb=U7382
X70326	H.sapiens MacMarcks mRNA /cds=(13,600) /gb=X70326 /gi=		U28424	Human protein kinase inhibitor p58 mRNA, complete cds
U61538	Human calcium-binding protein chp mRNA, complete cds /		J04617	HUMEF1A Human elongation factor EF-1-alpha ge
M27543	HUMGNAI1 Human guanine nucleotide-binding pro		U09877	Human helicase-like protein (HLP) mRNA, complete cds /
M15990	HUMCYES1 Human c-yes-1 mRNA		D12686	HUMEIF4G Human mRNA for eukaryotic initiation
M82809	Human annexin IV (ANX4) mRNA, complete cds /cds=(73,10		U29607	Human methionine aminopeptidase mRNA, complete cds /cd
M34181	Human testis-specific cAMP-dependent protein kinase ca		AF012072	Homo sapiens eIF4GII mRNA, complete cds /cds=(256,5013
M36340	Human ADP-ribosylation factor 1 (ARF1) mRNA, complete		AL080102	Homo sapiens mRNA; cDNA DKFZp564N1916 (from clone DKFZ
U05770	Human annexin V (ANX5) gene /cds=(164,1126) /gb=U05770		L34600	Human nuclear-encoded mitochondrial initiation factor
AF039656	Homo sapiens neuronal tissue-enriched acidic protein (		D26068	Human mRNA for KIAA0038 gene, partial cds /cds=(0,694)
AF043325	Homo sapiens N-myristoyltransferase 2 mRNA, complete c		X55733	H.sapiens initiation factor 4B cDNA /cds=(0,1835) /gb=
M20560	Human lipocortin-III mRNA, complete cds /cds=(46,1017)		D30655	HUMELF4AII Homo sapiens mRNA for eukaryotic i
Y00097	Human mRNA for protein p68 /cds=(100,2121) /gb=Y00097		U36764	HSU36764 Human TGF-beta receptor interacting
Y16521	Homo sapiens mRNA for CDS2 protein /cds=(258,1595) /gb		U50648	HSIIPKR17 Human interferon-inducible RNA-depe
D50840	Homo sapiens mRNA for ceramide glucosyltransferase, co		U39067	Homo sapiens translation initiation factor eIF3 p36 su
AB018356	Homo sapiens mRNA for GM3 synthase, complete cds /cds=		X70649	Homo sapiens DDX1 gene, complete CDS /cds=(288,2510) /
U11037	Human Sel-1 like mRNA, complete cds /cds=(11,298) /gb=		X95384	Homo sapiens mRNA for translational inhibitor protein
U58766	Human FX protein mRNA, complete cds /cds=(74,1039) /gb		X81625	H.sapiens mRNA for Cl1 protein /cds=(135,1448) /gb=X81
D50645	Homo sapiens mRNA for SDF2, complete cds /cds=(39,674)		AF054186	Homo sapiens p18 protein mRNA, complete cds /cds=(28,5
M58597	Human ELAM-1 ligand fucosyltransferase (ELFT) mRNA, co		S75463	P43=mitochondrial elongation factor homolog [human, li
X74837	H.sapiens HUMM9 mRNA /cds=(689,2566) /gb=X74837 /gi=41		X70940	H.sapiens mRNA for elongation factor 1 alpha-2 /cds=(8
D15057	Human mRNA for DAD-1, complete cds /cds=(66,407) /gb=D		L37936	Human nuclear-encoded mitochondrial elongation factor
D86967	Human mRNA for KIAA0212 gene, complete cds /cds=(58,20		X17644	Human GST1-Hs mRNA for GTP-binding protein /cds=(648,2
L38961	Human putative transmembrane protein precursor (B5) mR		AF013758	Homo sapiens polyadenylate binding protein-interacting
D29643	Human mRNA for KIAA0115 gene, complete cds /cds=(106,1		D32002	Human mRNA for nuclear cap binding protein, complete c
M90657	HUML6A Human tumor antigen (L6) mRNA, complet		X81788	Homo sapiens ICT1 (alias DS-1) mRNA /cds=(2,622) /gb=X
J04182	Homo sapiens lysosomal membrane glycoprotein-1 (LAMP1)		D86324	Homo sapiens mRNA for CMP-N-acetylneuraminic acid hydr
X53795	Human R2 mRNA for an inducible membrane protein /cds=(		M63167	HUMRACPC Human rac protein kinase alpha mRNA,
U15128	Human beta-1,2-N-acetylglucosaminyltransferase II (MGA		M81600	Human NAD(P)H-quinone oxireductase gene /cds=(111,935)
X12496	Human mRNA for erythrocyte membrane sialoglycoprotein		AF112219	Homo sapiens esterase D mRNA, complete cds /cds=(183,1
X62654	H.sapiens gene for Me491/CD63 antigen /cds=(69,785) /g		AB007882	Homo sapiens KIAA0422 mRNA, partial cds /cds=(0,2926)
U36336	Human lysosome-associated membrane protein-2b (LAMP2)		D25538	Human mRNA for KIAA0037 gene, complete cds /cds=(265,3
X77196	H.sapiens mRNA for lysosome-associated membrane protei		D14874	Homo sapiens mRNA for adrenomedullin precursor, comple
X76488	H.sapiens mRNA for lysosomal acid lipase /cds=(145,134		AF036927	Homo sapiens adenylyl cyclase type IX mRNA, complete c
X52151	Homo sapiens arylsulphatase A mRNA, complete cds /cds=		X02308	Human mRNA for thymidylate synthase (EC 2.1.1.45) /cds
U77413	Human O-linked GlcNAc transferase mRNA, complete cds /		D00596	HUMTS1 Homo sapiens gene for thymidylate synt
M97347	Human beta-1,6-N-acetylglucosaminyltransferase mRNA, c		Y00971	Human mRNA for phosphoriobosyl pyrophosphate synthetas
L00352	Human low density lipoprotein receptor gene /cds=(93,2		D00860	Homo sapiens mRNA for phosphoribosyl pyrophosphate syn
U41514	Human UDP-GalNAc-polypeptide N-acetylgalactosaminyltra		X15331	Human mRNA for phosphoribosylpyrophosphate synthetase
U07424	Human putative tRNA synthetase-like protein mRNA, comp		U56417	Human lysophosphatidic acid acyltransferase-alpha mRNA
U97188	Homo sapiens putative RNA binding protein KOC (koc) mR		U56418	Human lysophosphatidic acid acyltransferase-beta mRNA,
AF054187	Homo sapiens alpha NAC mRNA, complete cds /cds=(309,95		X75252	H.sapiens phosphatidylethanolamine binding protein mRN
AB016869	Homo sapiens mRNA for p70 ribosomal S6 kinase beta, co		U81802	HSU81802 Human PtdIns 4-kinase (PI4Kb) mRNA,
AF027302	Homo sapiens TNF-alpha stimulated ABC protein (ABC50)		L36151	Homo sapiens phosphatidylinositol 4-kinase mRNA, compl
AF006751	Homo sapiens ES/130 mRNA, complete cds /cds=(70,3003)		AF014398	Homo sapiens myo-inositol monophosphatase 2 mRNA, comp
U75686	Homo sapiens polyadenylate binding protein mRNA, compl		AF042729	Homo sapiens lithium-sensitive myo-inositol monophosph
U89436	Human tyrosyl-tRNA synthetase mRNA, complete cds /cds=		M95623	Homo sapiens hydroxymethylbilane synthase gene, comple
L38941	Homo sapiens ribosomal protein L34 (RPL34) mRNA, compl		AF104421	Homo sapiens isolate normal patient 1 uroporphyrinogen
D87735	Homo sapiens mRNA for ribosomal protein L14, complete		D16611	Human mRNA for coproporphyrinogen oxidase, complete cd
X94754	H.sapiens mRNA for yeast methionyl-tRNA synthetase hom		J03824	Human uroporphyrinogen III synthase mRNA, complete cds
M31520	Human ribosomal protein S24 mRNA /cds=(142,543) /gb=M3		Y00451	Human mRNA for 5-aminolevulinate synthase /cds=(83,201
U19796	HSU19796 Human melanoma antigen p15 mRNA, com		X53793	H.sapiens ADE2H1 mRNA showing homologies to SAICAR syn
X59892	H.sapiens mRNA for IFN-inducible gamma2 protein /cds=(		U10860	Human guanosine 5-monophosphate synthase mRNA, complet
X63527	H.sapiens mRNA for ribosomal protein L19 /cds=(28,618)		X52142	Human mRNA for CTP synthetase (EC 6.3.4.2) /cds=(75,18
U96074	Human translation initiation factor eIF3 p44 subunit m		D90282	Human carbamyl phosphate synthetase I (EC 6.3.4.16) mR
M64241	HUMQM Human Wilm s tumor-related protein (QM)		U20938	Human lymphocyte dihydropyrimidine dehydrogenase mRNA,
U37230	Human ribosomal protein L23a mRNA, complete cds /cds=(		J03626	Human UMP synthase mRNA, complete cds /cds=UNKNOWN /gb
D13748	HUM4AI Human mRNA for eukaryotic initiation f		D16532	Human gene for very low density lipoprotein receptor,
Z26876	H.sapiens gene for ribosomal protein L38 /cds=(110,322		AF027974	Homo sapiens clone LM1955 H105e3 gene, partial cds /cd
M58458	Human ribosomal protein S4 (RPS4X) isoform mRNA, compl		M14565	Human cholesterol side-chain cleavage enzyme P450scc m
X69391	H.sapiens mRNA for ribosomal protein L6 /cds=(26,892)		U22526	Human 2,3-oxidosqualene-lanosterol cyclase mRNA, compl
X69392	H.sapiens mRNA for ribosomal protein L26 /cds=(6,443)		AF034544	Homo sapiens delta7-sterol reductase mRNA, complete cd
M14199	HUMLAMR Human laminin receptor (2H5 epitope)		U19765	Human nucleic acid binding protein gene, complete cds
Z11692	H.sapiens mRNA for elongation factor 2 /cds=(0,2576) /		D14697	Human mRNA for KIAA0003 gene, complete cds /cds=(114,1
X91257	H.sapiens mRNA for seryl-tRNA synthetase /cds=(75,1619		U23942	Human lanosterol 14-demethylase cytochrome P450 (CYP51
D84273	Homo sapiens mRNA for Asparaginyl tRNA Synthetase, com		U79528	Human SR31747 binding protein 1 mRNA, complete cds /cd
X03342	Human mRNA for ribosomal protein L32 /cds=(34,441) /gb		D78130	Homo sapiens mRNA for squalene epoxidase, complete cds
X67247	H.sapiens rpS8 gene for ribosomal protein S8 /cds=(23,		L25931	HUMLBR Human lamin B receptor (LBR) mRNA, com
D14530	HUMRSPT Human homolog of yeast ribosomal prot		AB011004	Homo sapiens HuUAP1 mRNA for UDP-N-acetylglucosamine p
M17885	Human acidic ribosomal phosphoprotein P0 mRNA, complet		X56667	Human mRNA for calretinin /cds=(43,858) /gb=X56667 /gi
M17886	Human acidic ribosomal phosphoprotein P1 mRNA, complet			
L11566	Homo sapiens ribosomal protein L18 (RPL18) mRNA, compl			
D21853	Human mRNA for KIAA0111 gene, complete cds /cds=(214,1			

Supplementary Table 9: Decay Rates (hour^-1) for Accessions in HepG2 Experiments

The following accessions were listed as "present" (p<.04) in the initial and final condition in at least one of four HepG2 experiments Rate constant (hour^-1) and standard deviation are shown for each accession. A probe set from a pair of HepG2 hybridizations "contributes" to the shown decay rate inferences if, and only if, it is "Present" (p<0.04) in both experiments of the pair. The number of contributing probe sets for each accession is shown under "Probe set Information". If the intersection of the 99% probability intervals from the posterior distributions of all contributing probe sets is non-empty, the set of probe sets is called "consistent".

Note: inclusion of Present-->Absent probe sets does not alter the qualitative outcome of our analysis and number of probe sets included can exceed four due to the presence of accessions with multiple probe sets. Also, it is very important to realize that this table is only provided for reference. In the actual analysis, we weighted all probe sets equally to capture the patterns in Figure 1 and 2 (ie did not use this list for analysis directly).

Accession	Brief Description	Rate	StdDev	Probe Set Information
AB013382	Homo sapiens mRNA for DUSP6, complete cds /cds=(351,1	2.250	0.605	1 consistent probesets
M69043	HUMMAD3A Homo sapiens MAD-3 mRNA encoding IkB	2.000	0.302	1 consistent probesets
U39840	Human hepatocyte nuclear factor-3 alpha (HNF-3 alpha)	1.850	0.303	2 consistent probesets
V00568	HSMYC1 Human mRNA encoding the c-myc oncogene	1.840	0.317	5 inconsistent probesets
AF041259	Homo sapiens breast cancer putative transcription fact	1.750	0.353	1 consistent probesets
L76571	HUMSHP Homo sapiens nuclear hormone receptor	1.717	0.410	3 consistent probesets
AF050110	Homo sapiens TGFb inducible early protein and early gr	1.700	0.363	2 consistent probesets
X64318	H.sapiens E4BP4 gene /cds=(213,1601) /gb=X64318 /gi=30	1.583	0.268	3 consistent probesets
U07802	Human Tis11d gene, complete cds /cds=(291,1739) /gb=U0	1.504	0.298	4 inconsistent probesets
U76247	Human hSIAH1 mRNA, complete cds /cds=(186,1034) /gb=	1.475	0.497	2 consistent probesets
X77956	H.sapiens Id1 mRNA /cds=(35,499) /gb=X77956 /gi=457784	1.417	0.090	8 inconsistent probesets
U43842	Homo sapiens bone morphogenetic protein-4 (hBMP-4) gen	1.375	0.410	2 consistent probesets
X84373	H.sapiens mRNA for nuclear factor RIP140 /cds=(287,376	1.346	0.194	4 inconsistent probesets
D13891	Human mRNA for Id-2H, complete cds /cds=(96,500) /gb=D	1.317	0.205	4 consistent probesets
U61234	Human tubulin-folding cofactor C mRNA, complete cds /c	1.317	0.438	3 consistent probesets
L43821	Homo sapiens enhancer of filamentation (HEF1) mRNA, co	1.300	0.958	1 consistent probesets
AL049471	Homo sapiens mRNA; cDNA DKFZp586N012 (from clone D	1.275	0.248	4 consistent probesets
D50917	Human mRNA for KIAA0127 gene, complete cds /cds=(297	1.250	0.504	1 consistent probesets
U73960	Human ADP-ribosylation factor-like protein 4 mRNA, com	1.250	0.655	1 consistent probesets
AF041037	Homo sapiens novel antagonist of FGF signaling (sprout	1.250	0.806	1 consistent probesets
U23070	Human putative transmembrane protein (nma) mRNA, com	1.250	0.857	1 consistent probesets
X52560	Human gene for nuclear factor NF-IL6 /cds=(0,1037) /gb	1.242	0.149	4 inconsistent probesets
D86550	D86550 Human mRNA for serine/threonine protei	1.229	0.199	7 inconsistent probesets
AB004066	Homo sapiens mRNA for DEC1, complete cds /cds=(196,14	1.225	0.713	2 consistent probesets
X78992	H.sapiens ERF-2 mRNA /cds=(66,1544) /gb=X78992 /gi=50	1.200	0.353	1 consistent probesets
S81439	S81439 EGR alpha=early growth response gene a	1.200	0.857	1 consistent probesets
U80736	Homo sapiens CAGF9 mRNA, partial cds /cds=(0,995) /gb=	1.179	0.187	4 consistent probesets
Z46629	Homo sapiens SOX9 mRNA /cds=(359,1888) /gb=Z46629 /	1.171	0.184	4 inconsistent probesets
AJ222700	Homo sapiens mRNA for TSC-22 protein /cds=(191,625) /g	1.167	0.212	3 consistent probesets
L00352	Human low density lipoprotein receptor gene /cds=(93,2	1.163	0.170	4 inconsistent probesets
M82882	Human cis-acting sequence /cds=UNKNOWN /gb=M82882	1.150	0.504	1 consistent probesets
M22490	HUMBMP2B Human bone morphogenetic protein-2B	1.150	0.857	1 consistent probesets
U03105	Human B4-2 protein mRNA, complete cds /cds=(113,1096)	1.146	0.213	4 inconsistent probesets
X89750	H.sapiens mRNA for TGIF protein /cds=(311,1129) /gb=X8	1.138	0.184	4 consistent probesets
X79536	H.sapiens mRNA for hnRNPcore protein A1 /cds=(26,988)	1.125	0.215	4 inconsistent probesets
AI304854	qo19f03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-190	1.117	0.216	3 consistent probesets
AF032885	Homo sapiens forkhead protein (FKHR) mRNA, complete c	1.117	0.502	3 consistent probesets
M64497	Human apolipoprotein AI regulatory protein (ARP-1) mRN	1.100	0.263	3 consistent probesets
AJ002572	Homo sapiens mRNA; transcriptional unit N143 /cds=UNKN	1.100	0.554	1 consistent probesets
M59465	HUMA20 Human tumor necrosis factor alpha indu	1.083	0.242	4 consistent probesets
D15050	Human mRNA for transcription factor AREB6, complete cd	1.067	0.353	4 consistent probesets
AB023206	Homo sapiens mRNA for KIAA0989 protein, partial cds /c	1.050	0.504	1 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AB000584	AB000584 Homo sapiens mRNA for TGF-beta super	1.050	0.289	4 consistent probesets
D38305	Human mRNA for Tob, complete cds /cds=(43,1080) /gb=D	0.992	0.151	4 consistent probesets
AB007938	Homo sapiens mRNA for KIAA0469 protein, complete cds /	0.983	0.276	3 consistent probesets
M62831	Human transcription factor ETR101 mRNA, complete cds /	0.983	0.288	3 consistent probesets
AB028943	Homo sapiens mRNA for KIAA1020 protein, partial cds /c	0.975	0.463	2 consistent probesets
U78305	Homo sapiens protein phosphatase Wip1 mRNA, complete	0.975	0.241	4 consistent probesets
L16499	Human orphan homeobox protein (PRH) mRNA, complete c	0.971	0.200	4 consistent probesets
M60974	HUMGADD45 Human growth arrest and DNA-damage-	0.963	0.263	4 consistent probesets
M59287	Human protein kinase mRNA /cds=UNKNOWN /gb=M5928	0.950	0.403	1 consistent probesets
M63256	Human major Yo paraneoplastic antigen (CDR2) mRNA, 3	0.950	0.449	3 consistent probesets
X84709	H.sapiens mRNA for mediator of receptor-induced toxici	0.950	1.177	1 consistent probesets
X61123	Human BTG1 mRNA /cds=(308,823) /gb=X61123 /gi=2950	0.933	0.303	3 consistent probesets
L08246	HUMMCL1X Human myeloid cell differentiation p	0.931	0.125	7 inconsistent probesets
AB014515	Homo sapiens mRNA for KIAA0615 protein, complete cds /	0.925	0.294	2 consistent probesets
D86985	Human mRNA for KIAA0232 gene, complete cds /cds=(596	0.900	0.417	3 consistent probesets
D42040	Human mRNA for KIAA9001 gene, complete cds /cds=(170	0.900	0.437	3 consistent probesets
AB020631	Homo sapiens mRNA for KIAA0824 protein, partial cds /c	0.900	0.783	2 consistent probesets
U71267	Human potential transcriptional repressor NOT4Hp (NOT4	0.883	0.607	3 consistent probesets
Y10313	Homo sapiens mRNA IFRD1 (PC4) interferon-related devel	0.875	0.469	2 consistent probesets
D63877	Human mRNA for KIAA0157 gene, partial cds /cds=(0,1261	0.875	0.497	2 consistent probesets
M74558	Human SIL mRNA, complete cds /cds=(380,4243) /gb=M74	0.875	0.736	2 consistent probesets
S78771	NAT=CpG island-associated gene [human, mRNA, 1741 nt]	0.858	0.146	8 inconsistent probesets
U69274	Human zinc finger protein mRNA, complete cds /cds=(161	0.850	0.370	3 consistent probesets
AF004292	Homo sapiens clone IMAGE-30181 mRNA sequence /cds=	0.850	0.424	3 consistent probesets
S83308	SOX5=Sry-related HMG box gene {alternatively spliced}	0.850	0.441	2 consistent probesets
M65217	HUMHSF2 Human heat shock factor 2 (HSF2) mRNA	0.850	0.454	2 consistent probesets
X78925	H.sapiens HZF2 mRNA for zinc finger protein /cds=(0,21	0.846	0.356	4 consistent probesets
M57730	Human B61 mRNA, complete cds /cds=(73,690) /gb=M577	0.844	0.330	3 consistent probesets
AJ224819	Homo sapiens mRNA for candidate tumor suppressor invol	0.842	0.319	4 consistent probesets
AL096858	Novel human gene mapping to chomosome 1 /cds=(331,10	0.833	0.378	3 consistent probesets
Z25535	H.sapiens mRNA for nuclear pore complex protein hnup15	0.829	0.174	4 consistent probesets
J04102	HUMETS2A Human erythroblastosis virus oncogen	0.825	0.258	4 consistent probesets
M95724	HUMCENPRO H.sapiens centromere autoantigen C	0.825	0.665	2 consistent probesets
AB002345	Human mRNA for KIAA0347 gene, complete cds /cds=(122	0.817	0.398	2 consistent probesets
AB029039	Homo sapiens mRNA for KIAA1116 protein, complete cds /	0.813	0.179	4 consistent probesets
S73591	brain-expressed HHCPA78 homolog [human, HL-60 acute p	0.813	0.291	4 consistent probesets
AB006968	Homo sapiens mRNA for CIS4, complete cds /cds=(0,1607)	0.800	0.353	1 consistent probesets
AL080088	Homo sapiens mRNA; cDNA DKFZp564K2062 (from clone	0.800	0.439	1 consistent probesets
U75308	HSU75308 Human TBP-associated factor (hTAFII1	0.800	0.554	1 consistent probesets
D87685	Human mRNA for KIAA0244 gene, partial cds /cds=(0,5173	0.796	0.246	4 inconsistent probesets
AL079310	Novel human gene mapping to chomosome 22 /cds=(565,2	0.783	0.343	4 consistent probesets
AF048731	Homo sapiens cyclin T2a mRNA, complete cds /cds=(32,20	0.783	0.392	3 consistent probesets
X79067	H.sapiens ERF-1 mRNA 3 end /cds=UNKNOWN /gb=X790	0.783	0.602	3 consistent probesets
D31716	Human mRNA for GC box bindig protein, complete cds /cd	0.775	0.575	2 consistent probesets
X17576	HSNCK Human melanoma mRNA for nck protein, sh	0.775	0.351	2 consistent probesets
M57763	Human ADP-ribosylation factor (hARF6) mRNA, complete c	0.771	0.098	4 consistent probesets
AB028964	Homo sapiens mRNA for KIAA1041 protein, complete cds /	0.767	0.149	4 consistent probesets
AL050171	Homo sapiens mRNA; cDNA DKFZp586F1122 (from clone	0.758	0.157	4 consistent probesets
AF025770	Homo sapiens C2H2 zinc finger protein (ZNF189) mRNA, c	0.750	0.353	3 consistent probesets
U03644	Human recepin mRNA, complete cds /cds=(32,1387) /gb=U	0.750	0.504	1 consistent probesets
AF052099	Homo sapiens clone 23632 mRNA sequence /cds=UNKNO	0.750	0.706	1 consistent probesets
AJ131245	Homo sapiens mRNA for Sec24 protein (Sec24B isoform) /	0.737	0.213	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AA115140	zl10d12.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-501	0.733	0.229	3 consistent probesets
AB023199	Homo sapiens mRNA for KIAA0982 protein, complete cds /	0.733	0.418	3 consistent probesets
AB018327	Homo sapiens mRNA for KIAA0784 protein, partial cds /c	0.717	0.196	4 consistent probesets
U00115	Human zinc-finger protein (bcl-6) mRNA, complete cds /	0.717	0.259	4 consistent probesets
AB020670	Homo sapiens mRNA for KIAA0863 protein, complete cds /	0.717	0.471	3 consistent probesets
D50926	Human mRNA for KIAA0136 gene, partial cds /cds=(0,2854	0.708	0.129	4 consistent probesets
S62539	S62539 insulin receptor substrate-1 [human, s	0.707	0.119	10 inconsistent probesets
AF072250	Homo sapiens methyl-CpG binding protein MBD4 (MBD4) r	0.704	0.221	4 consistent probesets
U72649	Human BTG2 (BTG2) mRNA, complete cds /cds=(71,547) /	0.700	0.344	4 consistent probesets
AL049423	Homo sapiens mRNA; cDNA DKFZp586B211 (from clone D	0.700	0.359	3 inconsistent probesets
D31770	Human osteosarcoma mRNA for activin typeII A receptor,	0.700	0.605	1 consistent probesets
AB020680	Homo sapiens mRNA for KIAA0873 protein, partial cds /c	0.683	0.265	3 consistent probesets
M37197	HUMCBF Human CCAAT-box-binding factor (CBF) m	0.681	0.179	8 consistent probesets
AF006513	Homo sapiens CHD1 mRNA, complete cds /cds=(163,5292	0.679	0.161	4 consistent probesets
L40411	Homo sapiens thyroid receptor interactor (TRIP8) mRNA,	0.675	0.489	2 consistent probesets
U77664	Human RNaseP protein p38 (RPP38) mRNA, complete cds	0.675	0.608	2 consistent probesets
AI126004	qc50e12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-17	0.675	0.139	4 consistent probesets
AB004884	Homo sapiens mRNA for PKU-alpha, partial cds /cds=(0,2	0.671	0.205	4 consistent probesets
AB002803	Homo sapiens BACH1 mRNA, complete cds /cds=(118,232	0.667	0.247	3 consistent probesets
U09366	Human zinc finger protein ZNF133 /cds=(445,2409) /gb=U	0.667	0.282	3 consistent probesets
W63793	zc55a10.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-326	0.663	0.098	4 consistent probesets
AA114830	zk88e06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-489	0.656	0.198	3 consistent probesets
D85429	D85429 Homo sapiens gene for heat shock prote	0.650	0.100	4 inconsistent probesets
W26521	32g11 Homo sapiens cDNA /gb=W26521 /gi=1307382 /	0.650	0.215	4 consistent probesets
AB002369	Human mRNA for KIAA0371 gene, complete cds /cds=(247	0.650	0.403	1 consistent probesets
AF008442	Homo sapiens RNA polymerase I subunit hRPA39 mRNA, c	0.650	0.403	1 consistent probesets
AF150247	AF150247 Homo sapiens cDNA /clone=CBFBCC09 /gb=AF	0.650	0.504	1 consistent probesets
AF001461	Homo sapiens Kruppel-like zinc finger protein Zf9 mRNA	0.650	0.706	1 consistent probesets
X62048	H.sapiens Wee1 hu gene /cds=(170,2110) /gb=X62048 /gi=	0.646	0.145	4 consistent probesets
U66469	HSU66469 Human cell growth regulator CGR19 mR	0.646	0.215	8 consistent probesets
D79994	Human mRNA for KIAA0172 gene, partial cds /cds=(0,3923	0.642	0.091	4 consistent probesets
S74017	S74017 Nrf2=NF-E2-like basic leucine zipper t	0.638	0.078	4 inconsistent probesets
AF011468	Homo sapiens serine/threonine kinase (BTAK) mRNA, com	0.635	0.095	8 inconsistent probesets
AF059611	Homo sapiens nuclear matrix protein NRP/B (NRPB) mRNA	0.633	0.077	4 inconsistent probesets
U76638	HSU76638 Human BRCA1-associated RING domain p	0.633	0.276	3 consistent probesets
U06632	Homo sapiens p80-coilin mRNA, complete cds /cds=(22,17	0.633	0.431	3 consistent probesets
AB020661	Homo sapiens mRNA for KIAA0854 protein, complete cds /	0.629	0.179	4 consistent probesets
AF030339	Homo sapiens receptor for viral semaphorin protein (VE	0.629	0.209	4 inconsistent probesets
U47741	Human CREB-binding protein (CBP) mRNA, complete cds /	0.625	0.299	4 consistent probesets
X66360	HSSTHPKC H.sapiens mRNA PCTAIRE-2 for serine/	0.625	0.484	2 consistent probesets
AF071771	Homo sapiens SPH-binding factor mRNA, partial cds /cds	0.625	0.644	2 consistent probesets
D87467	Human mRNA for KIAA0277 gene, complete cds /cds=(55,1	0.621	0.352	4 consistent probesets
AF020762	Homo sapiens clone 1400 unknown protein mRNA, partial	0.617	0.124	4 consistent probesets
U25435	Human transcriptional repressor (CTCF) mRNA, complete	0.617	0.165	4 consistent probesets
AB002306	Human mRNA for KIAA0308 gene, partial cds /cds=(0,3895	0.617	0.326	3 consistent probesets
D87077	Human mRNA for KIAA0240 gene, partial cds /cds=(0,2953	0.617	0.337	3 consistent probesets
AB007949	Homo sapiens mRNA for KIAA0480 protein, complete cds /	0.617	0.337	3 consistent probesets
W27541	32c12 Homo sapiens cDNA /gb=W27541 /gi=1307345 /	0.617	0.527	3 consistent probesets
AB007930	Homo sapiens mRNA for KIAA0461 perotein, partial cds /	0.613	0.268	4 consistent probesets
W21827	57E11 Homo sapiens cDNA /clone=(not-directional) /gb=W	0.612	0.146	4 consistent probesets
U68494	Human hbc647 mRNA sequence /cds=UNKNOWN /gb=U68	0.612	0.277	4 consistent probesets
D76444	Homo sapiens hkf-1 mRNA, complete cds /cds=(922,2979)	0.608	0.177	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AA418437	zv92d11.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-767	0.608	0.292	4 consistent probesets
M21154	HUMAMD Human S-adenosylmethionine decarboxyla	0.607	0.061	12 inconsistent probesets
AL021977	bK447C4.1 (novel MAFF (v-maf musculoaponeurotic fibros	0.604	0.208	4 inconsistent probesets
D31888	Human mRNA for KIAA0071 gene, partial cds /cds=(0,1190	0.604	0.213	4 consistent probesets
D87450	Human mRNA for KIAA0261 gene, partial cds /cds=(0,3865	0.604	0.238	4 consistent probesets
AL080090	Homo sapiens mRNA; cDNA DKFZp564L0562 (from clone	0.600	0.161	2 consistent probesets
AF035119	Homo sapiens deleted in liver cancer-1 (DLC-1) mRNA, c	0.600	0.180	3 consistent probesets
AL080149	Homo sapiens mRNA; cDNA DKFZp434B094 (from clone D	0.600	0.194	3 consistent probesets
X70683	H.sapiens mRNA for SOX-4 protein /cds=(350,1774) /gb=X	0.600	0.223	3 consistent probesets
U90912	Human clone 23865 mRNA sequence /cds=UNKNOWN /gb	0.600	0.287	2 consistent probesets
AB000461	Homo sapiens mRNA, complete cds, clone-RES4-22C /cds	0.600	0.303	3 consistent probesets
L04282	HUMTB Human CACCC box-binding protein mRNA, c	0.600	0.353	1 consistent probesets
AB023221	Homo sapiens mRNA for KIAA1004 protein, partial cds /c	0.600	0.545	2 consistent probesets
AL041879	DKFZp434H0419_r1 Homo sapiens cDNA, 5 end /clone=D	0.600	0.554	1 consistent probesets
AB007883	Homo sapiens KIAA0423 mRNA, partial cds /cds=(0,5090)	0.600	0.605	1 consistent probesets
AL050064	Homo sapiens mRNA; cDNA DKFZp566L033 (from clone D	0.600	0.655	1 consistent probesets
X51435	Human PRDII-BF1 gene for a DNA-binding protein /cds=(3	0.600	0.655	1 consistent probesets
AB014575	Homo sapiens mRNA for KIAA0675 protein, complete cds /	0.600	0.806	1 consistent probesets
X98494	H.sapiens mRNA for M phase phosphoprotein 10 /cds=(0,2	0.596	0.141	4 consistent probesets
AL040137	DKFZp434D1813_s1 Homo sapiens cDNA, 3 end /clone=D	0.592	0.268	4 consistent probesets
AL049432	Homo sapiens mRNA; cDNA DKFZp586J231 (from clone D	0.588	0.170	4 consistent probesets
J03407	Human rfp transforming protein mRNA, complete cds /cds	0.587	0.229	4 consistent probesets
U20240	Human C/EBP gamma mRNA, complete cds /cds=(250,702	0.583	0.164	4 consistent probesets
AB011178	Homo sapiens mRNA for KIAA0606 protein, partial cds /c	0.583	0.257	3 consistent probesets
AF031383	Homo sapiens hMed7 (MED7) mRNA, complete cds /cds=(	0.583	0.313	4 consistent probesets
AL035447	Human DNA sequence from clone 1183I21 on chromosome	0.583	0.385	3 consistent probesets
AB014597	Homo sapiens mRNA for KIAA0697 protein, partial cds /c	0.579	0.102	4 consistent probesets
AL050164	Homo sapiens mRNA; cDNA DKFZp586C1622 (from clone	0.579	0.163	4 consistent probesets
U76248	Human hSIAH2 mRNA, complete cds /cds=(526,1500) /gb=	0.575	0.081	4 inconsistent probesets
AJ010014	Homo sapiens mRNA for M96A protein /cds=(243,2024) /gb	0.575	0.180	4 consistent probesets
D38550	Human mRNA for KIAA0075 gene, partial cds /cds=(0,524)	0.575	0.451	2 consistent probesets
AF038951	Homo sapiens DNA-binding protein mRNA, complete cds /c	0.575	0.500	2 consistent probesets
AF048755	Homo sapiens HsPex13p (PEX13) mRNA, complete cds /cd	0.575	0.626	2 consistent probesets
AB028999	Homo sapiens mRNA for KIAA1076 protein, partial cds /c	0.575	0.814	2 consistent probesets
M26880	HUMUBI13 Human ubiquitin mRNA, complete cds"	0.567	0.113	8 inconsistent probesets
AB002333	Human mRNA for KIAA0335 gene, complete cds /cds=(226	0.567	0.170	4 consistent probesets
AL050102	Homo sapiens mRNA; cDNA DKFZp586F1019 (from clone	0.567	0.409	3 consistent probesets
U15655	HSU15655 Human ets domain protein ERF mRNA, c	0.565	0.193	8 inconsistent probesets
AF070570	Homo sapiens clone 24473 mRNA sequence /cds=UNKNO	0.563	0.103	4 consistent probesets
AI671547	wb33e07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.563	0.205	4 consistent probesets
D50911	Human mRNA for KIAA0121 gene, complete cds /cds=(410	0.558	0.186	4 consistent probesets
AB018285	Homo sapiens mRNA for KIAA0742 protein, partial cds /c	0.554	0.111	4 inconsistent probesets
AI434146	ti36g07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-213	0.554	0.193	4 consistent probesets
U51166	Human G/T mismatch-specific thymine DNA glycosylase m	0.554	0.297	4 consistent probesets
AL049940	Homo sapiens mRNA; cDNA DKFZp564E1922 (from clone	0.550	0.085	4 inconsistent probesets
J03626	Human UMP synthase mRNA, complete cds /cds=UNKNO	0.550	0.138	3 consistent probesets
AI830496	wh51h03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.550	0.227	4 consistent probesets
AF038177	Homo sapiens clone 23899 mRNA sequence /cds=UNKNO	0.550	0.252	2 consistent probesets
D87930	Homo sapiens mRNA for myosin phosphatase target subun	0.550	0.293	3 consistent probesets
AB007855	Homo sapiens KIAA0395 mRNA, partial cds /cds=(0,1381)	0.550	0.510	2 consistent probesets
U78524	Human Gu binding protein mRNA, partial cds /cds=(0,193	0.550	0.605	1 consistent probesets
AB020626	Homo sapiens mRNA for KIAA0819 protein, partial cds /c	0.550	0.655	1 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D89859	Homo sapiens mRNA for zinc finger 5 protein, complete	0.550	0.756	1 consistent probesets
U09412	Human zinc finger protein ZNF134 mRNA, complete cds /c	0.550	0.806	1 consistent probesets
AL049321	Homo sapiens mRNA; cDNA DKFZp564D156 (from clone D	0.550	0.137	4 consistent probesets
AF080227	Homo sapiens embryonic ectoderm development protein m	0.546	0.151	4 consistent probesets
D43951	Human mRNA for KIAA0099 gene, complete cds /cds=(56,3	0.542	0.096	4 inconsistent probesets
U77914	Human soluble protein Jagged mRNA, partial cds /cds=(0	0.542	0.127	4 inconsistent probesets
AF038844	Homo sapiens MKP-1 like protein tyrosine phosphatase m	0.538	0.130	4 consistent probesets
U84487	HSU84487 Human CX3C chemokine precursor, mRNA	0.538	0.165	4 consistent probesets
AB002311	Human mRNA for KIAA0313 gene, complete cds /cds=(62,4	0.538	0.183	4 consistent probesets
U48251	HSU48251 Homo sapiens protein kinase C-bindin	0.533	0.213	3 consistent probesets
AB018342	Homo sapiens mRNA for KIAA0799 protein, partial cds /c	0.533	0.235	3 consistent probesets
AB018309	Homo sapiens mRNA for KIAA0766 protein, complete cds /	0.533	0.254	3 consistent probesets
AI475497	tj92g12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-214	0.533	0.259	3 consistent probesets
AA037278	zc52c04.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-325	0.533	0.288	3 consistent probesets
AL049842	Human DNA sequence from clone 129L7 on chromosome 6	0.533	0.397	3 consistent probesets
W22296	65A11 Homo sapiens cDNA /clone=(not-directional) /gb=W	0.529	0.118	4 consistent probesets
AB020315	Homo sapiens Dickkopf-1 (hdkk-1) gene /cds=(0,800) /gb	0.529	0.162	4 consistent probesets
AI123426	qa49c09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.529	0.167	4 consistent probesets
AF017790	Homo sapiens retinoblastoma-associated protein HEC mRN	0.525	0.214	4 consistent probesets
AL096880	Novel human mRNA containing Zinc finger C2H2 type dom	0.525	0.241	4 consistent probesets
L03426	Human XE7 mRNA, complete alternate coding regions /cds	0.525	0.135	4 consistent probesets
X51804	Human PMI gene for a putative receptor protein /cds=(2	0.525	0.215	4 consistent probesets
AB009010	Homo sapiens mRNA for polyubiquitin UbC, complete cds	0.519	0.141	8 inconsistent probesets
AF091263	Homo sapiens RNA binding motif protein 5 (RBM5) mRNA,	0.517	0.173	3 consistent probesets
X99050	H.sapiens mRNA; UV Radiation Resistance Associated Ger	0.517	0.448	3 consistent probesets
AL096842	Homo sapiens mRNA; cDNA DKFZp586D1519 (from clone	0.513	0.211	4 consistent probesets
D45906	Homo sapiens mRNA for LIMK-2, complete cds /cds=(114,2	0.508	0.282	4 consistent probesets
L19872	Human AH-receptor mRNA, complete cds /cds=(375,2921)	0.508	0.287	4 consistent probesets
AB014458	Homo sapiens hUBP mRNA for ubiquitin specific protease	0.504	0.140	4 consistent probesets
AB018283	Homo sapiens mRNA for KIAA0740 protein, complete cds /	0.500	0.183	4 consistent probesets
U09825	Human acid finger protein mRNA, complete cds /cds=(555	0.500	0.186	3 consistent probesets
AB005754	Homo sapiens mRNA for LAK-1, complete cds /cds=(126,17	0.500	0.253	3 consistent probesets
AB020700	Homo sapiens mRNA for KIAA0893 protein, complete cds /	0.500	0.273	2 consistent probesets
AL118582	DKFZp761B0810_r1 Homo sapiens cDNA, 5 end /clone=D	0.500	0.275	3 consistent probesets
AF083105	Homo sapiens HMG box factor SOX-13 mRNA, complete c	0.500	0.282	3 consistent probesets
AB023234	Homo sapiens mRNA for KIAA1017 protein, complete cds /	0.500	0.410	2 consistent probesets
AL050374	Homo sapiens mRNA; cDNA DKFZp586C1619 (from clone	0.500	0.439	1 consistent probesets
X80200	H.sapiens MLN62 mRNA /cds=(85,1497) /gb=X80200 /gi=9	0.500	0.443	3 consistent probesets
M16750	HUMPIM1 Human pim-1 oncogene mRNA, complete c	0.500	0.504	1 consistent probesets
D26018	Human mRNA for KIAA0039 gene, partial cds /cds=(0,1475	0.500	0.554	1 consistent probesets
U13948	Human zinc finger/leucine zipper protein (AF10) mRNA,	0.500	0.554	1 consistent probesets
Y10936	H.sapiens mRNA for hypothetical protein downstream of	0.500	0.655	1 consistent probesets
Z16411	HSPLPC H.sapiens mRNA encoding phospholipase	0.500	0.857	1 consistent probesets
X83492	HSFAS47 H.sapiens mRNA for Fas/Apo-1 (clone p	0.500	0.958	1 consistent probesets
X80821	H.sapiens mRNA for ribosomal protein L18a homologue /c	0.500	1.109	1 consistent probesets
AF055001	Homo sapiens clone 24560 unknown mRNA, complete cds	0.496	0.095	4 consistent probesets
AF052178	Homo sapiens clone 24523 mRNA sequence /cds=UNKNO	0.496	0.135	4 consistent probesets
AF009353	Homo sapiens transcription intermediary factor 1 (TIF1	0.496	0.210	4 consistent probesets
U23946	HSU23946 Human putative tumor suppressor (LUC	0.492	0.151	4 consistent probesets
D86961	Human mRNA for KIAA0206 gene, partial cds /cds=(0,581)	0.492	0.167	4 consistent probesets
U26710	HSU26710 Human cbl-b mRNA, complete cds"	0.492	0.211	4 consistent probesets
AL039458	DKFZp434N0910_s1 Homo sapiens cDNA, 3 end /clone=D	0.492	0.239	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
M99435	Human transducin-like enhancer protein (TLE1) mRNA, co	0.488	0.215	4 consistent probesets
AI950382	wp10g06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.488	0.238	4 consistent probesets
X77743	HSCDKAK H.sapiens CDK activating kinase mRNA	0.483	0.105	4 consistent probesets
U85430	Human transcription factor NFATx4 mRNA, complete cds /	0.483	0.159	3 inconsistent probesets
AB007889	Homo sapiens KIAA0429 mRNA, complete cds /cds=(2373,	0.483	0.184	4 consistent probesets
AB015342	Homo sapiens HRIHFB2436 mRNA, partial cds /cds=(0,674	0.483	0.282	3 consistent probesets
U39817	HSU39817 Human Bloom s syndrome protein (BLM)	0.483	0.292	3 consistent probesets
Y11525	H.sapiens mRNA for CCAAT/enhancer binding protein alph	0.483	0.296	3 consistent probesets
X63468	H.sapiens mRNA for transcription factor TFIIE alpha /c	0.483	0.315	3 consistent probesets
AF055993	Homo sapiens mSin3A associated polypeptide p30 mRNA,	0.483	0.485	3 consistent probesets
AL096744	Homo sapiens mRNA; cDNA DKFZp566H033 (from clone D	0.479	0.101	4 inconsistent probesets
AJ238764	Homo sapiens mRNA for UDP-N-acetylglucosamine-2-epim	0.479	0.145	4 consistent probesets
AL080063	Homo sapiens mRNA; cDNA DKFZp564I052 (from clone D	0.475	0.217	4 consistent probesets
AB023148	Homo sapiens mRNA for KIAA0931 protein, partial cds /c	0.475	0.220	4 inconsistent probesets
D43772	HUMGRB7 Human squamous cell carcinama of esop	0.475	0.228	2 consistent probesets
AB028987	Homo sapiens mRNA for KIAA1064 protein, partial cds /c	0.475	0.323	2 consistent probesets
AJ132440	Homo sapiens mRNA for PLU-1 protein /cds=(89,4723) /gb	0.475	0.385	2 consistent probesets
M97676	HUMHOX7 Homo sapiens (region 7) homeobox prot	0.474	0.144	7 consistent probesets
X68560	H.sapiens SPR-2 mRNA for GT box binding protein /cds=(	0.471	0.108	4 consistent probesets
AB007927	Homo sapiens mRNA for KIAA0458 protein, complete cds /	0.471	0.117	4 consistent probesets
U44060	Human homeodomain protein (Prox 1) mRNA, complete cd	0.471	0.154	4 inconsistent probesets
X55504	HSP120A H.sapiens mRNA for P120 antigen	0.471	0.186	4 consistent probesets
X79204	H.sapiens SCA1 mRNA for ataxin /cds=(935,3385) /gb=X79	0.471	0.246	4 consistent probesets
L02547	Homo sapiens (clone pZ50-19) cleavage stimulation fact	0.471	0.264	4 consistent probesets
D50683	D50683 Homo sapiens mRNA for TGF-betaIIR alph	0.469	0.079	8 inconsistent probesets
D14041	Homo sapiens mRNA for H-2K binding factor-2, complete	0.467	0.179	4 consistent probesets
AF039945	Homo sapiens synaptojanin 2B mRNA, partial cds /cds=(0	0.467	0.198	4 consistent probesets
AB002337	Human mRNA for KIAA0339 gene, complete cds /cds=(686	0.467	0.216	3 consistent probesets
AJ132917	Homo sapiens mRNA for methyl-CpG-binding protein 2 /cd	0.467	0.225	4 consistent probesets
AB014593	Homo sapiens mRNA for KIAA0693 protein, partial cds /c	0.467	0.231	3 consistent probesets
AI538172	ti75f08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2137	0.467	0.256	3 consistent probesets
AB028976	Homo sapiens mRNA for KIAA1053 protein, partial cds /c	0.467	0.316	3 consistent probesets
X70394	H.sapiens OZF mRNA /cds=(856,1734) /gb=X70394 /gi=468	0.463	0.095	4 consistent probesets
X78947	H.sapiens mRNA for connective tissue growth factor /cd	0.463	0.102	4 inconsistent probesets
AW020536	df11b12.y1 Homo sapiens cDNA, 5 end /clone=IMAGE-248	0.463	0.136	4 consistent probesets
AB000468	Homo sapiens mRNA for zinc finger protein, complete cd	0.463	0.162	4 consistent probesets
AI191768	qd62e10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-17	0.463	0.249	4 consistent probesets
AB018312	Homo sapiens mRNA for KIAA0769 protein, complete cds /	0.458	0.191	4 consistent probesets
Y13247	Homo sapiens fb19 mRNA /cds=(539,3361) /gb=Y13247 /gi	0.458	0.289	4 consistent probesets
U07664	HSHB9HB2 Human HB9 homeobox gene, exons 2 and	0.458	0.302	4 consistent probesets
Z93930	Human DNA sequence from clone 292E10 on chromosome	0.454	0.051	8 inconsistent probesets
AB020710	Homo sapiens mRNA for KIAA0903 protein, partial cds /c	0.454	0.080	4 consistent probesets
AF022375	Homo sapiens vascular endothelial growth factor mRNA,	0.454	0.097	4 consistent probesets
X87613	H.sapiens mRNA for skeletal muscle abundant protein /c	0.454	0.119	4 consistent probesets
M83667	HUMNFIL6BA Human NF-IL6-beta protein mRNA, co	0.454	0.143	4 inconsistent probesets
AF099935	Homo sapiens MDC-3.13 isoform 2 mRNA, complete cds /c	0.454	0.166	4 consistent probesets
AB002353	Human mRNA for KIAA0355 gene, complete cds /cds=(838	0.454	0.217	4 consistent probesets
AL050071	Homo sapiens mRNA; cDNA DKFZp566B0846 (from clone	0.454	0.245	4 consistent probesets
AF023917	Homo sapiens protein tyrosine phosphatase PIR1 mRNA, c	0.450	0.083	4 consistent probesets
L36720	Homo sapiens bystin mRNA, complete cds /cds=(64,984) /	0.450	0.139	4 consistent probesets
AI743299	wg91b04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.450	0.159	3 consistent probesets
AB020689	Homo sapiens mRNA for KIAA0882 protein, partial cds /c	0.450	0.178	2 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
X56687	H.sapiens mRNA for autoantigen NOR-90 /cds=(179,2362)	0.450	0.188	3 consistent probesets
AB007939	Homo sapiens mRNA for KIAA0470 protein, complete cds /	0.450	0.252	1 consistent probesets
AB020697	Homo sapiens mRNA for KIAA0890 protein, complete cds /	0.450	0.252	1 consistent probesets
M68941	HUMPTYPH Human protein-tyrosine phosphatase m	0.450	0.299	4 consistent probesets
AB028069	Homo sapiens mRNA for activator of S phase Kinase, com	0.450	0.353	1 consistent probesets
D79998	Human mRNA for KIAA0176 gene, partial cds /cds=(0,797)	0.450	0.353	1 consistent probesets
U97198	Homo sapiens CG1 mRNA, complete cds /cds=(219,1490) /	0.450	0.353	1 consistent probesets
AB002304	Human mRNA for KIAA0306 gene, partial cds /cds=(0,4357	0.450	0.375	2 consistent probesets
L05144	Homo sapiens (clone lamda-hPEC-3) phosphoenolpyruvate	0.450	0.403	1 consistent probesets
Y18264	Homo sapiens SALL1 gene, exon 1 and joined CDS /cds=(2	0.450	0.454	1 consistent probesets
D50923	Human mRNA for KIAA0133 gene, complete cds /cds=(136	0.450	0.454	1 consistent probesets
J04178	Human abnormal beta-hexosaminidase alpha chain (HEXA)	0.450	0.454	1 consistent probesets
X59656	HSCRKL H.sapiens crk-like gene CRKL	0.450	0.521	3 consistent probesets
AA203246	zx54h03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-446	0.450	0.706	1 consistent probesets
W28614	49b5 Homo sapiens cDNA /gb=W28614 /gi=1308562 /	0.450	0.806	1 consistent probesets
X98175	H.sapiens mRNA for MACH-beta-2 protein /cds=(0,277) /g	0.450	0.857	1 consistent probesets
U44378	HSU44378 Human homozygous deletion target in	0.447	0.094	12 consistent probesets
AL050144	Homo sapiens mRNA; cDNA DKFZp586C1620 (from clone	0.446	0.183	4 consistent probesets
M86699	HUMTTK Human kinase (TTK) mRNA, complete cds"	0.446	0.198	4 consistent probesets
U43899	HSU43899 Human signal transducing adaptor mol	0.442	0.137	4 consistent probesets
AL049246	Homo sapiens mRNA; cDNA DKFZp564C053 (from clone D	0.442	0.147	4 consistent probesets
U28946	HSU28946 Human G/T mismatch binding protein (	0.442	0.160	4 consistent probesets
M54915	HUMPIM1LE Human h-pim-1 protein (h-pim-1) mRN	0.442	0.189	4 consistent probesets
AB023207	Homo sapiens mRNA for KIAA0990 protein, complete cds /	0.442	0.200	4 consistent probesets
W21884	58c2 Homo sapiens cDNA /clone=(not-directional) /gb=W2	0.442	0.205	4 consistent probesets
D80004	Human mRNA for KIAA0182 gene, partial cds /cds=(0,3474	0.439	0.185	3 consistent probesets
L13773	Human AF-4 mRNA, complete cds /cds=(420,4052) /gb=L1	0.438	0.112	4 consistent probesets
AB011151	Homo sapiens mRNA for KIAA0579 protein, partial cds /c	0.438	0.132	4 consistent probesets
AB023195	Homo sapiens mRNA for KIAA0978 protein, partial cds /c	0.438	0.151	4 consistent probesets
AB020638	Homo sapiens mRNA for KIAA0831 protein, complete cds /	0.438	0.170	4 consistent probesets
AL049450	Homo sapiens mRNA; cDNA DKFZp586B1922 (from clone	0.438	0.224	4 consistent probesets
X59268	Human mRNA for general transcription factor IIB /cds=(	0.438	0.264	4 consistent probesets
AF012108	Homo sapiens Amplified in Breast Cancer (AIB1) mRNA, c	0.433	0.058	4 inconsistent probesets
AB011148	Homo sapiens mRNA for KIAA0576 protein, partial cds /c	0.433	0.150	4 consistent probesets
AJ012755	Homo sapiens mRNA for TL132 /cds=(1241,2305) /gb=AJ0	0.433	0.153	4 consistent probesets
AI935420	wo84c08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.433	0.157	4 consistent probesets
U61167	HSU61167 Human SH3 domain-containing protein	0.433	0.176	3 consistent probesets
X67155	H.sapiens mRNA for mitotic kinesin-like protein-1 /cds	0.433	0.247	3 consistent probesets
X96401	H.sapiens mRNA for ROX protein /cds=(212,1960) /gb=X96	0.433	0.283	3 consistent probesets
AB011088	Homo sapiens mRNA for KIAA0516 protein, partial cds /c	0.433	0.378	3 consistent probesets
AL046961	DKFZp586J0417_r1 Homo sapiens cDNA, 5 end /clone=DI	0.433	0.514	1 consistent probesets
AB028969	Homo sapiens mRNA for KIAA1046 protein, complete cds /	0.429	0.093	4 inconsistent probesets
D28118	HUMDB1 Human mRNA for DB1, complete cds"	0.429	0.124	8 inconsistent probesets
AF055029	Homo sapiens clone 24711 mRNA sequence /cds=UNKNO	0.429	0.154	4 consistent probesets
U92538	Homo sapiens origin recognition complex subunit 5 homo	0.429	0.197	4 inconsistent probesets
Y13115	HSSTPKSAK Homo sapiens mRNA for serine/threon	0.427	0.243	5 consistent probesets
U79263	Human clone 23760 mRNA, partial cds /cds=(0,1021) /gb=	0.425	0.098	4 inconsistent probesets
D87445	Human mRNA for KIAA0256 gene, complete cds /cds=(142	0.425	0.230	4 consistent probesets
AF016270	Homo sapiens thyroid hormone receptor coactivating pro	0.425	0.259	2 consistent probesets
AF052102	Homo sapiens clone 23926 mRNA sequence /cds=UNKNO	0.425	0.308	2 consistent probesets
Y18206	Homo sapiens mRNA for protein phosphatase 1 (PPP1R6) /	0.425	0.363	2 consistent probesets
S66427	S66427 RBP1=retinoblastoma binding protein 1	0.425	0.454	2 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U43746	HSU43746 Human breast cancer susceptibility (	0.425	0.489	2 consistent probesets
W28612	49b3 Homo sapiens cDNA /gb=W28612 /gi=1308560 /	0.421	0.145	4 consistent probesets
AF019214	Homo sapiens HMG box containing protein 1 mRNA, comp	0.421	0.264	4 consistent probesets
AB024057	Homo sapiens mRNA for vascular Rab-GAP/TBC-containin	0.417	0.114	4 consistent probesets
D63476	Human mRNA for KIAA0142 gene, complete cds /cds=(473	0.417	0.124	4 consistent probesets
AF024710	AF024710 Homo sapiens vascular endothelial gr	0.417	0.135	4 consistent probesets
U10564	HSU10564 Human CDK tyrosine 15-kinase WEE1Hu	0.417	0.147	3 consistent probesets
X92493	H.sapiens mRNA for STM-7 protein /cds=(419,2041) /gb=X	0.417	0.231	4 consistent probesets
AB007925	Homo sapiens mRNA for KIAA0456 protein, partial cds /c	0.417	0.303	3 consistent probesets
M55654	Human TATA-binding protein mRNA, complete cds /cds=(2	0.417	0.348	3 consistent probesets
X95152	HSBRCA22 H.sapiens brca2 gene exon 2 (and joi	0.417	0.373	3 consistent probesets
U51990	Human hPrp18 mRNA, complete cds /cds=(72,1100) /gb=U	0.417	0.385	4 consistent probesets
AF058918	Homo sapiens unknown mRNA /cds=(212,634) /gb=AF0589	0.417	0.473	3 consistent probesets
M23379	HUMGAPA Human GTPase-activating protein ras p	0.415	0.079	8 inconsistent probesets
M27691	HUMCREB Human transactivator protein (CREB) m	0.413	0.212	5 consistent probesets
U34249	Human putative zinc finger protein (ZNFB7) mRNA, compl	0.413	0.152	4 consistent probesets
AB018273	Homo sapiens mRNA for KIAA0730 protein, partial cds /c	0.413	0.162	4 consistent probesets
D26488	Human mRNA for KIAA0007 gene, partial cds /cds=(0,2062	0.413	0.187	4 consistent probesets
AF027150	Homo sapiens survival of motor neuron protein interact	0.413	0.189	4 consistent probesets
AL050089	Homo sapiens mRNA; cDNA DKFZp586E0518 (from clone	0.413	0.199	4 consistent probesets
D13644	Human mRNA for KIAA0019 gene, complete cds /cds=(279	0.411	0.197	3 consistent probesets
AB028942	Homo sapiens mRNA for KIAA1019 protein, partial cds /c	0.411	0.432	3 consistent probesets
AL050190	Homo sapiens mRNA; cDNA DKFZp586B0923 (from clone	0.408	0.124	4 consistent probesets
M58286	HUMTNFRB Homo sapiens tumor necrosis factor r	0.408	0.137	4 consistent probesets
AB002374	Human mRNA for KIAA0376 gene, partial cds /cds=(0,2657	0.408	0.161	4 consistent probesets
L20320	Human protein serine/threonine kinase stk1 mRNA, compl	0.408	0.162	4 consistent probesets
AB028948	Homo sapiens mRNA for KIAA1025 protein, partial cds /c	0.408	0.207	4 consistent probesets
AB029001	Homo sapiens mRNA for KIAA1078 protein, partial cds /c	0.408	0.315	4 consistent probesets
M64929	HUMPROP2AA Human protein phosphatase 2A alpha	0.407	0.130	7 consistent probesets
U47414	HSU47414 Human cyclin G2 mRNA, complete cds"	0.403	0.213	5 consistent probesets
AL050141	Homo sapiens mRNA; cDNA DKFZp586O031 (from clone D	0.400	0.094	4 inconsistent probesets
AI655015	wb66a10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.400	0.101	1 consistent probesets
U08023	HSU08023 Human cellular proto-oncogene (c-mer	0.400	0.138	8 consistent probesets
AB011165	Homo sapiens mRNA for KIAA0593 protein, partial cds /c	0.400	0.178	4 consistent probesets
H10776	ym07h11.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-47	0.400	0.213	3 inconsistent probesets
AB011170	Homo sapiens mRNA for KIAA0598 protein, complete cds /	0.400	0.215	2 consistent probesets
AJ006591	Homo sapiens mRNA for cysteine-rich protein /cds=(180,	0.400	0.281	3 consistent probesets
AF043897	Homo sapiens C90RF3 large isoform, mRNA sequence /cds	0.400	0.308	2 consistent probesets
AI553745	tn28c11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-216	0.400	0.353	1 consistent probesets
U70370	Human hindlimb expressed homeobox protein backfoot (Bf	0.400	0.353	1 consistent probesets
Z22534	H.sapiens ALK-2 mRNA /cds=(103,1632) /gb=Z22534 /gi=4	0.400	0.356	3 consistent probesets
Z11773	Homo sapiens mRNA for SRE-ZBP /cds=(0,1226) /gb=Z117	0.400	0.358	2 consistent probesets
AL049229	Homo sapiens mRNA; cDNA DKFZp564O1016 (from clone	0.400	0.378	3 consistent probesets
X91868	H.sapiens mRNA for SIX1 protein /cds=(275,1129) /gb=X9	0.400	0.390	3 consistent probesets
AF097441	Homo sapiens phenylalanine-tRNA synthetase (FARS1) mR	0.400	0.403	1 consistent probesets
AF104304	Homo sapiens Smad anchor for receptor activation (SARA	0.400	0.419	2 consistent probesets
U65002	Human zinc finger protein PLAG1 mRNA, complete cds /cd	0.400	0.454	1 consistent probesets
W25798	13f12 Homo sapiens cDNA /gb=W25798 /gi=1305939 /	0.400	0.485	2 consistent probesets
AB028997	Homo sapiens mRNA for KIAA1074 protein, complete cds /	0.400	0.504	1 consistent probesets
L32164	Homo sapiens zinc finger protein mRNA, 3 end /cds=(0,	0.400	0.642	2 consistent probesets
AB014537	Homo sapiens mRNA for KIAA0637 protein, complete cds /	0.400	0.706	1 consistent probesets
AB023188	Homo sapiens mRNA for KIAA0971 protein, complete cds /	0.400	0.706	1 consistent probesets

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AB023183	Homo sapiens mRNA for KIAA0966 protein, complete cds /	0.400	0.756	2 consistent probesets
AF070594	Homo sapiens clone 24570 HNK-1 sulfotransferase mRNA,	0.400	0.857	1 consistent probesets
L76937	Homo sapiens Werner syndrome gene, complete cds /cds=	0.400	1.008	1 consistent probesets
L41067	HUMHFAT4A Homo sapiens NF-AT4c mRNA, complete	0.400	0.082	5 consistent probesets
M11058	Human 3-hydroxy-3-methylglutaryl coenzyme A reductase	0.396	0.114	4 consistent probesets
AB020669	Homo sapiens mRNA for KIAA0862 protein, complete cds /	0.396	0.116	4 consistent probesets
D38552	Human mRNA for KIAA0073 gene, partial cds /cds=(0,1938	0.396	0.142	4 consistent probesets
AA255502	zr85b06.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-682	0.396	0.146	4 consistent probesets
AF010607	HASMAD5S6 Homo sapiens SMAD5 (Smad5) gene, ex	0.396	0.178	4 inconsistent probesets
AI670788	tz10c02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-228	0.396	0.191	4 consistent probesets
X98001	HSGGII H.sapiens mRNA for geranylgeranyl tran	0.392	0.093	4 inconsistent probesets
AF055012	Homo sapiens clone 24615 mRNA sequence /cds=UNKNO	0.392	0.165	4 consistent probesets
AB023229	Homo sapiens mRNA for KIAA1012 protein, complete cds /	0.392	0.168	4 inconsistent probesets
X52425	HSIL4R Human IL-4-R mRNA for the interleukin	0.392	0.208	4 consistent probesets
X98834	H.sapiens mRNA for zinc finger protein, Hsal2 /cds=UNK	0.392	0.230	4 consistent probesets
U89278	Human polyhomeotic 2 homolog (HPH2) mRNA, complete (	0.388	0.095	4 consistent probesets
U53446	HSU53446 Human mitogen-responsive phosphoprot	0.388	0.129	4 inconsistent probesets
AL050050	Homo sapiens mRNA; cDNA DKFZp566D133 (from clone D	0.387	0.153	4 consistent probesets
U13695	HSU13695 Human homolog of yeast mutL (hPMS1)	0.385	0.106	12 inconsistent probesets
AB023169	Homo sapiens mRNA for KIAA0952 protein, complete cds /	0.383	0.108	4 consistent probesets
L10379	Human (clone CTG-B45d) mRNA sequence /cds=UNKNOW	0.383	0.143	4 consistent probesets
AA477576	zu44b03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-740	0.383	0.150	4 consistent probesets
M55422	Human Krueppel-related zinc finger protein (H-plk) mRN	0.383	0.175	4 consistent probesets
U84404	Human E6-associated protein E6-AP/ubiquitin-protein li	0.383	0.179	4 consistent probesets
Y09631	H.sapiens mRNA for PIBF1 protein, complete /cds=(0,227	0.383	0.190	4 consistent probesets
D87448	Human mRNA for KIAA0259 gene, partial cds /cds=(0,4653	0.383	0.228	3 consistent probesets
Z68747	H.sapiens mRNA for imogen 38 /cds=(5,1192) /gb=Z68747	0.383	0.235	4 consistent probesets
AL050081	Homo sapiens mRNA; cDNA DKFZp566J2146 (from clone	0.383	0.238	3 consistent probesets
U19969	Human two-handed zinc finger protein ZEB mRNA, partial	0.383	0.266	3 consistent probesets
AL049538	Human DNA sequence from clone 1175I6 on chromosome	0.383	0.282	3 consistent probesets
AL035291	H.sapiens gene from PACs 125H23 and 105D12 /cds=(219	0.383	0.304	3 consistent probesets
M74099	Human displacement protein (CCAAT) mRNA /cds=UNKNO	0.383	0.314	3 consistent probesets
M10901	HUMGCRA Human glucocorticoid receptor alpha m	0.380	0.187	5 consistent probesets
D63881	Human mRNA for KIAA0160 gene, partial cds /cds=(0,2413	0.379	0.204	4 consistent probesets
AB023164	Homo sapiens mRNA for KIAA0947 protein, partial cds /c	0.379	0.249	4 consistent probesets
Z36714	HSCYCLF H.sapiens mRNA for cyclin F	0.379	0.107	7 inconsistent probesets
AL049941	Homo sapiens mRNA; cDNA DKFZp564E2222 (from clone	0.375	0.112	4 consistent probesets
X16416	Human c-abl mRNA encoding p150 protein /cds=(147,3539	0.375	0.123	4 consistent probesets
AB007899	Homo sapiens KIAA0439 mRNA, partial cds /cds=(0,2989)	0.375	0.159	4 consistent probesets
AB007891	Homo sapiens KIAA0431 mRNA, partial cds /cds=(963,296	0.375	0.217	2 consistent probesets
R59697	yh11b03.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-42	0.375	0.221	4 consistent probesets
AA643063	nr95e11.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-117	0.375	0.257	4 consistent probesets
U68233	Human farnesol receptor HRR-1 (HRR-1) mRNA, complete	0.375	0.287	2 consistent probesets
AB018322	Homo sapiens mRNA for KIAA0779 protein, partial cds /c	0.375	0.429	2 consistent probesets
M91670	HUME2EPI Human ubiquitin carrier protein (E2-	0.374	0.066	12 inconsistent probesets
AB020694	Homo sapiens mRNA for KIAA0887 protein, partial cds /c	0.371	0.257	4 consistent probesets
AB018262	Homo sapiens mRNA for KIAA0719 protein, complete cds /	0.367	0.096	4 consistent probesets
L32832	Homo sapiens zinc finger homeodomain protein (ATBF1-A)	0.367	0.141	3 consistent probesets
X63547	H.sapiens mRNA for tre oncogene (clone 213) /cds=UNKNO	0.367	0.158	4 consistent probesets
AW005997	wz91c01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.367	0.179	4 consistent probesets
U37547	Human IAP homolog B (MIHB) mRNA, complete cds /cds=(	0.367	0.179	4 consistent probesets
M59371	HUMECK Human protein tyrosine kinase mRNA, co	0.367	0.234	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U90920	Human PTPL1-associated RhoGAP mRNA, complete cds /c	0.367	0.237	4 consistent probesets
U47742	HSU47742 Human monocytic leukaemia zinc finge	0.367	0.270	3 consistent probesets
X68487	H.sapiens mRNA for A2b adenosine receptor /cds=(332,13	0.367	0.303	3 consistent probesets
U43318	Human putative transmembrane receptor (frizzled 5) mRN	0.367	0.313	3 consistent probesets
X63417	H.sapiens irlB mRNA /cds=(0,575) /gb=X63417 /gi=33968	0.367	0.349	3 consistent probesets
U26424	Human Ste20-like kinase (MST2) mRNA, complete cds /cds	0.367	0.372	3 consistent probesets
AL080109	Homo sapiens mRNA; cDNA DKFZp586G1822 (from clone	0.363	0.082	4 consistent probesets
L14076	Human pre-mRNA splicing factor SRp75 mRNA, complete c	0.363	0.087	4 inconsistent probesets
D83032	Homo sapiens mRNA for nuclear protein, NP220, complete	0.363	0.120	4 inconsistent probesets
AB020633	Homo sapiens mRNA for KIAA0826 protein, partial cds /c	0.363	0.258	4 consistent probesets
X59244	Human ZNF43 mRNA /cds=(337,2748) /gb=X59244 /gi=380	0.363	0.274	4 consistent probesets
AF047438	Homo sapiens GOS28/P28 protein mRNA, complete cds /c	0.358	0.118	4 consistent probesets
D87078	Human mRNA for KIAA0235 gene, partial cds /cds=(1,2553	0.358	0.141	4 consistent probesets
Y08262	H.sapiens mRNA for SCA2 protein /cds=(0,2746) /gb=Y082	0.358	0.178	4 consistent probesets
U59863	Human TRAF-interacting protein I-TRAF mRNA, complete c	0.358	0.196	4 consistent probesets
X97324	H.sapiens mRNA for adipophilin /cds=(0,1313) /gb=X9732	0.354	0.090	4 consistent probesets
U19247	HSINFGRA7 Homo sapiens interferon-gamma recep	0.354	0.092	4 consistent probesets
W26477	30b5 Homo sapiens cDNA /gb=W26477 /gi=1307176 /	0.354	0.124	4 consistent probesets
M76766	HUMTFIIB Human transcription factor (TFIIB) m	0.354	0.145	4 consistent probesets
AB028949	Homo sapiens mRNA for KIAA1026 protein, partial cds /c	0.354	0.146	4 inconsistent probesets
X06409	Human mRNA fragment for activated c-raf-1 (exons 8-17)	0.354	0.154	4 consistent probesets
AJ224901	Homo sapiens mRNA for ZNF198 protein /cds=(184,4317) /	0.354	0.160	4 consistent probesets
AL096857	Novel human mRNA from chromosome 1, which has simila	0.350	0.098	4 consistent probesets
AB018328	Homo sapiens mRNA for KIAA0785 protein, complete cds /	0.350	0.111	4 consistent probesets
AF109134	Homo sapiens 7-60 mRNA, complete cds /cds=(205,2061) /	0.350	0.112	4 consistent probesets
AJ236876	Homo sapiens mRNA for poly(ADP-ribose) polymerase-2 /c	0.350	0.112	8 consistent probesets
U79252	Human clone 23679 mRNA, complete cds /cds=(973,1449)	0.350	0.136	4 consistent probesets
AB015019	Homo sapiens mRNA for BAP2-alpha protein, complete cds	0.350	0.140	4 consistent probesets
AF042378	Homo sapiens spindle pole body protein spc98 homolog G	0.350	0.145	4 consistent probesets
D80012	Human mRNA for KIAA0190 gene, partial cds /cds=(0,2442	0.350	0.149	4 consistent probesets
D90150	Human Gx-alpha gene /cds=(619,1686) /gb=D90150 /gi=21	0.350	0.159	3 consistent probesets
AF032886	Homo sapiens forkhead protein (FKHRL1) mRNA, complete	0.350	0.180	3 consistent probesets
AB007881	Homo sapiens KIAA0421 mRNA, partial cds /cds=(0,3908)	0.350	0.216	3 inconsistent probesets
M99578	Human lymphocyte surface protein exons 1-5, complete c	0.350	0.221	4 consistent probesets
X64116	H.sapiens PVR gene for poliovirus receptor (exon 1) /c	0.350	0.252	1 consistent probesets
D87076	Human mRNA for KIAA0239 gene, partial cds /cds=(0,1716	0.350	0.252	1 consistent probesets
U59321	Human DEAD-box protein p72 (P72) mRNA, complete cds ,	0.350	0.264	3 inconsistent probesets
AB014542	Homo sapiens mRNA for KIAA0642 protein, partial cds /c	0.350	0.266	3 consistent probesets
AB003698	AB003698 Homo sapiens mRNA for Cdc7-related k	0.350	0.288	3 consistent probesets
U50062	HSU50062 Homo sapiens RIP protein kinase mRNA	0.350	0.302	1 consistent probesets
U07000	HSU07000 Human breakpoint cluster region (BCR	0.350	0.302	1 consistent probesets
AF038203	Homo sapiens clone 23596 mRNA sequence /cds=UNKNO	0.350	0.302	1 consistent probesets
D86958	Human mRNA for KIAA0203 gene, complete cds /cds=(515	0.350	0.313	4 consistent probesets
AB021179	Homo sapiens mRNA for HEXIM1 protein, complete cds /c	0.350	0.316	2 consistent probesets
AB014549	Homo sapiens mRNA for KIAA0649 protein, complete cds /	0.350	0.353	1 consistent probesets
AI028290	ov84f11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-164	0.350	0.353	1 consistent probesets
AL050007	Homo sapiens mRNA; cDNA DKFZp564A043 (from clone D	0.350	0.363	2 consistent probesets
Y07595	H.sapiens mRNA for 52 kD subunit of transcription fact	0.350	0.376	3 consistent probesets
AB007928	Homo sapiens mRNA for KIAA0459 protein, partial cds /c	0.350	0.378	2 consistent probesets
X95826	H.sapiens ART4 gene /cds=(0,803) /gb=X95826 /gi=149542	0.350	0.417	2 consistent probesets
M74297	HUMHOX14 Human homeobox 1.4 protein mRNA, com	0.350	0.454	1 consistent probesets
U09410	Human zinc finger protein ZNF131 mRNA, partial cds /cd	0.350	0.454	1 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
M34667	HUMPLCA Human phospholipase C-gamma mRNA, com	0.350	0.463	2 consistent probesets
H53921	yq87g03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-202	0.350	0.510	2 consistent probesets
U79298	Human clone 23803 mRNA, partial cds /cds=(0,1123) /gb=	0.350	0.559	3 consistent probesets
AL050385	Homo sapiens mRNA; cDNA DKFZp564L2416 (from clone	0.350	0.706	1 consistent probesets
U13044	Human nuclear respiratory factor-2 subunit alpha mRNA,	0.350	0.756	1 consistent probesets
W27666	36a5 Homo sapiens cDNA /gb=W27666 /gi=1307614 /	0.350	0.756	1 consistent probesets
AB011102	Homo sapiens mRNA for KIAA0530 protein, partial cds /c	0.350	0.857	1 consistent probesets
D45132	HUMHOXY1 Homo sapiens mRNA for zinc-finger DN	0.350	1.058	1 consistent probesets
U28042	HSU28042 Human DEAD box RNA helicase-like pro	0.350	0.146	3 consistent probesets
AF034803	Homo sapiens liprin-beta2 mRNA, partial cds /cds=(0,23	0.350	0.175	3 consistent probesets
AB014540	Homo sapiens mRNA for KIAA0640 protein, partial cds /c	0.350	0.340	3 consistent probesets
AF002715	Homo sapiens MAP kinase kinase kinase (MTK1) mRNA, c	0.348	0.123	7 consistent probesets
U24169	Human JTV-1 (JTV-1) mRNA, complete cds /cds=(113,105	0.346	0.101	4 consistent probesets
Y08319	H.sapiens mRNA for kinesin-2 /cds=(18,2057) /gb=Y08319	0.346	0.105	4 inconsistent probesets
AB023198	Homo sapiens mRNA for KIAA0981 protein, partial cds /c	0.346	0.112	4 consistent probesets
AL080216	Homo sapiens mRNA; cDNA DKFZp586K1123 (from clone	0.346	0.120	4 consistent probesets
D87434	Human mRNA for KIAA0247 gene, complete cds /cds=(268	0.346	0.165	4 consistent probesets
AF064607	Homo sapiens GC20 protein mRNA, complete cds /cds=(70	0.346	0.168	4 consistent probesets
N98667	yy66d05.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-278	0.346	0.190	4 consistent probesets
AL049288	Homo sapiens mRNA; cDNA DKFZp564M053 (from clone D	0.342	0.072	8 inconsistent probesets
D83784	Human mRNA for KIAA0198 gene, partial cds /cds=(0,1690	0.342	0.085	4 inconsistent probesets
M23115	Homo sapiens calcium-ATPase (HK2) mRNA, complete cds	0.342	0.097	4 consistent probesets
AF042357	AF042357 Homo sapiens cDNA /gb=AF042357 /gi=410491	0.342	0.108	4 consistent probesets
AL022312	dJ1104E15.2 (activating transcription factor 4 (tax-re	0.342	0.118	4 inconsistent probesets
L13385	Homo sapiens(clone 71) Miller-Dieker lissencephaly pro	0.342	0.132	4 consistent probesets
AI341565	qq94g11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-19	0.342	0.140	4 consistent probesets
U49020	Human myocyte-specific enhancer factor 2A (MEF2A) gene	0.342	0.195	4 consistent probesets
AF053551	Homo sapiens metaxin 2 (MTX2) mRNA, nuclear gene enco	0.342	0.205	4 consistent probesets
D50927	Human mRNA for KIAA0137 gene, complete cds /cds=(108	0.338	0.081	4 consistent probesets
AB014547	Homo sapiens mRNA for KIAA0647 protein, partial cds /c	0.338	0.108	4 consistent probesets
M28215	HUMRAB5A Homo sapiens GTP-binding protein (RA	0.338	0.111	8 inconsistent probesets
Y09703	H.sapiens mRNA for MEMA protein /cds=(406,2166) /gb=Y(	0.338	0.133	4 consistent probesets
U14518	HSU14518 Human centromere protein-A (CENP-A)	0.338	0.134	4 consistent probesets
D83785	Human mRNA for KIAA0200 gene, complete cds /cds=(263	0.338	0.167	4 consistent probesets
AL035699	Human DNA sequence from clone 73H22 on chromosome (	0.338	0.175	4 consistent probesets
D25218	Human mRNA for KIAA0112 gene, partial cds /cds=(0,1201	0.338	0.175	4 consistent probesets
D13645	Human mRNA for KIAA0020 gene, complete cds /cds=(418	0.338	0.177	4 consistent probesets
AL049422	Homo sapiens mRNA; cDNA DKFZp586A191 (from clone D	0.338	0.185	4 consistent probesets
AF012086	Homo sapiens Ran binding protein 2 (RanBP2alpha) mRNA	0.338	0.201	4 consistent probesets
U31556	HSU31556 Human transcription factor E2F-5 mRN	0.336	0.127	7 consistent probesets
Y08201	Homo sapiens mRNA for rab geranylgeranyl transferase,	0.333	0.087	4 consistent probesets
AF017786	Homo sapiens phosphatidic acid phosphohydrolase homolo	0.333	0.104	4 inconsistent probesets
D14678	HUMMHCB Human mRNA for kinesin-related protei	0.333	0.144	5 consistent probesets
AB014533	Homo sapiens mRNA for KIAA0633 protein, partial cds /c	0.333	0.172	3 consistent probesets
AB023194	Homo sapiens mRNA for KIAA0977 protein, complete cds /	0.333	0.178	4 consistent probesets
AF051850	Homo sapiens supervillin mRNA, complete cds /cds=(450,	0.333	0.185	4 consistent probesets
U13897	Human homolog of Drosophila discs large protein, isofo	0.333	0.241	3 consistent probesets
X95263	H.sapiens mRNA for PWP2 protein /cds=(31,2790) /gb=X95	0.333	0.290	3 consistent probesets
AJ005814	Homo sapiens mRNA for hoxA7 protein /cds=(106,798) /gb	0.333	0.303	2 consistent probesets
AL050287	Homo sapiens mRNA; cDNA DKFZp586C021 (from clone D	0.333	0.325	3 consistent probesets
AL049987	Homo sapiens mRNA; cDNA DKFZp564F112 (from clone D	0.333	0.334	3 consistent probesets
U07223	Human beta2-chimaerin mRNA, complete cds /cds=(444,18	0.329	0.081	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
X12791	Human mRNA for 19kD protein of signal recognition part	0.329	0.121	4 consistent probesets
AB011118	Homo sapiens mRNA for KIAA0546 protein, partial cds /c	0.329	0.122	4 consistent probesets
D85418	D85418 Homo sapiens mRNA for phosphatidylinos	0.329	0.155	4 consistent probesets
AB028956	Homo sapiens mRNA for KIAA1033 protein, partial cds /c	0.329	0.164	4 consistent probesets
AL049944	Homo sapiens mRNA; cDNA DKFZp564G2022 (from clone	0.329	0.170	4 consistent probesets
D14659	Human mRNA for KIAA0103 gene, complete cds /cds=(6,8	0.329	0.181	4 consistent probesets
M34677	Human nested gene protein gene, complete cds /cds=(0,1	0.329	0.213	4 consistent probesets
AB002347	Human mRNA for KIAA0349 gene, partial cds /cds=(0,3827	0.328	0.284	3 consistent probesets
AF002020	AF002020 Homo sapiens Niemann-Pick C disease	0.327	0.092	8 consistent probesets
U17989	Homo sapiens nuclear autoantigen GS2NA mRNA, complet	0.325	0.111	4 consistent probesets
U08377	Human homolog of Drosophila splicing regulator suppres	0.325	0.169	2 consistent probesets
D80003	Human mRNA for KIAA0181 gene, partial cds /cds=(0,6019	0.325	0.171	4 consistent probesets
AL050261	Homo sapiens mRNA; cDNA DKFZp547E2110 (from clone	0.325	0.187	4 consistent probesets
AB029006	Homo sapiens mRNA for KIAA1083 protein, complete cds /	0.325	0.199	4 consistent probesets
AL049996	Homo sapiens mRNA; cDNA DKFZp564K112 (from clone D	0.325	0.217	4 consistent probesets
U26032	Human translation initiation factor eIF-2alpha mRNA, 3	0.325	0.220	4 consistent probesets
U57629	Human retinitis pigmentosa GTPase regulator (RPGR) mRN	0.325	0.265	4 consistent probesets
X16155	Human mRNA for chicken ovalbumin upstream promoter tra	0.325	0.268	2 consistent probesets
AJ012375	Homo sapiens mRNA for SUI1 protein translation initiat	0.325	0.101	4 inconsistent probesets
D86971	Human mRNA for KIAA0217 gene, partial cds /cds=(0,2021	0.325	0.158	4 consistent probesets
AB014525	Homo sapiens mRNA for KIAA0625 protein, partial cds /c	0.325	0.173	4 consistent probesets
AF061261	Homo sapiens zinc finger protein (MBLL) mRNA, complete	0.325	0.339	2 consistent probesets
AB017915	Homo sapiens mRNA for condoroitin 6-sulfotransferase,	0.322	0.256	3 consistent probesets
N36997	yy39g07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-27	0.321	0.081	4 consistent probesets
AF052174	Homo sapiens clone 24630 mRNA sequence /cds=UNKNO	0.321	0.136	4 consistent probesets
AB007858	Homo sapiens KIAA0398 mRNA, complete cds /cds=(196,1	0.321	0.137	4 consistent probesets
S79639	S79639 EXT1=putative tumour suppressor/heredi	0.321	0.152	8 consistent probesets
N42007	yw69e06.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-25	0.321	0.162	4 consistent probesets
AJ131693	Homo sapiens mRNA for AKAP450 protein /cds=(222,1194	0.321	0.195	4 consistent probesets
X76104	H.sapiens DAP-kinase mRNA /cds=(336,4631) /gb=X76104	0.321	0.226	4 consistent probesets
AF062346	Homo sapiens zinc finger protein 216 splice variant 1	0.321	0.241	4 consistent probesets
AI951046	wx62g06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.321	0.252	4 consistent probesets
AL080186	Homo sapiens mRNA; cDNA DKFZp564B0769 (from clone	0.321	0.342	4 consistent probesets
U68723	HSU68723 Human checkpoint suppressor 1 mRNA,	0.317	0.091	4 consistent probesets
AB018259	Homo sapiens mRNA for KIAA0716 protein, complete cds /	0.317	0.137	4 consistent probesets
AB018319	Homo sapiens mRNA for KIAA0776 protein, partial cds /c	0.317	0.139	4 consistent probesets
AL050371	Homo sapiens mRNA; cDNA DKFZp566G2246 (from clone	0.317	0.141	4 consistent probesets
U15173	Homo sapiens BCL2/adenovirus E1B 19kD-interacting prot	0.317	0.141	4 consistent probesets
AB007866	Homo sapiens KIAA0406 mRNA, complete cds /cds=(195,2	0.317	0.148	4 consistent probesets
L10910	Homo sapiens splicing factor (CC1.3) mRNA, complete cd	0.317	0.161	4 consistent probesets
AB023231	Homo sapiens mRNA for KIAA1014 protein, partial cds /c	0.317	0.163	4 consistent probesets
AI561196	tq27a01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-220	0.317	0.211	4 consistent probesets
AF055581	Homo sapiens adaptor protein Lnk mRNA, complete cds /c	0.317	0.213	3 consistent probesets
M11717	HUMHSP70D Human heat shock protein (hsp 70) g	0.317	0.215	3 consistent probesets
AI382123	te30a09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-208	0.317	0.252	4 inconsistent probesets
AF025840	Homo sapiens DNA polymerase epsilon subunit B (DPE2) m	0.317	0.302	3 consistent probesets
AL080215	Homo sapiens mRNA; cDNA DKFZp586J0323 (from clone	0.317	0.305	4 consistent probesets
AB018345	Homo sapiens mRNA for KIAA0802 protein, partial cds /c	0.317	0.311	3 consistent probesets
AB011090	Homo sapiens mRNA for KIAA0518 protein, partial cds /c	0.317	0.363	3 consistent probesets
AB014543	Homo sapiens mRNA for KIAA0643 protein, partial cds /c	0.317	0.372	3 consistent probesets
AB023235	Homo sapiens mRNA for KIAA1018 protein, complete cds /	0.317	0.381	3 consistent probesets
U58334	HSU58334 Human Bcl2, p53 binding protein Bbp/	0.317	0.386	3 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF001175	Homo sapiens ribonuclease P protein subunit p14 (Rpp14	0.317	0.539	3 consistent probesets
Z72499	H.sapiens mRNA for herpesvirus associated ubiquitin-sp	0.313	0.085	4 consistent probesets
U01038	Human pLK mRNA, complete cds /cds=(63,1874) /gb=U010	0.313	0.101	4 consistent probesets
D31762	Human mRNA for KIAA0057 gene, complete cds /cds=(75,1	0.313	0.116	4 consistent probesets
AB014527	Homo sapiens mRNA for KIAA0627 protein, partial cds /c	0.313	0.151	4 consistent probesets
X81889	H.sapiens mRNA for p0071 protein /cds=(141,3776) /gb=X	0.313	0.170	4 consistent probesets
U59912	Human chromosome 4 Mad homolog Smad1 mRNA, comp	0.313	0.228	4 consistent probesets
AF055024	Homo sapiens clone 24763 mRNA sequence /cds=UNKNO	0.311	0.288	3 consistent probesets
U85773	Human phosphomannomutase (PMM2) mRNA, complete c	0.308	0.079	4 inconsistent probesets
U90911	Human clone 23652 mRNA sequence /cds=UNKNOWN /gb	0.308	0.081	4 consistent probesets
Z24725	H.sapiens mitogen inducible gene mig-2, complete CDS /	0.308	0.100	4 consistent probesets
M87339	HUMACT1A Human replication factor C, 37-kDa s	0.308	0.111	8 consistent probesets
AB019494	Homo sapiens IDN3 mRNA, partial cds /cds=(706,7503) /g	0.308	0.192	4 consistent probesets
X98260	H.sapiens mRNA for M-phase phosphoprotein, mpp11 /cds=	0.308	0.217	4 consistent probesets
X59841	Human PBX3 mRNA /cds=UNKNOWN /gb=X59841 /gi=353	0.308	0.222	4 consistent probesets
Y17829	Homo sapiens mRNA for Homer-related protein Syn47 /cds	0.308	0.259	4 consistent probesets
M30704	Human amphiregulin (AR) mRNA, complete cds, clones lan	0.306	0.246	3 consistent probesets
M59830	Human MHC class III HSP70-2 gene (HLA), complete cds /	0.304	0.145	4 consistent probesets
AF107463	Homo sapiens splicing factor mRNA, complete cds /cds=(	0.304	0.183	4 consistent probesets
AA595596	nk92d08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-10	0.304	0.223	4 consistent probesets
M97388	HUMDR1TATA Human TATA binding protein-associa	0.302	0.130	7 consistent probesets
X98172	H.sapiens mRNA for MACH-alpha-1 protein /cds=(291,1730	0.300	0.122	4 inconsistent probesets
U29332	Homo sapiens heart protein (FHL-2) mRNA, complete cds	0.300	0.124	4 consistent probesets
U77735	HSU77735 Human pim-2 protooncogene homolog pi	0.300	0.130	4 consistent probesets
D10704	Human mRNA for choline kinase /cds=(27,1397) /gb=D107	0.300	0.132	2 consistent probesets
AI184802	qd24g04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-17	0.300	0.133	4 consistent probesets
AL049404	Homo sapiens mRNA; cDNA DKFZp586F0219 (from clone	0.300	0.136	3 consistent probesets
AL080115	Homo sapiens mRNA; cDNA DKFZp564G0222 (from clone	0.300	0.137	4 consistent probesets
AI796944	we25b03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.300	0.154	4 consistent probesets
D86982	Human mRNA for KIAA0229 gene, partial cds /cds=(0,3543	0.300	0.156	4 consistent probesets
AB023216	Homo sapiens mRNA for KIAA0999 protein, partial cds /c	0.300	0.186	3 consistent probesets
AF112972	Homo sapiens TJ6 mRNA, complete cds /cds=(5,2575) /gb=	0.300	0.190	4 consistent probesets
Z35491	H.sapiens mRNA for novel glucocorticoid receptor-assoc	0.300	0.197	2 consistent probesets
AB014591	Homo sapiens mRNA for KIAA0691 protein, complete cds /	0.300	0.205	3 consistent probesets
AF041381	Homo sapiens putative transcriptional repressor E2F-6	0.300	0.270	3 consistent probesets
AB028639	Homo sapiens CAPN7 mRNA for PalBH, complete cds /cds	0.300	0.293	3 consistent probesets
S76638	S76638 p50-NF-kappa B homolog [human, periphe	0.300	0.302	1 consistent probesets
AF038182	Homo sapiens clone 23860 mRNA sequence /cds=UNKNO	0.300	0.365	3 consistent probesets
AL080198	Homo sapiens mRNA; cDNA DKFZp434D222 (from clone D	0.300	0.372	2 consistent probesets
L35251	Homo sapiens extracellular matrix protein (MFAP3) gene	0.300	0.375	2 consistent probesets
Y08991	H.sapiens mRNA for adaptor protein p150 /cds=(567,4643	0.300	0.382	3 consistent probesets
AL109722	Homo sapiens mRNA full length insert cDNA clone EUROIN	0.300	0.403	1 consistent probesets
L36529	Human (clone N5-4) protein p84 mRNA, complete cds /cds	0.300	0.403	1 consistent probesets
AF017418	Homo sapiens homeobox protein MEIS2 (MEIS2) mRNA, p	0.300	0.407	3 consistent probesets
AF055006	Homo sapiens clone 24666 sec6 homolog mRNA, partial co	0.300	0.454	1 consistent probesets
Y11997	H.sapiens mRNA for A-kinase anchoring protein AKAP95 /	0.300	0.454	1 consistent probesets
W37606	zc12a03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-322	0.300	0.496	3 consistent probesets
L40399	Homo sapiens (clone S240ii117/zap112) mRNA, complete c	0.300	0.504	1 consistent probesets
U45974	Human phosphatidylinositol (4,5) bisphosphate 5-phosph	0.300	0.504	1 consistent probesets
U92981	Homo sapiens clone DT1P1B6 mRNA, CAG repeat region /	0.300	0.504	1 consistent probesets
W27545	32c4 Homo sapiens cDNA /gb=W27545 /gi=1307349 /	0.300	0.553	2 consistent probesets
AJ133115	Homo sapiens mRNA for TSC-22-like protein /cds=(340,15	0.300	0.605	1 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AB016898	Homo sapiens HGC6.4 mRNA, complete cds /cds=(254,101	0.300	0.646	2 consistent probesets
AJ009770	Homo sapiens mRNA for putative transcription factor, p	0.300	0.706	1 consistent probesets
U29671	HSU29671 Human MEK kinase (Mekk) gene, partia	0.300	0.706	1 consistent probesets
AF077954	Homo sapiens protein inhibitor of activated STAT prote	0.300	0.706	1 consistent probesets
AB011422	Homo sapiens mRNA for Trad, complete cds /cds=(117,398	0.300	0.706	1 consistent probesets
L39060	Homo sapiens transcription factor SL1 mRNA, complete c	0.300	0.756	1 consistent probesets
U06454	HSU06454 Human AMP-activated protein kinase (	0.300	0.756	1 consistent probesets
AA442799	zv69b10.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-758	0.296	0.105	4 consistent probesets
U15782	Human cleavage stimulation factor 77kDa subunit mRNA,	0.296	0.107	4 consistent probesets
U80760	Homo sapiens CAGH1 alternate open reading frame mRNA	0.296	0.128	4 consistent probesets
U19523	Human GTP cyclohydrolase I mRNA, complete cds /cds=(1	0.296	0.154	4 consistent probesets
AJ011679	Homo sapiens mRNA for Rab6 GTPase activating protein /	0.296	0.158	4 consistent probesets
D21089	HUMXPCR Human mRNA for XP-C repair complement	0.296	0.195	4 consistent probesets
AB015051	Homo sapiens mRNA for Daxx, complete cds /cds=(112,23	0.296	0.196	4 consistent probesets
AB026118	Homo sapiens mRNA for MALT1, complete cds /cds=(65,25	0.296	0.199	4 consistent probesets
U13737	Human cysteine protease CPP32 isoform alpha mRNA, cor	0.296	0.230	4 consistent probesets
U13896	Human homolog of Drosophila discs large protein, isofo	0.296	0.236	4 consistent probesets
Y11312	H.sapiens mRNA for phosphoinositide 3-kinase /cds=(515	0.296	0.246	4 consistent probesets
L25441	HUMGGTBS Human geranylgeranyltransferase type	0.296	0.285	4 consistent probesets
U07563	HSABLGR3 Human proto-oncogene tyrosine-protei	0.292	0.079	12 inconsistent probesets
AB003103	AB003103 Homo sapiens mRNA for 26S proteasome	0.292	0.124	4 consistent probesets
AB011149	Homo sapiens mRNA for KIAA0577 protein, complete cds /	0.292	0.150	4 consistent probesets
X95525	H.sapiens mRNA for TAFII100 protein /cds=(23,2422) /gb	0.292	0.166	4 consistent probesets
AB014563	Homo sapiens mRNA for KIAA0663 protein, complete cds /	0.292	0.185	4 consistent probesets
U15306	Human cysteine-rich sequence-specific DNA-binding prot	0.292	0.193	4 consistent probesets
AF049460	Homo sapiens nuclear DEAF-1 related transcriptional re	0.292	0.196	4 consistent probesets
AB023153	Homo sapiens mRNA for KIAA0936 protein, complete cds /	0.292	0.220	4 consistent probesets
N95168	zb55f11.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-307	0.289	0.425	3 consistent probesets
AA977580	on61b02.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-15	0.288	0.085	4 consistent probesets
AF062536	Homo sapiens cullin 1 mRNA, complete cds /cds=(124,245	0.288	0.112	4 consistent probesets
AB007619	Homo sapiens mRNA for EBAG9, complete cds /cds=(283,9	0.288	0.112	4 consistent probesets
U61145	Human enhancer of zeste homolog 2 (EZH2) mRNA, compl	0.288	0.117	4 consistent probesets
M29960	HUMTR211 Human steroid receptor (TR2-11) mRNA	0.288	0.154	4 consistent probesets
U27459	Human origin recognition complex protein 2 homolog hOR	0.288	0.158	4 consistent probesets
L06147	Human (clone SY11) golgin-95 mRNA, complete cds /cds=(	0.288	0.158	4 consistent probesets
D29677	Human mRNA for KIAA0054 gene, complete cds /cds=(145	0.288	0.170	4 inconsistent probesets
M34065	HUMCDC25HS Human cdc25Hs mRNA, complete cds"	0.288	0.199	4 consistent probesets
AB025254	Homo sapiens mRNA for tudor repeat associator with PCT	0.288	0.233	4 consistent probesets
X56681	HSJUNDR Human junD mRNA	0.287	0.054	9 inconsistent probesets
X99720	H.sapiens TPRC gene /cds=(212,1687) /gb=X99720 /gi=186	0.283	0.147	3 consistent probesets
X89985	H.sapiens mRNA for BCL7B protein /cds=(37,645) /gb=X89	0.283	0.150	4 consistent probesets
AJ009771	Homo sapiens mRNA for putative RING finger protein, pa	0.283	0.154	4 consistent probesets
AF070528	Homo sapiens clone 24631 mRNA sequence /cds=UNKNO	0.283	0.163	4 consistent probesets
AB023137	Homo sapiens mRNA for KIAA0920 protein, complete cds /	0.283	0.208	3 consistent probesets
AI337901	qt34f05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-194	0.283	0.220	3 consistent probesets
AF016370	Homo sapiens U4/U6 small nuclear ribonucleoprotein hPr	0.283	0.232	4 consistent probesets
AF022116	Homo sapiens 5-AMP-activated protein kinase beta-1 mRN	0.283	0.263	3 consistent probesets
D79997	Human mRNA for KIAA0175 gene, complete cds /cds=(170	0.283	0.268	3 consistent probesets
AJ010840	Homo sapiens mRNA for ATP-dependent RNA helicase, pa	0.283	0.311	3 consistent probesets
AF006010	Human progestin induced protein (DD5) mRNA, complete c	0.281	0.134	8 consistent probesets
AI254524	qv48f07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-198	0.281	0.179	7 consistent probesets
M93425	HUMPTPPEST Human protein tyrosine phosphatase	0.279	0.087	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
M80627	Human HEB helix-loop-helix protein (HEB) mRNA, complet	0.279	0.127	4 consistent probesets
W27129	22g8 Homo sapiens cDNA /gb=W27129 /gi=1306663 /	0.279	0.133	4 consistent probesets
U77949	HSU77949 Human Cdc6-related protein (HsCDC6)	0.279	0.171	8 inconsistent probesets
AL049341	Homo sapiens mRNA; cDNA DKFZp566A163 (from clone D	0.279	0.322	4 inconsistent probesets
U09584	Human PL6 protein (PL6) mRNA, complete cds /cds=(211,1	0.275	0.104	4 consistent probesets
Y12065	Homo sapiens mRNA for nucleolar protein hNop56 /cds=(2	0.275	0.107	4 consistent probesets
L35263	Human CSaids binding protein (CSBP1) mRNA, complete c	0.275	0.116	4 consistent probesets
AB007931	Homo sapiens mRNA for KIAA0462 protein, partial cds /c	0.275	0.128	4 consistent probesets
U79266	Human clone 23627 mRNA, complete cds /cds=(184,1041)	0.275	0.141	4 consistent probesets
U90919	Human clones 23667 and 23775 zinc finger protein mRNA,	0.275	0.151	4 consistent probesets
D42045	Human mRNA for KIAA0086 gene, complete cds /cds=(918	0.275	0.254	4 consistent probesets
AB018337	Homo sapiens mRNA for KIAA0794 protein, partial cds /c	0.275	0.333	2 consistent probesets
AB028957	Homo sapiens mRNA for KIAA1034 protein, partial cds /c	0.275	0.413	2 consistent probesets
X89984	H.sapiens mRNA for BCL7A protein /cds=(953,1648) /gb=X	0.275	0.478	2 consistent probesets
U44798	Human U1-snRNP binding protein homolog mRNA, comple	0.275	0.665	2 consistent probesets
D86988	Human mRNA for KIAA0221 gene, complete cds /cds=(209	0.275	0.093	4 consistent probesets
AJ010346	Homo sapiens mRNA for RING-H2 protein RNF6, alternativ	0.275	0.131	4 consistent probesets
U79267	Human clone 23840 mRNA, partial cds /cds=(0,521) /gb=U	0.275	0.283	4 consistent probesets
L13286	HUMDHVH Human mitochondrial 1,25-dihydroxyvit	0.272	0.170	3 consistent probesets
L20046	HUMERCC5A Human ERCC5 excision repair protein	0.271	0.193	7 consistent probesets
AJ000644	Homo sapiens mRNA for SPOP /cds=(157,1281) /gb=AJ00	0.271	0.102	4 consistent probesets
M19481	Human follistatin gene /cds=(0,953) /gb=M19481 /gi=182	0.271	0.113	4 inconsistent probesets
AL030996	dJ1189B24.4 (novel PUTATIVE protein similar to hypothe	0.271	0.116	4 inconsistent probesets
D83776	Human mRNA for KIAA0191 gene, partial cds /cds=(0,4552	0.271	0.117	4 consistent probesets
AF075587	Homo sapiens protein associated with Myc mRNA, complet	0.271	0.125	4 consistent probesets
X03484	HSRAFR Human mRNA for raf oncogene	0.271	0.125	4 consistent probesets
AI656421	tt50h10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-224	0.271	0.148	4 consistent probesets
AB018271	Homo sapiens mRNA for KIAA0728 protein, partial cds /c	0.271	0.149	4 consistent probesets
M85169	Human homologue of yeast sec7 mRNA, complete cds /cds	0.271	0.169	4 consistent probesets
AI884738	wl84b02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.271	0.207	4 consistent probesets
M31516	Human decay-accelerating factor mRNA, complete cds /cd	0.271	0.228	4 consistent probesets
U88629	HSU88629 Human RNA polymerase II elongation f	0.269	0.110	8 inconsistent probesets
U04209	Human associated microfibrillar protein mRNA, complete	0.267	0.098	4 consistent probesets
X63753	H.sapiens son-a mRNA /cds=(414,4985) /gb=X63753 /gi=36	0.267	0.101	4 consistent probesets
AB029020	Homo sapiens mRNA for KIAA1097 protein, partial cds /c	0.267	0.112	4 consistent probesets
X97335	H.sapiens mRNA for kinase A anchor protein /cds=(124,2	0.267	0.114	4 consistent probesets
AB020714	Homo sapiens mRNA for KIAA0907 protein, complete cds /	0.267	0.118	4 consistent probesets
U18062	HSU18062 Human TFIID subunit TAFII55 (TAFII55	0.267	0.144	4 consistent probesets
D50922	Human mRNA for KIAA0132 gene, complete cds /cds=(112	0.267	0.148	3 consistent probesets
U00238	Homo sapiens glutamine PRPP amidotransferase (GPAT) r	0.267	0.159	3 inconsistent probesets
S81752	DPH2L=candidate tumor suppressor gene {ovarian cancer	0.267	0.163	3 consistent probesets
U94836	Human ERPROT 213-21 mRNA, complete cds /cds=(88,27	0.267	0.163	3 consistent probesets
S66431	S66431 RBP2=retinoblastoma binding protein 2	0.267	0.168	4 consistent probesets
AB019987	Homo sapiens mRNA for chromosome-associated polypept	0.267	0.177	4 consistent probesets
AB002330	Human mRNA for KIAA0332 gene, partial cds /cds=(0,3087	0.267	0.194	4 consistent probesets
D50487	HUMHRH1 Human mRNA for RNA helicase (HRH1), c	0.267	0.194	3 consistent probesets
AF070582	Homo sapiens clone 24766 mRNA sequence /cds=UNKNO	0.267	0.200	3 consistent probesets
D87454	Human mRNA for KIAA0265 gene, partial cds /cds=(0,1205	0.267	0.219	4 consistent probesets
U78575	Human 68 kDa type I phosphatidylinositol-4-phosphate 5	0.267	0.239	4 consistent probesets
AB002357	Human mRNA for KIAA0359 gene, complete cds /cds=(167	0.267	0.241	3 consistent probesets
D87440	Human mRNA for KIAA0252 gene, partial cds /cds=(0,1994	0.267	0.266	3 consistent probesets
AB011139	Homo sapiens mRNA for KIAA0567 protein, partial cds /c	0.267	0.286	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D11958	HUM0S11H12 Homo sapiens cDNA /clone=s11h12 /clone_	0.267	0.304	3 consistent probesets
D87455	Human mRNA for KIAA0266 gene, complete cds /cds=(733	0.267	0.305	3 consistent probesets
AB023139	Homo sapiens mRNA for KIAA0922 protein, partial cds /c	0.267	0.316	3 consistent probesets
D14533	HUMXPAC Human mRNA for XPAC protein	0.264	0.152	7 consistent probesets
W29087	56b8 Homo sapiens cDNA /gb=W29087 /gi=1309053 /	0.263	0.067	4 consistent probesets
AF016266	Homo sapiens TRAIL receptor 2 mRNA, complete cds /cds=	0.263	0.105	4 consistent probesets
AF017445	Homo sapiens GDP-L-fucose pyrophosphorylase (GFPP) m	0.263	0.131	4 consistent probesets
AF053306	Homo sapiens mitotic checkpoint kinase Mad3L (MAD3L) m	0.263	0.133	4 consistent probesets
U61836	HSU61836 Human putative cyclin G1 interacting	0.263	0.146	4 consistent probesets
D50930	Human mRNA for KIAA0140 gene, complete cds /cds=(212	0.263	0.199	4 consistent probesets
U89387	Human RNA polymerase II subunit hsRPB4 gene, complete	0.262	0.149	4 consistent probesets
Z24724	H.sapiens polyA site DNA /cds=UNKNOWN /gb=Z24724 /g	0.262	0.162	4 consistent probesets
AB017546	Homo sapiens Pex14 mRNA for peroxisomal membrane an	0.262	0.194	4 consistent probesets
AF038009	Homo sapiens tyrosylprotein sulfotransferase-1 mRNA, c	0.262	0.259	4 consistent probesets
X51688	HSCYCLINA Human mRNA for cyclin A	0.258	0.070	8 inconsistent probesets
AF007149	Homo sapiens clone 23568, 23621, 23795, 23873 and 2387	0.258	0.102	4 consistent probesets
U58087	Human Hs-cul-1 mRNA, complete cds /cds=(124,2382) /gb=	0.258	0.116	4 inconsistent probesets
L13616	Human focal adhesion kinase (FAK) mRNA, complete cds /	0.258	0.130	4 consistent probesets
AB014599	Homo sapiens mRNA for KIAA0699 protein, partial cds /c	0.258	0.139	4 consistent probesets
L35013	Human spliceosomal protein (SAP 49) gene, complete cds	0.258	0.149	4 consistent probesets
AL049389	Homo sapiens mRNA; cDNA DKFZp586O0118 (from clone	0.258	0.157	4 consistent probesets
D29956	Human mRNA for KIAA0055 gene, complete cds /cds=(317	0.258	0.163	4 consistent probesets
L08488	HUMINOS Human inositol polyphosphate 1-phosph	0.258	0.169	6 consistent probesets
Z75331	H.sapiens mRNA for nuclear protein SA-2 /cds=(649,4137	0.258	0.176	4 consistent probesets
AF074606	Homo sapiens histone acetyltransferase (HBO1) mRNA, co	0.258	0.179	4 consistent probesets
AB011164	Homo sapiens mRNA for KIAA0592 protein, partial cds /c	0.258	0.180	4 consistent probesets
D83702	Homo sapiens mRNA for photolyase, complete cds /cds=(5	0.258	0.182	4 consistent probesets
AB011125	Homo sapiens mRNA for KIAA0553 protein, partial cds /c	0.258	0.198	4 consistent probesets
Y09723	H.sapiens mRNA for Miz-1 protein /cds=(126,2537) /gb=Y	0.258	0.243	4 consistent probesets
AB014562	Homo sapiens mRNA for KIAA0662 protein, partial cds /c	0.258	0.245	4 consistent probesets
U65416	Human MHC class I molecule (MICB) gene, complete cds /	0.254	0.058	4 inconsistent probesets
Y08614	Homo sapiens mRNA for CRM1 protein /cds=(38,3253) /gb=	0.254	0.072	4 consistent probesets
M14083	Human beta-migrating plasminogen activator inhibitor I	0.254	0.078	4 inconsistent probesets
AF072810	Homo sapiens transcription factor WSTF mRNA, complete	0.254	0.107	4 consistent probesets
AI688299	wc87h10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.254	0.109	4 consistent probesets
X68836	H.sapiens mRNA for S-adenosylmethionine synthetase /cd	0.254	0.118	4 inconsistent probesets
AB007890	Homo sapiens KIAA0430 mRNA, complete cds /cds=(0,317	0.254	0.121	4 consistent probesets
AF017789	Homo sapiens putative transcription factor CA150 mRNA,	0.254	0.122	4 consistent probesets
AL043470	DKFZp434A0327_s1 Homo sapiens cDNA, 3 end /clone=D	0.254	0.142	4 consistent probesets
AB022785	Homo sapiens ASH2L gene, complete cds, similar to Dros	0.254	0.144	4 consistent probesets
U20979	Human chromatin assembly factor-I p150 subunit mRNA, c	0.254	0.205	4 consistent probesets
X83490	HSFAS34 H.sapiens mRNA for Fas/Apo-1 (clone p	0.254	0.247	4 consistent probesets
AB023196	Homo sapiens mRNA for KIAA0979 protein, partial cds /c	0.254	0.289	4 consistent probesets
U43286	Human selenophosphate synthetase 2 (SPS2) mRNA, com	0.250	0.072	4 inconsistent probesets
U45976	Human clathrin assembly protein lymphoid myeloid leuke	0.250	0.082	4 consistent probesets
AB018276	Homo sapiens mRNA for KIAA0733 protein, partial cds /c	0.250	0.093	4 consistent probesets
AB002321	Human mRNA for KIAA0323 gene, partial cds /cds=(0,2175	0.250	0.102	4 consistent probesets
D79996	Human mRNA for KIAA0174 gene, complete cds /cds=(63,1	0.250	0.108	4 consistent probesets
D87446	Human mRNA for KIAA0257 gene, partial cds /cds=(0,5418	0.250	0.111	4 consistent probesets
X85753	HSCDK8 Homo sapiens mRNA for CDK8 protein kin	0.250	0.122	4 consistent probesets
L41887	Homo sapiens splicing factor, arginine/serine-rich 7 (	0.250	0.130	4 consistent probesets
M18533	Homo sapiens dystrophin (DMD) mRNA, complete cds /cds	0.250	0.139	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF038966	Homo sapiens secretory carrier-associated membrane pro	0.250	0.143	4 inconsistent probesets
D63997	Homo sapiens mRNA for GCP170, complete cds /cds=(269	0.250	0.147	4 consistent probesets
U53204	Human plectin (PLEC1) mRNA, complete cds /cds=(51,137	0.250	0.151	1 consistent probesets
AB006630	Homo sapiens mRNA for KIAA0292 gene, partial cds /cds=	0.250	0.176	3 consistent probesets
AI829890	wj47a06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.250	0.185	3 consistent probesets
AL033538	Human DNA sequence from clone 477H23 on chromosome	0.250	0.186	3 consistent probesets
AB029017	Homo sapiens mRNA for KIAA1094 protein, complete cds /	0.250	0.197	3 consistent probesets
AB006179	Homo sapiens mRNA for heparan-sulfate 6-sulfotransfera	0.250	0.202	1 consistent probesets
AB028978	Homo sapiens mRNA for KIAA1055 protein, partial cds /c	0.250	0.202	1 consistent probesets
AB020735	Homo sapiens ENDOGL-2 (alias ENGL-b) mRNA for endon	0.250	0.202	1 consistent probesets
M91083	Human DNA-binding protein (HRC1) mRNA, complete cds /	0.250	0.203	2 consistent probesets
AF064606	Homo sapiens KB07 protein mRNA, partial cds /cds=(0,73	0.250	0.205	3 consistent probesets
AB014535	Homo sapiens mRNA for KIAA0635 protein, complete cds /	0.250	0.215	3 consistent probesets
X92841	H.sapiens MICA gene /cds=(39,1196) /gb=X92841 /gi=1405	0.250	0.216	3 consistent probesets
AB011113	Homo sapiens mRNA for KIAA0541 protein, partial cds /c	0.250	0.217	2 consistent probesets
AB011089	Homo sapiens mRNA for KIAA0517 protein, partial cds /c	0.250	0.220	4 consistent probesets
AL080214	Homo sapiens mRNA; cDNA DKFZp586I2223 (from clone D	0.250	0.231	3 consistent probesets
U32645	Human myeloid elf-1 like factor (MEF) mRNA, complete c	0.250	0.234	3 consistent probesets
AF069987	Homo sapiens nitrilase 1 (NIT1) mRNA, complete cds /cd	0.250	0.235	3 consistent probesets
AB002308	Human mRNA for KIAA0310 gene, complete cds /cds=(165	0.250	0.238	2 consistent probesets
W28205	43g5 Homo sapiens cDNA /gb=W28205 /gi=1308171 /	0.250	0.252	1 consistent probesets
X72790	Human endogenous retrovirus mRNA for ORF /cds=(524,85	0.250	0.252	1 consistent probesets
X97675	H.sapiens mRNA for plakophilin 2a and b /cds=(25,2670)	0.250	0.257	3 consistent probesets
AI761647	wg66h09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.250	0.294	2 consistent probesets
AF098462	Homo sapiens stanniocalcin-related protein mRNA, compl	0.250	0.301	3 consistent probesets
AF038176	Homo sapiens clone 23859 mRNA sequence /cds=UNKNO	0.250	0.302	1 consistent probesets
AI141670	ot08b12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-161	0.250	0.302	1 consistent probesets
U94703	Homo sapiens mitochondrial DNA polymerase accessory su	0.250	0.314	3 consistent probesets
U31501	Human fragile X mental retardation syndrome related pr	0.250	0.338	2 consistent probesets
T53900	yb83e08.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-77	0.250	0.343	2 consistent probesets
U15172	Homo sapiens BCL2/adenovirus E1B 19kD-interacting prot	0.250	0.343	2 consistent probesets
X54938	HSHIP3K Human mRNA for inositol 1,4,5-triphos	0.250	0.353	1 consistent probesets
S71018	cyclophilin C [human, kidney, mRNA, 883 nt] /cds=(164,	0.250	0.385	2 consistent probesets
AB007892	Homo sapiens KIAA0432 mRNA, complete cds /cds=(0,225	0.250	0.398	2 consistent probesets
AF074264	Homo sapiens LDL receptor-related protein 6 (LRP6) mRN	0.250	0.398	2 consistent probesets
L40904	H. sapiens peroxisome proliferator activated receptor	0.250	0.403	1 consistent probesets
M15025	HUMBCRABL Human BCR/ABL mRNA (product of tran	0.250	0.403	1 consistent probesets
AB014589	Homo sapiens mRNA for KIAA0689 protein, partial cds /c	0.250	0.403	1 consistent probesets
X63717	H.sapiens mRNA for APO-1 cell surface antigen /cds=(22	0.250	0.454	1 consistent probesets
U72621	Human LOT1 mRNA, complete cds /cds=(657,2048) /gb=U	0.250	0.454	1 consistent probesets
L01042	Human HIV1 tata element modulatory factor mRNA sequen	0.250	0.500	2 consistent probesets
AI525962	DU145-2.B11.r Homo sapiens cDNA, 5 end /clone_end=5	0.250	0.552	3 consistent probesets
AL049280	Homo sapiens mRNA; cDNA DKFZp564K143 (from clone D	0.250	0.554	1 consistent probesets
M61199	Human cleavage signal 1 protein mRNA, complete cds /cd	0.250	0.566	2 consistent probesets
AI742087	wg38g10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.250	0.683	1 consistent probesets
AL080140	Homo sapiens mRNA; cDNA DKFZp434L243 (from clone D	0.250	0.806	1 consistent probesets
D13317	Human mRNA for transcription factor, E4TF1-53, complet	0.250	0.907	1 consistent probesets
U00957	Human clone KDB1.2 (CAC)n/(GTG)n repeat-containing mR	0.250	1.058	1 consistent probesets
U89358	Human l(3)mbt protein homolog mRNA, complete cds /cds=	0.250	1.224	1 consistent probesets
AW024285	wt69d06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.250	0.113	4 consistent probesets
D80009	Human mRNA for KIAA0187 gene, complete cds /cds=(27,3	0.250	0.144	4 consistent probesets
U73379	HSU73379 Human cyclin-selective ubiquitin car	0.246	0.085	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AI808712	wf57c05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.246	0.094	4 consistent probesets
AL080113	Homo sapiens mRNA; cDNA DKFZp586K2322 (from clone	0.246	0.107	4 consistent probesets
AB020658	Homo sapiens mRNA for KIAA0851 protein, complete cds /	0.246	0.113	4 consistent probesets
U41315	Human ring zinc-finger protein (ZNF127-Xp) gene and 5	0.246	0.122	4 consistent probesets
AA447263	zw93f01.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-784	0.246	0.129	4 consistent probesets
Y13467	Homo sapiens mRNA for RB18A protein /cds=(235,4935) /g	0.246	0.131	4 consistent probesets
AI040324	oy33a12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.246	0.132	4 consistent probesets
M88338	Human serum constituent protein (MSE55) mRNA, complet	0.246	0.134	4 consistent probesets
AL031685	dJ963K23.2 (novel protein) /cds=(2,688) /gb=AL031685 /	0.246	0.140	4 consistent probesets
AB011166	Homo sapiens mRNA for KIAA0594 protein, partial cds /c	0.246	0.141	4 consistent probesets
M54968	HUMKRASM Human K-ras oncogene protein mRNA, c	0.246	0.146	4 consistent probesets
AA630312	ac08f05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-855	0.246	0.153	4 consistent probesets
U49844	Human FRAP-related protein (FRP1) mRNA, complete cds	0.246	0.174	4 consistent probesets
AF046024	Homo sapiens UBA3 (UBA3) mRNA, complete cds /cds=(84	0.246	0.209	4 consistent probesets
Z22535	H.sapiens ALK-3 mRNA /cds=(309,1907) /gb=Z22535 /gi=4	0.246	0.288	4 consistent probesets
W26655	34c9 Homo sapiens cDNA /gb=W26655 /gi=1307498 /	0.242	0.070	4 consistent probesets
AI535653	P9-C4.T3.P9.D4 Homo sapiens cDNA, 3 end /clone_end=3	0.242	0.086	4 consistent probesets
D21853	Human mRNA for KIAA0111 gene, complete cds /cds=(214	0.242	0.089	4 consistent probesets
AB007885	Homo sapiens KIAA0425 mRNA, complete cds /cds=(172,3	0.242	0.104	4 consistent probesets
AF036927	Homo sapiens adenylyl cyclase type IX mRNA, complete c	0.242	0.105	4 consistent probesets
AB029022	Homo sapiens mRNA for KIAA1099 protein, complete cds /	0.242	0.111	4 consistent probesets
AJ002030	Homo sapiens mRNA for putative progesterone binding pr	0.242	0.125	4 consistent probesets
AF051160	Homo sapiens tyrosine phosphatase (PRL-1) gene, comple	0.242	0.133	4 consistent probesets
Z46376	H.sapiens HK2 mRNA for hexokinase II /cds=(1490,4243)	0.242	0.136	4 consistent probesets
Z34975	H.sapiens LDLC mRNA /cds=(95,2311) /gb=Z34975 /gi=575	0.242	0.139	4 consistent probesets
AL080102	Homo sapiens mRNA; cDNA DKFZp564N1916 (from clone	0.242	0.145	4 consistent probesets
N95443	zb81c12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-31	0.242	0.153	4 consistent probesets
M80783	Human B12 protein mRNA, complete cds /cds=(153,1103) /	0.242	0.162	4 consistent probesets
U25165	Human fragile X mental retardation protein 1 homolog F	0.242	0.165	4 consistent probesets
M74089	Human TB1 gene mRNA, 3 end /cds=(0,1305) /gb=M7408	0.242	0.166	4 consistent probesets
AB002299	Human mRNA for KIAA0301 gene, partial cds /cds=(0,6144	0.242	0.167	4 consistent probesets
AL049933	Homo sapiens mRNA; cDNA DKFZp564K1216 (from clone	0.242	0.170	4 consistent probesets
AI610467	tp42a10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-219	0.242	0.191	4 inconsistent probesets
AL050260	Homo sapiens mRNA; cDNA DKFZp547E1010 (from clone	0.242	0.200	4 consistent probesets
U12897	Homo sapiens IPW mRNA sequence /cds=UNKNOWN /gb=	0.242	0.203	4 consistent probesets
AI310002	qo77c11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-19	0.238	0.060	4 consistent probesets
AB028986	Homo sapiens mRNA for KIAA1063 protein, partial cds /c	0.238	0.062	4 consistent probesets
D21851	Human mRNA for KIAA0028 gene, partial cds /cds=(185,28	0.238	0.092	4 consistent probesets
D42063	Human mRNA for RanBP2 (Ran-binding protein 2), comple	0.238	0.122	4 consistent probesets
X17025	Human homolog of yeast IPP isomerase /cds=(50,736) /gb	0.238	0.123	4 consistent probesets
AL050073	Homo sapiens mRNA; cDNA DKFZp566E2346 (from clone	0.238	0.133	4 consistent probesets
AI146846	qb92h04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-17	0.238	0.151	4 consistent probesets
H97470	yw11b04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.238	0.174	4 inconsistent probesets
AB001636	Homo sapiens mRNA for ATP-dependent RNA helicase #46	0.238	0.207	4 consistent probesets
AF071185	Homo sapiens formin binding protein 21 mRNA, complete	0.238	0.223	4 consistent probesets
AB023163	Homo sapiens mRNA for KIAA0946 protein, partial cds /c	0.238	0.224	4 consistent probesets
U83115	Human non-lens beta gamma-crystallin like protein (AIM	0.238	0.247	4 consistent probesets
M80359	Human protein p78 mRNA, complete cds /cds=(171,2312) /	0.238	0.255	4 consistent probesets
U93869	Human RNA polymerase III subunit (RPC39) mRNA, compl	0.238	0.280	4 consistent probesets
M90656	Human gamma-glutamylcysteine synthetase (GCS) mRNA,	0.237	0.122	4 consistent probesets
X54942	H.sapiens ckshs2 mRNA for Cks1 protein homologue /cds=	0.233	0.041	4 inconsistent probesets
AF005043	AF005043 Homo sapiens poly(ADP-ribose) glycoh	0.233	0.098	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U51334	Human putative RNA binding protein (RBP56) mRNA, comp	0.233	0.102	4 inconsistent probesets
D30655	HUMELF4AII Homo sapiens mRNA for eukaryotic i	0.233	0.102	4 consistent probesets
J03853	Human kidney alpha-2-adrenergic receptor mRNA, complet	0.233	0.120	4 consistent probesets
D80001	Human mRNA for KIAA0179 gene, partial cds /cds=(0,2288	0.233	0.122	4 consistent probesets
AB011144	Homo sapiens mRNA for KIAA0572 protein, partial cds /c	0.233	0.125	4 consistent probesets
AB011155	Homo sapiens mRNA for KIAA0583 protein, partial cds /c	0.233	0.128	4 consistent probesets
U30894	Human N-sulphoglucosamine sulphohydrolase mRNA, com	0.233	0.135	4 consistent probesets
X87843	HSCYCHASS H.sapiens mRNA for cyclin H assembl	0.233	0.144	6 consistent probesets
M91029	Human AMP deaminase (AMPD2) mRNA /cds=(0,2282) /gb	0.233	0.147	3 consistent probesets
AF020043	Homo sapiens chromosome-associated polypeptide (HCAP	0.233	0.150	4 consistent probesets
AB023209	Homo sapiens mRNA for KIAA0992 protein, partial cds /c	0.233	0.158	4 inconsistent probesets
U58198	Human interleukin enhancer binding factor 3 mRNA, comp	0.233	0.159	5 consistent probesets
U79297	Human clone 23589 mRNA sequence /cds=UNKNOWN /gb	0.233	0.169	4 consistent probesets
AL050018	Homo sapiens mRNA; cDNA DKFZp564B116 (from clone D	0.233	0.169	4 consistent probesets
D10656	HUMCRK Human mRNA for CRK-II, complete cds"	0.233	0.173	6 consistent probesets
AF045184	Homo sapiens nuclear receptor coactivator NCoA-62 mRNA	0.233	0.194	3 consistent probesets
AF000560	Homo sapiens TTF-I interacting peptide 20 mRNA, partia	0.233	0.202	3 consistent probesets
M83738	HUMPTPSA Human protein-tyrosine phosphatase (	0.233	0.208	3 consistent probesets
AB020676	Homo sapiens mRNA for KIAA0869 protein, partial cds /c	0.233	0.215	3 consistent probesets
AJ006068	Homo sapiens mRNA for dTDP-D-glucose 4,6-dehydratase	0.233	0.224	4 consistent probesets
AI743090	wg87a11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.233	0.227	3 consistent probesets
U59423	HSU59423 Human Smad1 mRNA, complete cds"	0.233	0.229	3 consistent probesets
M55536	Human glucose transporter pseudogene /cds=UNKNOWN /	0.233	0.230	3 inconsistent probesets
X64838	H.sapiens mRNA for restin /cds=(132,4415) /gb=X64838 /	0.233	0.231	3 consistent probesets
AF006516	Homo sapiens eps8 binding protein e3B1 mRNA, complete	0.233	0.241	3 consistent probesets
AI935442	wo84e06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.233	0.242	4 consistent probesets
L31573	Human sulfite oxidase mRNA, complete cds /cds=(903,236	0.233	0.247	3 consistent probesets
AI890903	wm91f10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.233	0.254	3 consistent probesets
U48736	HSU48736 Human serine/threonine-protein kinas	0.233	0.266	3 inconsistent probesets
AB018338	Homo sapiens mRNA for KIAA0795 protein, partial cds /c	0.233	0.275	4 consistent probesets
D38583	Human mRNA for calgizzarin, complete cds /cds=(120,437	0.233	0.288	3 consistent probesets
AJ006778	Homo sapiens mRNA for DRIM protein /cds=(144,8501) /gb	0.233	0.295	3 consistent probesets
AA887480	oj54a12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-150	0.233	0.296	4 consistent probesets
AB020623	Homo sapiens DAM1 mRNA, complete cds /cds=(48,725) /g	0.233	0.301	3 consistent probesets
X91809	H.sapiens mRNA for GAIP protein /cds=(288,941) /gb=X91	0.233	0.303	3 consistent probesets
AF070590	Homo sapiens clones 24622 and 24623 mRNA sequence /c	0.233	0.415	3 consistent probesets
AL049266	Homo sapiens mRNA; cDNA DKFZp564F093 (from clone D	0.233	0.433	3 consistent probesets
U18543	Human zinc-finger protein mRNA, complete cds /cds=(110	0.233	0.449	3 consistent probesets
U50928	Human autosomal dominant polycystic kidney disease typ	0.233	0.526	3 consistent probesets
M74587	HUMIGFBP1A Human insulin-like growth factor b	0.229	0.064	4 consistent probesets
U49436	HSU49436 Human translation initiation factor	0.229	0.089	4 consistent probesets
M31899	HUMERCC3A Human DNA repair helicase (ERCC3) m	0.229	0.093	4 consistent probesets
L40410	Homo sapiens thyroid receptor interactor (TRIP3) mRNA,	0.229	0.098	4 consistent probesets
AF091078	Homo sapiens clone 559 unknown mRNA, complete sequer	0.229	0.105	8 consistent probesets
X05360	HSCDC2 Human CDC2 gene involved in cell cycle	0.229	0.118	4 consistent probesets
AF022789	Homo sapiens ubiquitin hydrolyzing enzyme I (UBH1) mRN	0.229	0.139	4 consistent probesets
AB011158	Homo sapiens mRNA for KIAA0586 protein, complete cds /	0.229	0.146	4 consistent probesets
L07648	HUMMXI1A Human MXI1 mRNA, complete cds"	0.229	0.147	8 inconsistent probesets
AB029005	Homo sapiens mRNA for KIAA1082 protein, partial cds /c	0.229	0.184	4 consistent probesets
AJ006267	Homo sapiens mRNA for ClpX-like protein /cds=(27,1826)	0.229	0.196	4 consistent probesets
AL080169	Homo sapiens mRNA; cDNA DKFZp434C171 (from clone D	0.229	0.230	4 consistent probesets
N32257	yy28d09.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-27	0.229	0.242	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
M95585	HUMHLF Human hepatic leukemia factor (HLF) mR	0.227	0.070	8 inconsistent probesets
AL008583	dJ327J16.3 (novel CHROMObox family protein) /cds=(29,1	0.226	0.141	7 consistent probesets
X91247	H.sapiens mRNA for thioredoxin reductase /cds=(439,193	0.225	0.058	4 inconsistent probesets
AF084928	Homo sapiens erythroblast macrophage protein EMP mRNA	0.225	0.070	4 consistent probesets
U37426	Human kinesin-like spindle protein HKSP (HKSP) mRNA, c	0.225	0.077	4 consistent probesets
D11466	Homo sapiens mRNA for PIG-A protein, complete cds /cds	0.225	0.101	4 consistent probesets
X98263	H.sapiens mRNA for M-phase phosphoprotein, mpp6 /cds=(	0.225	0.102	4 consistent probesets
D50645	Homo sapiens mRNA for SDF2, complete cds /cds=(39,674	0.225	0.104	4 consistent probesets
Y00451	Human mRNA for 5-aminolevulinate synthase /cds=(83,201	0.225	0.109	4 consistent probesets
AF044968	untitled /cds=(0,1351) /gb=AF044968 /gi=3941380 /	0.225	0.118	4 consistent probesets
AF042729	Homo sapiens lithium-sensitive myo-inositol monophosph	0.225	0.120	4 consistent probesets
AB014519	Homo sapiens mRNA for KIAA0619 protein, complete cds /	0.225	0.121	4 consistent probesets
M99701	Homo sapiens (pp21) mRNA, complete cds /cds=(164,637)	0.225	0.123	4 consistent probesets
X78338	Synthetic adenovirus transformed human retina cell lin	0.225	0.145	4 consistent probesets
AB029028	Homo sapiens mRNA for KIAA1105 protein, partial cds /c	0.225	0.168	4 consistent probesets
L19182	HUMMAC25X Human MAC25 mRNA, complete cds"	0.225	0.182	2 consistent probesets
AA059408	zl96e07.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-512	0.225	0.201	4 consistent probesets
AB029037	Homo sapiens mRNA for KIAA1114 protein, complete cds /	0.225	0.217	2 consistent probesets
L36870	HUMMKK4A Homo sapiens MAP kinase kinase 4 (MK	0.225	0.225	4 consistent probesets
AB011121	Homo sapiens mRNA for KIAA0549 protein, partial cds /c	0.225	0.229	4 consistent probesets
W25911	14h11 Homo sapiens cDNA /gb=W25911 /gi=1306034 /	0.225	0.237	4 consistent probesets
D87119	D87119 Homo sapiens mRNA for GS3955, complete	0.225	0.238	2 consistent probesets
D80007	Human mRNA for KIAA0185 gene, partial cds /cds=(0,5655	0.225	0.248	2 consistent probesets
AB029036	Homo sapiens mRNA for KIAA1113 protein, partial cds /c	0.225	0.273	2 consistent probesets
AJ011896	Homo sapiens mRNA for HIV-1, Nef-associated factor 1 b	0.225	0.273	2 consistent probesets
U80743	Homo sapiens CAGH32 mRNA, partial cds /cds=(0,1671) /g	0.225	0.304	2 consistent probesets
AB018272	Homo sapiens mRNA for KIAA0729 protein, partial cds /c	0.225	0.311	4 consistent probesets
AF070644	Homo sapiens clone 24742 mRNA sequence /cds=UNKNO	0.225	0.336	2 consistent probesets
M27543	HUMGNAI1 Human guanine nucleotide-binding pro	0.225	0.398	2 consistent probesets
M58603	HUMNFKB Human nuclear factor kappa-B DNA bind	0.224	0.101	11 consistent probesets
L10413	HUMFTA Human farnesyltransferase alpha-subuni	0.221	0.068	4 inconsistent probesets
X80199	H.sapiens MLN51 mRNA /cds=(233,1837) /gb=X80199 /gi=	0.221	0.072	4 consistent probesets
Y12781	Homo sapiens mRNA for transducin (beta) like 1 protein	0.221	0.081	4 consistent probesets
AJ132712	Homo sapiens mRNA for tip associating protein (TAP) /c	0.221	0.088	4 inconsistent probesets
S80343	S80343 ArgRS=arginyl-tRNA synthetase [human,	0.221	0.091	4 consistent probesets
AF055479	Homo sapiens lung cancer candidate FUS1 (FUS1) mRNA,	0.221	0.110	4 inconsistent probesets
D13636	Human mRNA for KIAA0011 gene, complete cds /cds=(39,2	0.221	0.111	4 consistent probesets
AL050105	Homo sapiens mRNA; cDNA DKFZp586H0519 (from clone	0.221	0.116	4 consistent probesets
AB014596	Homo sapiens mRNA for KIAA0696 protein, partial cds /c	0.221	0.116	4 consistent probesets
Y00978	Human mRNA for dihydrolipoamide acetyltransferase (PDC	0.221	0.127	4 consistent probesets
D89053	Homo sapiens mRNA for Acyl-CoA synthetase 3, complete	0.221	0.166	4 consistent probesets
W27466	31c9 Homo sapiens cDNA /gb=W27466 /gi=1307270 /	0.221	0.167	4 consistent probesets
X81788	Homo sapiens ICT1 (alias DS-1) mRNA /cds=(2,622) /gb=X	0.221	0.188	4 consistent probesets
AF007551	Homo sapiens Bet1p homolog (hbet1) mRNA, complete cds	0.221	0.189	4 consistent probesets
U22816	Human LAR-interacting protein 1b mRNA, complete cds /c	0.221	0.195	4 consistent probesets
X97795	HSRAD54 H.sapiens mRNA homologous to S. cerev	0.220	0.139	5 inconsistent probesets
AF054186	Homo sapiens p18 protein mRNA, complete cds /cds=(28,5	0.219	0.100	7 inconsistent probesets
U03911	HSU03911 Human mutator gene (hMSH2) mRNA, com	0.219	0.093	8 consistent probesets
AF058696	Homo sapiens cell cycle regulatory protein p95 (NBS1)	0.217	0.068	4 inconsistent probesets
X63563	H.sapiens mRNA for RNA polymerase II 140 kDa subunit /	0.217	0.074	4 inconsistent probesets
U37146	Human silencing mediator of retinoid and thyroid hormo	0.217	0.077	4 consistent probesets
AJ006291	Homo sapiens mRNA for leucine rich protein /cds=(63,92	0.217	0.085	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
W28865	53g9 Homo sapiens cDNA /gb=W28865 /gi=1308876 /	0.217	0.115	4 consistent probesets
Z74792	H.sapiens mRNA for CCAAT transcription binding factor	0.217	0.119	3 consistent probesets
U73704	Homo sapiens 48 kDa FKBP-associated protein FAP48 mR	0.217	0.132	8 consistent probesets
L43964	Homo sapiens (clone F-T03796) STM-2 mRNA, complete c	0.217	0.139	3 consistent probesets
AB018333	Homo sapiens mRNA for KIAA0790 protein, partial cds /c	0.217	0.141	4 consistent probesets
AA142964	zl43a07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-504	0.217	0.148	4 consistent probesets
X60287	HSMAXM H.sapiens max mRNA	0.217	0.153	3 inconsistent probesets
U79291	Human clone 23721 mRNA sequence /cds=UNKNOWN /gb	0.217	0.156	4 consistent probesets
AI582831	tn36c06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-216	0.217	0.158	3 consistent probesets
X66113	H.sapiens mRNA for PM/Scl 100kD nucleolar protein /cds	0.217	0.167	3 consistent probesets
AL047596	DKFZp586G0421_s1 Homo sapiens cDNA /clone=DKFZp5	0.217	0.175	4 consistent probesets
D79987	Human mRNA for KIAA0165 gene, complete cds /cds=(111	0.217	0.179	3 consistent probesets
AB007144	Homo sapiens mRNA for ZIP-kinase, complete cds /cds=(9	0.217	0.196	3 consistent probesets
U26914	HSU26914 Human ras-responsive element binding	0.217	0.202	4 consistent probesets
D86969	Human mRNA for KIAA0215 gene, complete cds /cds=(298	0.217	0.203	4 consistent probesets
D31886	Human mRNA for KIAA0066 gene, partial cds /cds=(0,2947	0.217	0.206	3 consistent probesets
AF045458	Homo sapiens serine/threonine kinase ULK1 (ULK1) mRNA	0.217	0.208	3 consistent probesets
X84002	HSTAFII20 H.sapiens TAFII20 mRNA for transcri	0.217	0.225	3 consistent probesets
AB020660	Homo sapiens mRNA for KIAA0853 protein, partial cds /c	0.217	0.244	3 consistent probesets
U60337	Homo sapiens beta-mannosidase mRNA, complete cds /cds	0.217	0.248	4 consistent probesets
AA126515	zn85c12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-564	0.217	0.280	4 consistent probesets
U62325	Human FE65-like protein (hFE65L) mRNA, partial cds /cd	0.217	0.282	3 consistent probesets
L00049	Human cellular c-Ki-ras2 proto-oncogene, 5 flank and /	0.217	0.282	4 consistent probesets
M73077	Human glucocorticoid receptor repression factor 1 (GRF	0.217	0.295	3 consistent probesets
AA044787	zk74c11.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-488	0.217	0.297	3 consistent probesets
U46024	Homo sapiens myotubularin (MTM1) mRNA, complete cds /	0.217	0.303	3 consistent probesets
AB023210	Homo sapiens mRNA for KIAA0993 protein, partial cds /c	0.217	0.388	3 consistent probesets
AB023162	Homo sapiens mRNA for KIAA0945 protein, complete cds /	0.217	0.406	3 consistent probesets
M25077	Human SS-A/Ro ribonucleoprotein autoantigen 60 kd subu	0.215	0.108	8 consistent probesets
L77886	HUMPTPC Human protein tyrosine phosphatase mR	0.213	0.055	4 consistent probesets
U07806	HSU07806 Human camptothecin resistant clone C	0.213	0.077	4 consistent probesets
AB006625	Homo sapiens mRNA for KIAA0287 gene, partial cds /cds=	0.213	0.082	4 consistent probesets
AI365215	qz41a06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-20	0.213	0.085	4 consistent probesets
M25753	HUMCYCB Human cyclin B mRNA, 3 end"	0.213	0.089	8 consistent probesets
AI056696	oz26h05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.213	0.096	8 inconsistent probesets
J02939	Human membrane glycoprotein 4F2 antigen heavy chain m	0.213	0.098	4 consistent probesets
D83777	Human mRNA for KIAA0193 gene, complete cds /cds=(352	0.213	0.102	4 inconsistent probesets
U02619	Human TFIIIC Box B-binding subunit mRNA, complete cds	0.213	0.102	4 inconsistent probesets
AL080162	Homo sapiens mRNA; cDNA DKFZp434N024 (from clone D	0.213	0.108	4 consistent probesets
U82328	Homo sapiens pyruvate dehydrogenase complex protein X	0.213	0.120	4 consistent probesets
W26019	18b9 Homo sapiens cDNA /gb=W26019 /gi=1306304 /	0.213	0.124	4 inconsistent probesets
L20859	HUMGLVR1X Human leukemia virus receptor 1 (GL	0.213	0.128	4 consistent probesets
AI307607	tb15h10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-205	0.213	0.130	4 consistent probesets
Y08698	H.sapiens mRNA for RanBP3 (59 kDa) /cds=(20,1708) /gb=	0.213	0.131	4 consistent probesets
AF025654	Homo sapiens mRNA capping enzyme (HCE) mRNA, comp	0.213	0.131	4 consistent probesets
X52015	H.sapiens mRNA for interleukin-1 receptor antagonist /	0.213	0.132	4 consistent probesets
U63743	Homo sapiens mitotic centromere-associated kinesin mRN	0.213	0.136	4 consistent probesets
D00591	HUMRCC1 Homo sapiens RCC1 gene, exons 1, 2, 3	0.213	0.136	4 inconsistent probesets
D80010	Human mRNA for KIAA0188 gene, partial cds /cds=(0,2700	0.213	0.141	4 inconsistent probesets
AF084513	Homo sapiens DNA repair exonuclease (REC1) mRNA, alte	0.213	0.147	4 consistent probesets
U09848	Human zinc finger protein (ZNF139) mRNA, partial cds /	0.213	0.149	4 consistent probesets
AL050390	Homo sapiens mRNA; cDNA DKFZp564O043 (from clone D	0.213	0.159	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U63809	Homo sapiens prostate apoptosis response protein par-4	0.213	0.166	4 consistent probesets
AB023197	Homo sapiens mRNA for KIAA0980 protein, partial cds /c	0.213	0.185	4 consistent probesets
AF026402	Homo sapiens U5 snRNP 100 kD protein mRNA, complete	0.213	0.225	4 consistent probesets
D63876	Human mRNA for KIAA0154 gene, partial cds /cds=(0,2080	0.213	0.318	4 consistent probesets
U96113	HSU96113 Homo sapiens Nedd-4-like ubiquitin-p	0.212	0.064	8 consistent probesets
AF037204	Homo sapiens RING zinc finger protein (RZF) mRNA, comp	0.212	0.209	4 consistent probesets
AF095791	Homo sapiens TACC2 protein (TACC2) mRNA, partial cds /	0.212	0.269	4 consistent probesets
U80017	Homo sapiens basic transcription factor 2 p44 (btf2p44	0.211	0.194	6 consistent probesets
L13689	HUMBMI1X Human prot-oncogene (BMI-1) mRNA, co	0.210	0.078	8 consistent probesets
X97544	H.sapiens mRNA for TIM17 preprotein translocase /cds=(	0.210	0.088	8 consistent probesets
AL031778	dJ34B21.4.1 (nuclear transcription factor Y, alpha (CC	0.210	0.092	8 inconsistent probesets
AF026166	Homo sapiens chaperonin-containing TCP-1 beta subunit	0.208	0.058	4 consistent probesets
AI526089	DU3.2-7.H07.r Homo sapiens cDNA, 5 end /clone_end=5	0.208	0.085	4 inconsistent probesets
AI659108	tu08c09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-225	0.208	0.085	4 inconsistent probesets
H16917	ym39e02.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-50	0.208	0.088	4 consistent probesets
U09577	Homo sapiens lysosomal hyaluronidase (LUCA2/HYAL2) m	0.208	0.093	4 consistent probesets
M31724	HUMPTPBX Human phosphotyrosyl-protein phospha	0.208	0.093	8 consistent probesets
U05875	Human clone pSK1 interferon gamma receptor accessory f	0.208	0.095	4 consistent probesets
L42373	HUMPP2A Homo sapiens phosphatase 2A B56-alpha	0.208	0.109	4 consistent probesets
AA780049	zj24f06.s1 Homo sapiens cDNA, 3 end /clone=451235 /cl	0.208	0.121	4 consistent probesets
AA926957	om68h06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-15	0.208	0.121	4 consistent probesets
M33509	Human HLA-B-associated transcript 2 (BAT2) mRNA, comp	0.208	0.127	4 consistent probesets
D79993	Human mRNA for KIAA0171 gene, complete cds /cds=(101	0.208	0.134	4 consistent probesets
AL050139	Homo sapiens mRNA; cDNA DKFZp586M141 (from clone D	0.208	0.135	4 consistent probesets
M15353	Homo sapiens cap-binding protein mRNA, complete cds /c	0.208	0.139	4 consistent probesets
U15642	HSU15642 Human transcription factor E2F-5 mRN	0.208	0.147	4 consistent probesets
U12535	HSU12535 Human epidermal growth factor recept	0.208	0.153	4 consistent probesets
U60519	HSU60519 Human apoptotic cysteine protease Mc	0.208	0.154	8 consistent probesets
U24576	U24576 Homo sapiens breast tumor autoantigen	0.208	0.159	4 consistent probesets
U90543	Human butyrophilin (BTF1) mRNA, complete cds /cds=(210	0.208	0.166	4 consistent probesets
X66435	H.sapiens mRNA for HMG-CoA-synthase /cds=(122,1684) /	0.208	0.172	4 inconsistent probesets
M29550	Human calcineurin A1 mRNA, complete cds /cds=(107,165	0.208	0.210	4 consistent probesets
AB014550	Homo sapiens mRNA for KIAA0650 protein, partial cds /c	0.208	0.214	4 consistent probesets
AL080190	Homo sapiens mRNA; cDNA DKFZp434A202 (from clone D	0.208	0.217	4 consistent probesets
D87459	Human mRNA for KIAA0269 gene, complete cds /cds=(242	0.208	0.264	4 consistent probesets
AF008915	Homo sapiens EVI5 homolog mRNA, complete cds /cds=(1	0.208	0.328	4 consistent probesets
AL080122	Homo sapiens mRNA; cDNA DKFZp564O123 (from clone D	0.208	0.448	4 consistent probesets
U93867	HSU93867 Human RNA polymerase III subunit (RP	0.207	0.173	5 consistent probesets
D84454	Human mRNA for UDP-galactose translocator, complete co	0.206	0.237	3 consistent probesets
X58288	HSHRPTPU H.sapiens hR-PTPu gene for protein t	0.205	0.144	10 consistent probesets
D12686	HUMEIF4G Human mRNA for eukaryotic initiation	0.204	0.070	4 consistent probesets
U84720	Homo sapiens mRNA export protein (RAE1) mRNA, comple	0.204	0.081	8 consistent probesets
AF052130	Homo sapiens clone 23704 mRNA sequence /cds=UNKNO	0.204	0.083	4 consistent probesets
Y18314	Homo sapiens mRNA for paraplegin-like protein /cds=(11	0.204	0.087	4 consistent probesets
W26023	18c3 Homo sapiens cDNA /gb=W26023 /gi=1306308 /	0.204	0.114	4 consistent probesets
D78130	Homo sapiens mRNA for squalene epoxidase, complete cds	0.204	0.121	4 consistent probesets
M64347	Human novel growth factor receptor mRNA, 3 cds /cds=(0	0.204	0.123	4 consistent probesets
AL080184	Homo sapiens mRNA; cDNA DKFZp434O071 (from clone D	0.204	0.137	4 consistent probesets
U68018	HSU68018 Human mad protein homolog (hMAD-2) m	0.204	0.152	4 consistent probesets
U11791	HSU11791 Human cyclin H mRNA, complete cds"	0.204	0.159	4 consistent probesets
AF006041	AF006041 Homo sapiens Fas-binding protein (DA	0.204	0.174	4 consistent probesets
D50840	Homo sapiens mRNA for ceramide glucosyltransferase, co	0.204	0.179	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D31889	Human mRNA for KIAA0072 gene, partial cds /cds=(0,1513	0.204	0.179	4 consistent probesets
U67058	Human proteinase activated receptor-2 mRNA, 3UTR /cds=	0.204	0.206	4 consistent probesets
AF072928	Homo sapiens myotubularin related protein 6 mRNA, part	0.204	0.220	4 consistent probesets
U37352	HSU37352 Human protein phosphatase 2A B alpha	0.202	0.107	8 consistent probesets
AA868898	ak55b08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-14	0.202	0.159	8 consistent probesets
X55005	HSCERBAR Homo sapiens mRNA for thyroid hormon	0.200	0.098	1 consistent probesets
AB014514	Homo sapiens mRNA for KIAA0614 protein, partial cds /c	0.200	0.101	1 consistent probesets
AB002378	Human mRNA for KIAA0380 gene, complete cds /cds=(745	0.200	0.101	1 consistent probesets
M80899	Human novel protein AHNAK mRNA, partial sequence /cds	0.200	0.105	4 consistent probesets
D50928	Human mRNA for KIAA0138 gene, complete cds /cds=(36,2	0.200	0.136	3 consistent probesets
D26362	Human mRNA for KIAA0043 gene, complete cds /cds=(139	0.200	0.137	4 consistent probesets
R16035	ya51h07.r2 Homo sapiens cDNA, 5 end /clone=IMAGE-664	0.200	0.138	4 consistent probesets
D87461	Human mRNA for KIAA0271 gene, complete cds /cds=(176	0.200	0.140	4 consistent probesets
M29893	Human low molecular mass GTP-binding protein (ral) mRN	0.200	0.142	4 consistent probesets
AB028980	Homo sapiens mRNA for KIAA1057 protein, partial cds /c	0.200	0.145	3 consistent probesets
AB011126	Homo sapiens mRNA for KIAA0554 protein, partial cds /c	0.200	0.156	3 consistent probesets
L36818	Human (clone 51C-3) 51C protein mRNA, complete cds /cd	0.200	0.161	2 consistent probesets
L42025	Homo sapiens cellular co-factor (RAB) gene, complete c	0.200	0.195	4 consistent probesets
L28010	Homo sapiens HnRNP F protein mRNA, complete cds /cds=	0.200	0.198	3 consistent probesets
M12625	Human lecithin-cholesterol acyltransferase mRNA, compl	0.200	0.202	1 consistent probesets
AI683743	tw53d12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-22	0.200	0.202	1 consistent probesets
AB007859	Homo sapiens KIAA0399 mRNA, partial cds /cds=(0,2352)	0.200	0.203	2 consistent probesets
X74594	HSRB2P130 H.sapiens mRNA for Rb2/p130 protein	0.200	0.222	3 inconsistent probesets
AL049309	Homo sapiens mRNA; cDNA DKFZp564B176 (from clone D	0.200	0.228	4 consistent probesets
AL121073	DKFZp762B235_r1 Homo sapiens cDNA, 5 end /clone=DK	0.200	0.231	3 consistent probesets
M63978	Human vascular endothelial growth factor gene /cds=(10	0.200	0.252	1 consistent probesets
AI627354	ty75d06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-228	0.200	0.252	1 consistent probesets
X52151	Homo sapiens arylsulphatase A mRNA, complete cds /cds=	0.200	0.268	2 consistent probesets
AF042169	Homo sapiens putative ATP-dependent mitochondrial RNA	0.200	0.276	3 consistent probesets
AB023203	Homo sapiens mRNA for KIAA0986 protein, partial cds /c	0.200	0.302	3 consistent probesets
AL049787	Novel human gene mapping to chomosome 13 /cds=(557,7	0.200	0.302	1 consistent probesets
AF015257	Homo sapiens flow-induced endothelial G protein-couple	0.200	0.302	1 consistent probesets
Y07846	H.sapiens mRNA for GAR22 protein /cds=(132,1145) /gb=Y	0.200	0.302	1 consistent probesets
X83441	HALIG4 H.sapiens mRNA for DNA ligase IV	0.200	0.302	1 consistent probesets
AL022316	Human DNA sequence from clone 126B4 on chromosome 2	0.200	0.323	2 consistent probesets
AL049387	Homo sapiens mRNA; cDNA DKFZp586N1918 (from clone	0.200	0.338	3 consistent probesets
Z70519	H.sapiens FAS/Apo 1 mRNA for FAS soluble protein (clon	0.200	0.343	2 consistent probesets
AB009598	Homo sapiens mRNA for glucuronyltransferase I, complet	0.200	0.353	1 consistent probesets
AF105424	Homo sapiens brush border myosin I (BBMI) mRNA, compl	0.200	0.353	1 consistent probesets
AA058762	zk65d06.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-487	0.200	0.400	2 consistent probesets
D31840	Human DRPLA mRNA for ORF, complete cds /cds=(238,37	0.200	0.403	1 consistent probesets
AJ001685	Homo sapiens NKG2E gene /cds=(45,767) /gb=AJ001685 /(	0.200	0.454	1 consistent probesets
AB014592	Homo sapiens mRNA for KIAA0692 protein, partial cds /c	0.200	0.463	2 consistent probesets
L35594	Human autotaxin mRNA, complete cds /cds=(49,2796) /gb=	0.200	0.504	1 consistent probesets
AB022663	Homo sapiens HFB30 mRNA, complete cds /cds=(236,1660	0.200	0.517	2 consistent probesets
AL049346	Homo sapiens mRNA; cDNA DKFZp566B213 (from clone D	0.200	0.544	3 consistent probesets
Y14155	Homo sapiens mRNA for Hmob33 protein, 3 untranslated r	0.200	0.554	1 consistent probesets
AF009425	Homo sapiens clone 22 mRNA, alternative splicing varia	0.200	0.554	1 consistent probesets
U15552	Human acidic 82 kDa protein mRNA, complete cds /cds=(2	0.200	0.554	1 consistent probesets
AA203527	zx56f09.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-446	0.200	0.555	2 consistent probesets
AB026190	Homo sapiens mRNA for Kelch motif containing protein,	0.200	0.571	2 consistent probesets
U22963	Human class I histocompatibility antigen-like protein	0.200	0.655	1 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U35376	Human repressor transcriptional factor (ZNF85) mRNA, c	0.200	0.655	1 consistent probesets
AB002351	Human mRNA for KIAA0353 gene, partial cds /cds=(0,4125	0.200	0.655	1 consistent probesets
AL021331	dJ366N23.2 (putative C. elegans UNC-93 (protein 1, C46	0.200	0.683	1 consistent probesets
W28734	51a1 Homo sapiens cDNA /gb=W28734 /gi=1308682 /	0.200	0.706	1 consistent probesets
W26640	34b6 Homo sapiens cDNA /gb=W26640 /gi=1307483 /	0.200	0.731	1 consistent probesets
L36051	HUMTHROMA Human thrombopoietin gene, complete	0.200	0.756	1 consistent probesets
AL078636	Homo sapiens mRNA full length insert cDNA clone EUROIN	0.200	0.806	1 consistent probesets
U18914	Human 19.8 kDa protein mRNA, complete cds /cds=(91,645	0.200	0.857	1 consistent probesets
X67337	H.sapiens HPBRII-4 mRNA /cds=(34,1689) /gb=X67337 /gi=	0.200	0.857	1 consistent probesets
X83300	H.sapiens SMA4 mRNA /cds=(66,488) /gb=X83300 /gi=603	0.200	0.857	1 consistent probesets
AB002319	Human mRNA for KIAA0321 gene, partial cds /cds=(0,4628	0.200	0.907	1 consistent probesets
X00737	HSPNP Human mRNA for purine nucleoside phosph	0.200	0.072	4 consistent probesets
U82130	HSU82130 Human tumor susceptiblity protein (T	0.200	0.085	4 inconsistent probesets
X63564	H.sapiens mRNA for RNA polymerase II largest subunit /	0.200	0.086	4 consistent probesets
U75679	Human histone stem-loop binding protein (SLBP) mRNA, c	0.200	0.118	4 consistent probesets
L76703	Homo sapiens protein phosphatase 2A B56-epsilon (PP2A)	0.200	0.170	4 consistent probesets
U18937	Human histidyl-tRNA synthetase homolog (HO3) mRNA, co	0.200	0.211	4 consistent probesets
AB023160	Homo sapiens mRNA for KIAA0943 protein, partial cds /c	0.200	0.403	2 consistent probesets
AL049851	Human DNA sequence from clone 889J22B on chromosom	0.198	0.084	8 inconsistent probesets
AL031775	dJ30M3.2 (novel protein) /cds=(0,260) /gb=AL031775 /gi	0.198	0.092	8 inconsistent probesets
AF104913	Homo sapiens eukaryotic protein synthesis initiation f	0.196	0.078	4 consistent probesets
AF048977	Homo sapiens Ser/Arg-related nuclear matrix protein (S	0.196	0.081	4 consistent probesets
AF000984	Homo sapiens dead box, Y isoform (DBY) mRNA, alternati	0.196	0.081	4 consistent probesets
AL049948	Homo sapiens mRNA; cDNA DKFZp564K0222 (from clone	0.196	0.082	4 consistent probesets
X87241	H.sapiens mRNA for hFat protein /cds=(186,13958) /gb=X	0.196	0.093	4 consistent probesets
X59434	Human rohu mRNA for rhodanese /cds=(34,924) /gb=X594	0.196	0.097	4 consistent probesets
U05237	Human fetal Alz-50-reactive clone 1 (FAC1) mRNA, compl	0.196	0.115	4 inconsistent probesets
AF032862	Homo sapiens intracellular hyaluronic acid binding pro	0.196	0.116	4 consistent probesets
U32315	Human syntaxin 3 mRNA, complete cds /cds=(38,907) /gb=	0.196	0.132	4 consistent probesets
AF032456	Homo sapiens ubiquitin conjugating enzyme G2 (UBE2G2)	0.196	0.141	4 consistent probesets
AL050205	Homo sapiens mRNA; cDNA DKFZp586F1323 (from clone	0.196	0.147	4 consistent probesets
D78275	D78275 Homo sapiens mRNA for proteasome subun	0.196	0.147	4 consistent probesets
AB007888	Homo sapiens KIAA0428 mRNA, complete cds /cds=(1414,	0.196	0.153	4 consistent probesets
X82834	H.sapiens mRNA for golgin /cds=(207,6764) /gb=X82834 /	0.196	0.153	4 consistent probesets
D29013	HUMLNCAP Human mRNA for DNA polymerase beta,	0.196	0.154	4 consistent probesets
U14391	Human myosin-IC mRNA, complete cds /cds=(375,3704) /g	0.196	0.175	4 consistent probesets
X80507	H.sapiens YAP65 mRNA /cds=(0,1364) /gb=X80507 /gi=51	0.196	0.177	4 consistent probesets
AJ007583	Homo sapiens mRNA for acetylglucosaminyltransferase-li	0.196	0.177	4 consistent probesets
M38690	Human CD9 antigen mRNA, complete cds /cds=(51,737) /g	0.196	0.208	4 consistent probesets
AJ007042	Homo sapiens mRNA for TRX5 protein /cds=(762,4859) /gb	0.196	0.241	4 consistent probesets
U79745	Homo sapiens monocarboxylate transporter homologue MC	0.196	0.257	4 consistent probesets
AI688640	wd40b07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.196	0.258	4 consistent probesets
U59305	Human ser-thr protein kinase PK428 mRNA, complete cds	0.194	0.375	3 consistent probesets
AB015331	Homo sapiens HRIHFB2017 mRNA, partial cds /cds=(0,650	0.194	0.378	3 consistent probesets
AW051579	wy87g03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.193	0.144	7 consistent probesets
M20681	Human glucose transporter-like protein-III (GLUT3), co	0.192	0.053	4 consistent probesets
M77698	HUMKRP Homo sapiens GLI-Krupple related prote	0.192	0.073	4 consistent probesets
D79986	Human mRNA for KIAA0164 gene, complete cds /cds=(253	0.192	0.080	4 consistent probesets
W27233	24b7 Homo sapiens cDNA /gb=W27233 /gi=1306749 /	0.192	0.081	4 consistent probesets
AF098799	Homo sapiens RanBP7/importin 7 mRNA, complete cds /cd	0.192	0.081	4 consistent probesets
J03805	HUMALPHLB Human phosphatase 2A mRNA, partial	0.192	0.085	4 inconsistent probesets
AL050196	Homo sapiens mRNA; cDNA DKFZp586D2223 (from clone	0.192	0.094	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
M23114	Homo sapiens calcium-ATPase (HK1) mRNA, complete cds	0.192	0.108	4 consistent probesets
AF056490	Homo sapiens cAMP-specific phosphodiesterase 8A (PDE8	0.192	0.110	4 consistent probesets
U01833	Human nucleotide-binding protein mRNA, complete cds /c	0.192	0.112	4 consistent probesets
W87466	zh67c05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-41	0.192	0.112	4 consistent probesets
U77718	Human desmosome associated protein pinin mRNA, compl	0.192	0.113	4 consistent probesets
W25933	15b2 Homo sapiens cDNA /gb=W25933 /gi=1306056 /	0.192	0.114	4 consistent probesets
AF083255	Homo sapiens RNA helicase-related protein mRNA, comple	0.192	0.121	4 consistent probesets
AI627877	ty20b09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-227	0.192	0.126	4 consistent probesets
U40038	Human GTP-binding protein alpha q subunit (GNAQ) mRNA	0.192	0.150	4 consistent probesets
AJ001625	Homo sapiens mRNA for Pex3 protein /cds=(63,1184) /gb=	0.192	0.163	4 inconsistent probesets
AB011133	Homo sapiens mRNA for KIAA0561 protein, partial cds /c	0.192	0.174	4 consistent probesets
Y10571	H.sapiens mRNA for dinG gene /cds=(12,1022) /gb=Y1057	0.192	0.232	4 consistent probesets
M14752	HUMABLA Human c-abl gene, complete cds"	0.192	0.315	4 consistent probesets
U68111	HSPPP1R2E6 Human protein phosphatase inhibito	0.190	0.100	8 inconsistent probesets
X15606	Human mRNA for ICAM-2, cell adhesion ligand for LFA-1	0.190	0.130	8 inconsistent probesets
AJ011916	Homo sapiens mRNA for hypothetical protein /cds=(30,76	0.188	0.058	4 consistent probesets
X80878	H.sapiens R kappa B mRNA /cds=(130,4104) /gb=X80878 /	0.188	0.077	4 consistent probesets
J03161	HUMSRF Human serum response factor (SRF) mRNA	0.188	0.098	8 consistent probesets
H15872	ym22b12.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-48	0.188	0.098	4 consistent probesets
U07695	HSU07695 Human tyrosine kinase (HTK) mRNA, co	0.188	0.102	4 inconsistent probesets
AA127624	zk89b09.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-489	0.188	0.104	4 inconsistent probesets
U48705	HSU48705 Human receptor tyrosine kinase DDR g	0.188	0.105	4 consistent probesets
N36295	yx99b12.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-269	0.188	0.123	4 consistent probesets
D13635	Human mRNA for KIAA0010 gene, complete cds /cds=(303	0.188	0.133	4 consistent probesets
AB000409	Homo sapiens mRNA for MNK1, complete cds /cds=(187,14	0.188	0.139	4 consistent probesets
AB029016	Homo sapiens mRNA for KIAA1093 protein, partial cds /c	0.188	0.139	4 consistent probesets
D83767	Human clone N9 Rep-8 mRNA, complete cds /cds=(15,827	0.188	0.142	4 consistent probesets
AF032437	Homo sapiens mitogen activated protein kinase activate	0.188	0.143	4 consistent probesets
L37936	Human nuclear-encoded mitochondrial elongation factor	0.188	0.146	4 consistent probesets
AF041080	Homo sapiens D15F37 pseudogene, S3 allele, mRNA sequ	0.188	0.148	4 consistent probesets
AL050289	Homo sapiens mRNA; cDNA DKFZp586G0522 (from clone	0.188	0.148	4 consistent probesets
AB014570	Homo sapiens mRNA for KIAA0670 protein, partial cds /c	0.188	0.155	4 consistent probesets
U46691	Human putative chromatin structure regulator (SUPT6H)	0.188	0.161	4 consistent probesets
AB018298	Homo sapiens mRNA for KIAA0755 protein, complete cds /	0.188	0.183	4 consistent probesets
AB007940	Homo sapiens mRNA for KIAA0471 protein, complete cds /	0.188	0.205	4 consistent probesets
AW016815	UI-H-BI0-aam-c-09-0-UI.s1 Homo sapiens cDNA, 3 end /c	0.188	0.395	4 consistent probesets
S78234	nuc2 homolog [human, fibroblasts, mRNA, 3320 nt] /cds=	0.187	0.225	4 consistent probesets
L32977	Homo sapiens (clone f17252) ubiquinol cytochrome c red	0.183	0.076	4 consistent probesets
AI445461	tj34g07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-214	0.183	0.078	4 consistent probesets
L36151	Homo sapiens phosphatidylinositol 4-kinase mRNA, compl	0.183	0.078	4 consistent probesets
X92720	H.sapiens mRNA for phosphoenolpyruvate carboxykinase /	0.183	0.079	4 consistent probesets
AB029023	Homo sapiens mRNA for KIAA1100 protein, complete cds /	0.183	0.091	4 consistent probesets
Z23090	H.sapiens mRNA for 28 kDa heat shock protein /cds=(491	0.183	0.093	4 consistent probesets
D79206	Homo sapiens gene for ryudocan core protein, exon1-5,	0.183	0.097	4 consistent probesets
AF052162	Homo sapiens clone 24655 mRNA sequence /cds=UNKNO	0.183	0.098	4 consistent probesets
D43642	Human YL-1 mRNA for YL-1 protein (nuclear protein with	0.183	0.101	3 consistent probesets
U79569	Human no arches (nar) mRNA, complete cds /cds=(36,845)	0.183	0.104	4 consistent probesets
AB029034	Homo sapiens mRNA for KIAA1111 protein, partial cds /c	0.183	0.112	4 consistent probesets
U64197	Homo sapiens chemokine exodus-1 mRNA, complete cds /	0.183	0.118	4 consistent probesets
X54199	Human mRNA for GARS-AIRS-GART /cds=UNKNOWN /gb	0.183	0.119	3 consistent probesets
X63469	H.sapiens mRNA for transcription factor TFIIE beta /cd	0.183	0.124	4 consistent probesets
U90268	Human Krit1 mRNA, complete cds /cds=(25,1614) /gb=U90	0.183	0.127	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AB000449	Homo sapiens mRNA for VRK1, complete cds /cds=(75,126	0.183	0.131	4 consistent probesets
U53209	Human transformer-2 alpha (htra-2 alpha) mRNA, complet	0.183	0.141	4 consistent probesets
AJ001381	Homo sapiens incomplete cDNA for a mutated allele of a	0.183	0.141	4 consistent probesets
U80735	Homo sapiens CAGF28 mRNA, partial cds /cds=(0,2235) /g	0.183	0.144	4 consistent probesets
AL080192	Homo sapiens mRNA; cDNA DKFZp434B102 (from clone D	0.183	0.144	4 inconsistent probesets
AF069250	Homo sapiens okadaic acid-inducible phosphoprotein (OA	0.183	0.150	4 consistent probesets
U77665	Human RNaseP protein p30 (RPP30) mRNA, complete cds	0.183	0.157	4 consistent probesets
M74093	Human cyclin mRNA /cds=UNKNOWN /gb=M74093 /gi=80	0.183	0.159	3 consistent probesets
D11428	Homo sapiens mRNA for PMP-22(PAS-II/SR13/Gas-3), cor	0.183	0.167	3 consistent probesets
U63541	Human mRNA expressed in HC/HCC livers and MolT-4 pro	0.183	0.167	3 consistent probesets
AF035812	Homo sapiens dynein light intermediate chain 2 (LIC2)	0.183	0.169	3 consistent probesets
M80629	HUMCHED Human cdc2-related protein kinase (CH	0.183	0.194	4 consistent probesets
AI685944	tu38g02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-225	0.183	0.202	4 consistent probesets
AL049970	Homo sapiens mRNA; cDNA DKFZp564B102 (from clone D	0.183	0.217	4 consistent probesets
AL046649	DKFZp434O088_s1 Homo sapiens cDNA, 3 end /clone=Dk	0.183	0.225	3 consistent probesets
D16815	Homo sapiens mRNA for EAR-1r, complete cds /cds=(305,2	0.183	0.245	3 consistent probesets
U12471	HSU12471 Human thrombospondin-1 gene, partial	0.183	0.251	3 consistent probesets
U26455	HSU26455 Human phosphatidylinositol 3-kinase	0.183	0.255	3 consistent probesets
AI862521	wj15a06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-240	0.183	0.259	4 consistent probesets
AB020699	Homo sapiens mRNA for KIAA0892 protein, partial cds /c	0.183	0.263	3 consistent probesets
U90907	Human clone 23907 mRNA sequence /cds=UNKNOWN /gb	0.183	0.266	3 consistent probesets
AF025794	Homo sapiens methionine synthase reductase (MTRR) mRI	0.183	0.282	3 consistent probesets
AL050198	Homo sapiens mRNA; cDNA DKFZp586D0823 (from clone	0.183	0.282	2 consistent probesets
AJ132545	Homo sapiens mRNA for protein kinase /cds=(395,2062) /	0.183	0.302	3 consistent probesets
Y00636	Human mRNA for lymphocyte function associated antigen-	0.183	0.341	3 consistent probesets
Z75330	H.sapiens mRNA for nuclear protein SA-1 /cds=(400,4176	0.183	0.432	3 consistent probesets
AF005775	AF005775 Homo sapiens caspase-like apoptosis	0.181	0.058	8 consistent probesets
L20852	HUMGLVR2X Human leukemia virus receptor 2 (GL	0.181	0.149	8 consistent probesets
D87071	Human mRNA for KIAA0233 gene, complete cds /cds=(2,61	0.179	0.053	4 inconsistent probesets
AI375913	tc14c08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-206	0.179	0.066	4 inconsistent probesets
AL050265	Homo sapiens mRNA; cDNA DKFZp564O1716 (from clone	0.179	0.078	4 consistent probesets
U73682	HSU73682 Human meningioma-expressed antigen 1	0.179	0.078	4 consistent probesets
D61391	Human mRNA for phosphoribosypyrophosphate synthetase	0.179	0.079	4 consistent probesets
AB002313	Human mRNA for KIAA0315 gene, partial cds /cds=(0,5526	0.179	0.086	4 inconsistent probesets
AB002382	Human mRNA for KIAA0384 gene, complete cds /cds=(465	0.179	0.089	4 consistent probesets
U63824	Human transcription factor RTEF-1 (RTEF1) mRNA, compl	0.179	0.093	4 consistent probesets
AB020705	Homo sapiens mRNA for KIAA0898 protein, partial cds /c	0.179	0.104	4 consistent probesets
AF052138	Homo sapiens clone 23718 mRNA sequence /cds=UNKNO	0.179	0.104	4 consistent probesets
Y00285	HSIGFIIR Human mRNA for insuline-like growth	0.179	0.107	4 consistent probesets
AL050136	Homo sapiens mRNA; cDNA DKFZp586L141 (from clone D	0.179	0.108	4 inconsistent probesets
AF099989	Homo sapiens Ste-20 related kinase SPAK mRNA, complet	0.179	0.112	4 consistent probesets
U84388	HSU84388 Human death domain containing protei	0.179	0.112	4 inconsistent probesets
U72066	Homo sapiens CtBP interacting protein CtIP (CtIP) mRNA	0.179	0.122	4 consistent probesets
W26854	17b4 Homo sapiens cDNA /gb=W26854 /gi=1306217 /	0.179	0.122	4 consistent probesets
AF001628	Homo sapiens interactor protein AblBP4 (AblBP4) mRNA,	0.179	0.124	4 consistent probesets
AF013758	Homo sapiens polyadenylate binding protein-interacting	0.179	0.125	4 consistent probesets
AI039880	ox97c12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.179	0.127	4 consistent probesets
D38521	Human mRNA for KIAA0077 gene, partial cds /cds=(0,5397	0.179	0.128	4 consistent probesets
Z46973	H.sapiens mRNA for phosphatidylinositol 3-kinase /cds=	0.179	0.129	4 consistent probesets
L07758	Human IEF SSP 9502 mRNA, complete cds /cds=(87,1592	0.179	0.129	4 consistent probesets
U88047	Homo sapiens DNA binding protein homolog (DRIL1) mRNA	0.179	0.130	4 consistent probesets
AB017642	Homo sapiens mRNA for oxidative-stress responsive 1, c	0.179	0.130	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AB002332	Human mRNA for KIAA0334 gene, complete cds /cds=(251	0.179	0.135	4 consistent probesets
M27492	HUMIL1RA Human interleukin 1 receptor mRNA, c	0.179	0.137	4 consistent probesets
S73885	AP-4=basic helix-loop-helix DNA-binding protein [human	0.179	0.143	4 consistent probesets
AF002999	Homo sapiens TTAGGG repeat binding factor 2 (hTRF2) m	0.179	0.146	4 consistent probesets
AF022385	Homo sapiens apoptosis-related protein TFAR15 (TFAR15)	0.179	0.167	4 consistent probesets
D50929	Human mRNA for KIAA0139 gene, complete cds /cds=(128	0.179	0.200	4 consistent probesets
Y11681	Homo sapiens mRNA for mitochondrial ribosomal protein	0.177	0.056	8 inconsistent probesets
AI800578	wg12b07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.177	0.128	8 inconsistent probesets
D26070	HUMINSP3R1 Human mRNA for type 1 inositol 1,4	0.177	0.155	8 consistent probesets
AI057607	oy31e07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.176	0.155	7 consistent probesets
U07424	Human putative tRNA synthetase-like protein mRNA, comp	0.175	0.062	4 inconsistent probesets
M80261	HUMAPE Human apurinic endonuclease (APE) mRNA	0.175	0.062	4 consistent probesets
AL096752	Homo sapiens mRNA; cDNA DKFZp434A012 (from clone D	0.175	0.082	4 consistent probesets
W28264	44c9 Homo sapiens cDNA /gb=W28264 /gi=1308212 /	0.175	0.083	4 consistent probesets
AF061741	Homo sapiens retinal short-chain dehydrogenase/reducta	0.175	0.085	4 consistent probesets
W27050	19f7 Homo sapiens cDNA /gb=W27050 /gi=1306422 /	0.175	0.085	4 consistent probesets
U90909	Human clone 23722 mRNA sequence /cds=UNKNOWN /gb	0.175	0.086	4 inconsistent probesets
AI743507	wf72a06.x2 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.175	0.090	4 consistent probesets
AI140114	qa95c06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.175	0.091	4 consistent probesets
AB023171	Homo sapiens mRNA for KIAA0954 protein, partial cds /c	0.175	0.094	4 consistent probesets
U85245	Human phosphatidylinositol-4-phosphate 5-kinase type I	0.175	0.096	4 consistent probesets
AB007916	Homo sapiens mRNA for KIAA0447 protein, complete cds /	0.175	0.102	4 consistent probesets
AF000152	Homo sapiens OS-4 protein (OS-4) mRNA, complete cds /c	0.175	0.104	4 consistent probesets
U66685	HSU66685 Homo sapiens cDNA /gb=U66685 /gi=1906570 /	0.175	0.107	2 consistent probesets
L04733	Homo sapiens kinesin light chain mRNA, complete cds /c	0.175	0.107	4 consistent probesets
U89896	HSU89896 Homo sapiens casein kinase I gamma 2	0.175	0.108	8 consistent probesets
U79287	Human clone 23867 mRNA sequence /cds=UNKNOWN /gb	0.175	0.109	4 consistent probesets
D43948	Human mRNA for KIAA0097 gene, complete cds /cds=(47,6	0.175	0.110	4 consistent probesets
AB029032	Homo sapiens mRNA for KIAA1109 protein, partial cds /c	0.175	0.115	4 consistent probesets
AF049105	Homo sapiens centrosomal Nek2-associated protein 1 (C-	0.175	0.126	2 consistent probesets
D87953	Human mRNA for RTP, complete cds /cds=(122,1306) /gb=	0.175	0.127	4 consistent probesets
D78132	D78132 Homo sapiens mRNA for ras-related GTP-	0.175	0.128	4 consistent probesets
AF007871	Homo sapiens torsinA (DYT1) mRNA, complete cds /cds=(4	0.175	0.132	4 consistent probesets
AB019036	Homo sapiens mRNA for geranylgeranyl pyrophosphate syn	0.175	0.142	4 consistent probesets
AL021396	Human DNA sequence from clone 971N18 on chromosome	0.175	0.145	4 inconsistent probesets
D88357	Homo sapiens mRNA for CDC2 delta T, complete cds /cds=	0.175	0.147	4 consistent probesets
AB009356	Homo sapiens mRNA for TGF-beta activated kinase 1a, co	0.175	0.170	4 inconsistent probesets
L05624	HUMMKK Homo sapiens MAP kinase kinase mRNA, c	0.175	0.175	4 consistent probesets
AF026445	Homo sapiens cofactor of initiator function (CIF150) m	0.175	0.175	4 consistent probesets
U83661	HSU83661 Homo sapiens multidrug resistance pr	0.175	0.178	2 consistent probesets
U71601	Human zinc finger protein zfp47 (zf47) mRNA, partial c	0.175	0.178	2 consistent probesets
U07681	Human NAD(H)-specific isocitrate dehydrogenase alpha s	0.175	0.181	4 consistent probesets
AL049942	Homo sapiens mRNA; cDNA DKFZp564F1422 (from clone	0.175	0.182	4 consistent probesets
X74496	H.sapiens mRNA for prolyl oligopeptidase /cds=(0,2132)	0.175	0.199	4 consistent probesets
L42243	HUMIFNAM08 Homo sapiens (clone 51H8) alternat	0.175	0.203	6 consistent probesets
X54936	HSPLGF H.sapiens mRNA for placenta growth fac	0.175	0.232	2 consistent probesets
AB014610	Homo sapiens mRNA for KIAA0710 protein, complete cds /	0.175	0.316	2 consistent probesets
AF019083	Homo sapiens phosphatase and tensin homolog 2 (PTH2) n	0.175	0.321	2 consistent probesets
Y14494	Homo sapiens mRNA for mitochondrial carrier protein AR	0.175	0.323	2 consistent probesets
Z80780	H.sapiens H2B/h gene /cds=(0,380) /gb=Z80780 /gi=15685	0.175	0.329	2 consistent probesets
AF041081	Homo sapiens D15F37 pseudogene, S4 allele, mRNA sequ	0.175	0.339	2 consistent probesets
W28567	51e2 Homo sapiens cDNA /gb=W28567 /gi=1308722 /	0.175	0.343	2 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D63480	Human mRNA for KIAA0146 gene, partial cds /cds=(0,2756	0.175	0.350	2 consistent probesets
U64871	HSU64871 Human putative G protein-coupled rec	0.175	0.358	2 consistent probesets
AB011132	Homo sapiens mRNA for KIAA0560 protein, complete cds /	0.175	0.519	2 consistent probesets
L07077	Human enyol-CoA- hydratase 3-hydroxyacyl-CoA dehydrog	0.175	0.576	2 consistent probesets
L19067	HUMNFKB65A Human NF-kappa-B transcription fac	0.174	0.060	12 inconsistent probesets
M60618	Human nuclear autoantigen (SP-100) mRNA, complete cds	0.172	0.163	6 consistent probesets
AF037448	Homo sapiens RRM RNA binding protein Gry-rbp (GRY-RB	0.171	0.066	4 inconsistent probesets
AF052113	Homo sapiens clone 23675 mRNA sequence /cds=UNKNO	0.171	0.078	4 consistent probesets
AB020724	Homo sapiens mRNA for KIAA0917 protein, partial cds /c	0.171	0.081	4 consistent probesets
AL041663	DKFZp434M0217_s1 Homo sapiens cDNA, 3 end /clone=D	0.171	0.086	4 consistent probesets
AB011105	Homo sapiens mRNA for KIAA0533 protein, partial cds /c	0.171	0.086	4 consistent probesets
AF108145	Homo sapiens MYLE mRNA, complete cds /cds=(52,258) /g	0.171	0.090	4 consistent probesets
AF052135	Homo sapiens clone 23625 mRNA sequence /cds=UNKNO	0.171	0.098	4 consistent probesets
M55630	Human topoisomerase I pseudogene 2 /cds=UNKNOWN /g	0.171	0.098	4 consistent probesets
AF042166	Homo sapiens beta-filamin mRNA, complete cds /cds=(131	0.171	0.101	4 consistent probesets
AB007945	Homo sapiens mRNA for KIAA0476 protein, complete cds /	0.171	0.101	4 consistent probesets
AF093670	Homo sapiens peroxisomal biogenesis factor (PEX11b) mR	0.171	0.102	4 consistent probesets
M34175	Human beta adaptin mRNA, complete cds /cds=(177,2990)	0.171	0.102	4 inconsistent probesets
X16354	HSTM1CEA Human mRNA for transmembrane carcino	0.171	0.104	4 consistent probesets
U52522	Human arfaptin 2, putative target protein of ADP-ribos	0.171	0.110	4 inconsistent probesets
D86966	Human mRNA for KIAA0211 gene, complete cds /cds=(570	0.171	0.113	4 consistent probesets
Y13350	Homo sapiens mRNA for DnaJ protein /cds=(0,975) /gb=Y1	0.171	0.113	4 consistent probesets
L11284	HUMMEK1NF Homosapiens ERK activator kinase (M	0.171	0.116	4 consistent probesets
AL050192	Homo sapiens mRNA; cDNA DKFZp586C1723 (from clone	0.171	0.121	4 consistent probesets
X90999	H.sapiens mRNA for Glyoxalase II /cds=(36,818) /gb=X90	0.171	0.122	4 consistent probesets
U57650	HSU57650 Human SH2-containing inositol 5-phos	0.171	0.124	4 consistent probesets
L27476	Human X104 mRNA, complete cds /cds=(79,3429) /gb=L27	0.171	0.128	4 consistent probesets
D85131	D85131 Homo sapiens mRNA for Myc-associated z	0.171	0.129	4 inconsistent probesets
AF003938	Homo sapiens thioredoxin-like protein mRNA, complete c	0.171	0.130	4 consistent probesets
AF089750	Homo sapiens flotillin-1 mRNA, complete cds /cds=(164,	0.171	0.131	4 consistent probesets
X74330	HSPRIM1 H.sapiens mRNA for DNA primase (subun	0.171	0.132	4 consistent probesets
AB002328	Human mRNA for KIAA0330 gene, partial cds /cds=(0,5710	0.171	0.135	4 consistent probesets
Y09765	Homo sapiens mRNA for putative GABA receptor epsilon s	0.171	0.144	4 consistent probesets
AB015633	Homo sapiens mRNA for type II membrane protein, comple	0.171	0.155	4 consistent probesets
AF101434	Homo sapiens Wolf-Hirschhorn syndrome candidate 2 prot	0.171	0.167	4 consistent probesets
AA453183	zx46b09.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-795	0.171	0.176	4 inconsistent probesets
W28948	54b12 Homo sapiens cDNA /gb=W28948 /gi=1308896 /	0.171	0.177	4 consistent probesets
N58115	yv65a01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.171	0.184	4 consistent probesets
AB018306	Homo sapiens mRNA for KIAA0763 protein, complete cds /	0.171	0.195	4 consistent probesets
AB023166	Homo sapiens mRNA for KIAA0949 protein, partial cds /c	0.171	0.231	4 consistent probesets
AI566877	tn24f02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-216	0.171	0.233	4 consistent probesets
U77129	Human SPS1/STE20 homolog KHS1 mRNA, complete cds	0.171	0.265	4 consistent probesets
X98296	HSUBIQHYD H.sapiens mRNA for ubiquitin hydrol	0.170	0.098	11 consistent probesets
L08069	HUMDNAJHOM Human heat shock protein, E. coli	0.169	0.071	8 inconsistent probesets
Z35102	HSPROKINX H.sapiens mRNA for Ndr protein kina	0.169	0.161	8 consistent probesets
N36842	yy35g03.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-27	0.167	0.053	4 consistent probesets
AL050254	Novel human gene mapping to chomosome 22 /cds=(205,1	0.167	0.064	4 consistent probesets
AJ005259	Homo sapiens mRNA for EDF-1 protein /cds=(34,480) /gb=	0.167	0.075	4 consistent probesets
AL050353	Homo sapiens mRNA; cDNA DKFZp547C0410 (from clone	0.167	0.085	4 consistent probesets
L40401	Homo sapiens (clone zap128) mRNA, 3 end of cds /cds=(	0.167	0.086	4 consistent probesets
M60483	HUMPP2AA Human protein phosphatase 2A catalyt	0.167	0.087	4 consistent probesets
J05032	Human aspartyl-tRNA synthetase alpha-2 subunit mRNA, c	0.167	0.093	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
L20817	Homo sapiens tyrosine protein kinase (CAK) gene, compl	0.167	0.093	4 consistent probesets
W28167	43a1 Homo sapiens cDNA /gb=W28167 /gi=1308115 /	0.167	0.095	4 consistent probesets
X64228	H.sapiens can mRNA /cds=(94,6366) /gb=X64228 /gi=2965	0.167	0.098	3 consistent probesets
D88153	Homo sapiens mRNA for HYA22, complete cds /cds=(147,1	0.167	0.098	4 consistent probesets
AF072860	Homo sapiens protein activator of the interferon-induc	0.167	0.098	4 consistent probesets
X72889	H.sapiens hbrm mRNA /cds=(222,4982) /gb=X72889 /gi=41	0.167	0.104	4 consistent probesets
AF084260	Homo sapiens signalosome subunit 2 (SGN2) mRNA, comp	0.167	0.107	4 consistent probesets
AF027957	Homo sapiens G protein-coupled receptor (GPR35) gene,	0.167	0.109	4 consistent probesets
AA149307	zl25h05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-503	0.167	0.111	4 consistent probesets
M15990	HUMCYES1 Human c-yes-1 mRNA	0.167	0.112	4 consistent probesets
L35253	HUMMAPKNS Human p38 mitogen activated protein	0.167	0.119	4 inconsistent probesets
D50640	D50625S16 Homo sapiens DNA for phosphodiester	0.167	0.120	8 consistent probesets
U78733	HSSMAD2S8 Homo sapiens mad protein homolog Sm	0.167	0.121	4 consistent probesets
AL050022	Homo sapiens mRNA; cDNA DKFZp564D116 (from clone D	0.167	0.121	4 consistent probesets
D86972	Human mRNA for KIAA0218 gene, complete cds /cds=(398	0.167	0.122	3 inconsistent probesets
D38498	HUMPMS1A Human PMS5 mRNA (yeast mismatch repa	0.167	0.126	8 consistent probesets
AB020880	Homo sapiens mRNA for squamous cell carcinoma antigen	0.167	0.127	4 consistent probesets
AL049417	Homo sapiens mRNA; cDNA DKFZp586O1919 (from clone	0.167	0.129	4 consistent probesets
AL031588	dJ1163J1.3 (novel protein similar to mouse B99) /cds=(	0.167	0.131	4 consistent probesets
AB029014	Homo sapiens mRNA for KIAA1091 protein, partial cds /c	0.167	0.131	4 consistent probesets
U81554	Homo sapiens CaM kinase II isoform mRNA, complete cds	0.167	0.136	4 consistent probesets
AJ222967	Homo sapiens mRNA for cystinosin /cds=(339,1442) /gb=A	0.167	0.136	3 consistent probesets
AI133727	Habcs0217 Homo sapiens cDNA /gb=AI133727 /gi=360292	0.167	0.142	4 consistent probesets
AF072242	Homo sapiens methyl-CpG binding protein MBD2 (MBD2) r	0.167	0.145	4 consistent probesets
L38928	Homo sapiens 5,10-methenyltetrahydrofolate synthetase	0.167	0.150	4 consistent probesets
AB006679	AB006679 Homo sapiens mRNA for ATP binding pr	0.167	0.153	4 consistent probesets
U00946	Human clone A9A2BRB5 (CAC)n/(GTG)n repeat-containing	0.167	0.154	4 consistent probesets
AF068868	Homo sapiens TNFR-related death receptor-6 (DR6) mRNA	0.167	0.158	3 consistent probesets
L34075	HUMFRAPX Human FKBP-rapamycin associated prot	0.167	0.158	3 consistent probesets
D50912	Human mRNA for KIAA0122 gene, partial cds /cds=(0,3033	0.167	0.165	3 consistent probesets
AB007962	Homo sapiens mRNA, chromosome 1 specific transcript KI	0.167	0.167	3 consistent probesets
M95809	Human basic transcription factor 62kD subunit (BTF2),	0.167	0.173	4 consistent probesets
L10347	Human pro-alpha1 type II collagen (COL2A1) gene exons	0.167	0.176	3 consistent probesets
AF070598	Homo sapiens clone 24410 ABC transporter mRNA, partial	0.167	0.176	3 consistent probesets
D42054	Human mRNA for KIAA0092 gene, complete cds /cds=(53,1	0.167	0.179	4 consistent probesets
AL050274	Homo sapiens mRNA; cDNA DKFZp566C243 (from clone D	0.167	0.188	3 consistent probesets
L38487	HUMHERRA1 Human estrogen receptor-related pro	0.167	0.196	3 consistent probesets
AJ010953	Homo sapiens mRNA for putative Ca2+-transporting ATPas	0.167	0.205	3 consistent probesets
AB011540	Homo sapiens mRNA for MEGF7, partial cds /cds=(0,4730)	0.167	0.215	4 consistent probesets
W27761	37c5 Homo sapiens cDNA /gb=W27761 /gi=1307709 /	0.167	0.215	4 consistent probesets
AB002303	Human mRNA for KIAA0305 gene, complete cds /cds=(248	0.167	0.217	4 consistent probesets
U80034	Human mitochondrial intermediate peptidase precursor (	0.167	0.221	3 consistent probesets
AB011104	Homo sapiens mRNA for KIAA0532 protein, partial cds /c	0.167	0.223	3 consistent probesets
U81802	HSU81802 Human PtdIns 4-kinase (PI4Kb) mRNA,	0.167	0.233	3 consistent probesets
AL049783	Novel human gene mapping to chomosome 13 /cds=(165,1	0.167	0.245	3 consistent probesets
W25793	13e7 Homo sapiens cDNA /gb=W25793 /gi=1305934 /	0.167	0.246	4 consistent probesets
AF084367	Homo sapiens inversin protein mRNA, complete cds /cds=	0.167	0.248	2 consistent probesets
AA873858	oh79b10.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-14	0.167	0.253	4 consistent probesets
AL050378	Homo sapiens mRNA; cDNA DKFZp586I1420 (from clone D	0.167	0.262	3 consistent probesets
AB007954	Homo sapiens mRNA, chromosome 1 specific transcript KI	0.167	0.273	4 consistent probesets
AL050151	Homo sapiens mRNA; cDNA DKFZp586J0720 (from clone	0.167	0.288	3 consistent probesets
AF038362	Homo sapiens TBP-associated factor 172 (TAF-172) mRNA	0.167	0.292	3 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AJ223948	Homo sapiens mRNA for putative RNA helicase, 3 end /c	0.167	0.296	4 consistent probesets
M94065	Human dihydroorotate dehydrogenase mRNA, 3 end /cds=	0.167	0.305	3 consistent probesets
AB020663	Homo sapiens mRNA for KIAA0856 protein, partial cds /c	0.167	0.331	3 consistent probesets
D85939	Homo sapiens mRNA for p97 homologous protein, complet	0.167	0.338	2 consistent probesets
AB011128	Homo sapiens mRNA for KIAA0556 protein, partial cds /c	0.167	0.358	3 consistent probesets
AB018324	Homo sapiens mRNA for KIAA0781 protein, partial cds /c	0.167	0.409	3 consistent probesets
AI955897	wt31b09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.167	0.432	3 consistent probesets
AL031985	Human DNA sequence from clone 228H13 on chromosome	0.167	0.491	3 consistent probesets
AF014837	Homo sapiens m6A methyltransferase (MT-A70) gene, com	0.165	0.087	8 inconsistent probesets
U16799	Human Na,K-ATPase beta-1 subunit mRNA, complete cds /	0.163	0.062	4 consistent probesets
U00968	Human SREBP-1 mRNA, complete cds /cds=(166,3609) /gb	0.163	0.081	4 consistent probesets
AI742846	wg46h09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.163	0.081	4 inconsistent probesets
D42084	Human mRNA for KIAA0094 gene, partial cds /cds=(0,1185	0.163	0.083	4 consistent probesets
AL049397	Homo sapiens mRNA; cDNA DKFZp586C1019 (from clone	0.163	0.091	4 consistent probesets
X52104	Human mRNA for p68 protein /cds=(175,2019) /gb=X52104	0.163	0.092	4 consistent probesets
Y08110	H.sapiens mRNA for mosaic protein LR11 /cds=(80,6724)	0.163	0.111	4 inconsistent probesets
U39400	Human NOF1 mRNA, complete cds /cds=(13,513) /gb=U39	0.163	0.111	4 consistent probesets
X53390	Human mRNA for upstream binding factor (hUBF) /cds=(14	0.163	0.116	4 consistent probesets
AB007934	Homo sapiens mRNA for KIAA0465 protein, partial cds /c	0.163	0.119	4 inconsistent probesets
X81198	H.sapiens mRNA (clone p5) for archain /cds=(78,1496) /	0.163	0.127	4 consistent probesets
AF039022	Homo sapiens exportin t mRNA, complete cds /cds=(0,288	0.163	0.133	4 consistent probesets
M73812	HUMCYCE Human cyclin E mRNA sequence	0.163	0.145	4 consistent probesets
D26069	Human mRNA for KIAA0041 gene, partial cds /cds=(0,1899	0.163	0.147	4 inconsistent probesets
AB007851	Homo sapiens mRNA for 41-kDa phosphoribosylpyrophospl	0.163	0.154	4 consistent probesets
AB028965	Homo sapiens mRNA for KIAA1042 protein, complete cds /	0.163	0.164	4 inconsistent probesets
AI057115	oz23g10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.163	0.200	4 consistent probesets
U72507	Human 40871 mRNA partial sequence /cds=UNKNOWN /g	0.163	0.231	4 consistent probesets
AI687419	tp95h03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-220	0.163	0.267	4 inconsistent probesets
M86917	Human oxysterol-binding protein (OSBP) mRNA, complete	0.163	0.393	4 consistent probesets
U77456	Human nucleosome assembly protein 2 mRNA, complete c	0.162	0.078	4 consistent probesets
D73409	Homo sapiens mRNA for diacylglycerol kinase delta, com	0.162	0.087	4 inconsistent probesets
AI827895	wf12b02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.162	0.094	4 consistent probesets
N25117	yx19c09.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-262	0.162	0.102	4 inconsistent probesets
Y07661	H.sapiens USF2 gene /cds=(0,1040) /gb=Y07661 /gi=18060	0.162	0.107	4 inconsistent probesets
Z15108	HSPKCZ H.sapiens mRNA for protein kinase C ze	0.162	0.108	4 consistent probesets
M55268	Human casein kinase II alpha subunit mRNA, complete cd	0.162	0.113	4 consistent probesets
D86978	Human mRNA for KIAA0225 gene, partial cds /cds=(0,6043	0.162	0.130	4 consistent probesets
AF045451	Homo sapiens transcriptional regulatory protein p54 mR	0.162	0.135	4 consistent probesets
L06845	Human cysteinyl-tRNA synthetase mRNA, partial cds /cds	0.162	0.143	4 consistent probesets
Y09048	H.sapiens PxF gene /cds=(10,909) /gb=Y09048 /gi=257002	0.162	0.146	4 consistent probesets
AF043325	Homo sapiens N-myristoyltransferase 2 mRNA, complete c	0.162	0.158	4 consistent probesets
U77970	Human neuronal PAS2 (NPAS2) mRNA, complete cds /cds	0.162	0.162	4 consistent probesets
X91648	H.sapiens mRNA for pur alpha extended 3untranslated re	0.162	0.199	4 consistent probesets
AB002438	Homo sapiens mRNA from chromosome 5q21-22, clone-FB	0.162	0.231	4 consistent probesets
AI263885	qi08d08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-185	0.162	0.246	4 consistent probesets
U88666	HSU88666 Homo sapiens serine kinase SRPK2 mRN	0.162	0.152	7 consistent probesets
X75940	H.sapiens beta glucuronidase pseudogene /cds=UNKNOWI	0.161	0.226	3 consistent probesets
U01062	HUMIP3R3 Human type 3 inositol 1,4,5-trisphos	0.160	0.063	8 inconsistent probesets
L22005	HUMCDC34H Human ubiquitin conjugating enzyme	0.160	0.084	5 consistent probesets
AB011173	Homo sapiens mRNA for KIAA0601 protein, partial cds /c	0.158	0.085	4 consistent probesets
AL049265	Homo sapiens mRNA; cDNA DKFZp564F053 (from clone D	0.158	0.087	4 inconsistent probesets
AI983043	wz30b11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.158	0.087	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AA524345	ng43a06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-93	0.158	0.088	4 consistent probesets
AB011153	Homo sapiens mRNA for KIAA0581 protein, partial cds /c	0.158	0.095	4 consistent probesets
X82103	H.sapiens mRNA for beta-COP /cds=(0,911) /gb=X82103 /g	0.158	0.097	4 consistent probesets
U70451	Human myeloid differentiation primary response protein	0.158	0.098	4 consistent probesets
L00634	HUMFPTA Human farnesyl-protein transferase al	0.158	0.100	4 consistent probesets
D28476	Human mRNA for KIAA0045 gene, complete cds /cds=(109	0.158	0.100	4 consistent probesets
AF086947	untitled /cds=(334,4119) /gb=AF086947 /gi=4139120 /	0.158	0.102	4 consistent probesets
W47047	zc38g10.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-324	0.158	0.102	4 consistent probesets
S59184	S59184 RYK=related to receptor tyrosine kinas	0.158	0.114	6 inconsistent probesets
U93237	HSU93237 Human menin (MEN1) gene, complete cd	0.158	0.116	4 consistent probesets
D21064	Human mRNA for KIAA0123 gene, partial cds /cds=(0,1586	0.158	0.116	4 consistent probesets
AF063605	Homo sapiens brain my047 protein mRNA, complete cds /c	0.158	0.118	4 consistent probesets
AB000450	Homo sapiens mRNA for VRK2, complete cds /cds=(130,16	0.158	0.121	4 consistent probesets
AB011175	Homo sapiens mRNA for KIAA0603 protein, complete cds /	0.158	0.132	4 consistent probesets
J04027	Human plasma membrane Ca2+ pumping ATPase mRNA,	0.158	0.143	4 consistent probesets
L76259	Homo sapiens PTS gene, complete cds /cds=(68,505) /gb=	0.158	0.161	4 consistent probesets
L36140	Homo sapiens (clone 1311) DNA helicase (RECQL) mRNA,	0.158	0.179	4 inconsistent probesets
U94905	Human diacylglycerol kinase zeta mRNA, alternatively s	0.158	0.186	4 consistent probesets
AL050272	Homo sapiens mRNA; cDNA DKFZp566B183 (from clone D	0.158	0.189	4 consistent probesets
AB022918	Homo sapiens mRNA for alpha2,3-sialyltransferase ST3Ga	0.158	0.205	4 consistent probesets
X55330	H.sapiens mRNA for aspartylglucosaminidase /cds=(170,1	0.158	0.214	4 consistent probesets
AB028951	Homo sapiens mRNA for KIAA1028 protein, partial cds /c	0.158	0.215	4 consistent probesets
X76388	H.sapiens mRNA for 2-5A binding protein /cds=(117,1916	0.158	0.254	4 consistent probesets
U29171	HSU29171 Human casein kinase I delta mRNA, co	0.156	0.067	8 inconsistent probesets
U72936	HSU72936 Homo sapiens putative DNA dependent	0.156	0.093	9 consistent probesets
AB007447	Homo sapiens mRNA for Fln29, complete cds /cds=(54,180	0.156	0.164	3 consistent probesets
AB014888	Homo sapiens mRNA for MRJ, complete cds /cds=(100,825	0.154	0.061	4 inconsistent probesets
M61764	Human gamma-tubulin mRNA, complete cds /cds=(24,1379	0.154	0.067	4 inconsistent probesets
AF035292	Homo sapiens clone 23584 mRNA sequence /cds=UNKNO	0.154	0.070	8 consistent probesets
U11861	Human G10 homolog (edg-2) mRNA, complete cds /cds=(3	0.154	0.075	4 consistent probesets
AB002405	Homo sapiens mRNA for LAK-4p, complete cds /cds=(109,1	0.154	0.085	4 consistent probesets
D86977	Human mRNA for KIAA0224 gene, complete cds /cds=(136	0.154	0.088	4 consistent probesets
AB018264	Homo sapiens mRNA for KIAA0721 protein, partial cds /c	0.154	0.089	4 consistent probesets
AA160724	zo72h04.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-592	0.154	0.089	4 inconsistent probesets
Y08612	Homo sapiens mRNA for nuclear pore complex protein 88	0.154	0.091	4 consistent probesets
AL080146	Homo sapiens mRNA; cDNA DKFZp434B174 (from clone D	0.154	0.091	4 consistent probesets
AJ245416	Homo sapiens mRNA for G7b protein (G7b gene, located i	0.154	0.096	4 consistent probesets
D88435	Homo sapiens mRNA for HsGAK, complete cds /cds=(0,393	0.154	0.097	4 consistent probesets
L37042	Homo sapiens casein kinase I alpha isoform (CSNK1A1) m	0.154	0.102	4 consistent probesets
M19650	HUMCNPDEA Human 2 ,3 -cyclic nucleotide 3 -ph	0.154	0.103	4 consistent probesets
AL050197	Homo sapiens mRNA; cDNA DKFZp586D0623 (from clone	0.154	0.107	4 consistent probesets
D80005	Human mRNA for KIAA0183 gene, partial cds /cds=(0,3190	0.154	0.116	4 consistent probesets
D50525	Human mRNA for TI-227H /cds=UNKNOWN /gb=D50525 /(	0.154	0.117	4 consistent probesets
D14661	Human mRNA for KIAA0105 gene, complete cds /cds=(124	0.154	0.120	4 consistent probesets
D42085	Human mRNA for KIAA0095 gene, complete cds /cds=(66,2	0.154	0.124	4 consistent probesets
AF041248	Homo sapiens cyclin-dependent kinase inhibitor (CDKN2C	0.154	0.127	4 consistent probesets
U09477	HSU09477 Human clone 53BP1 p53-binding protei	0.154	0.132	4 consistent probesets
AB023175	Homo sapiens mRNA for KIAA0958 protein, partial cds /c	0.154	0.136	4 consistent probesets
AL050062	Homo sapiens mRNA; cDNA DKFZp566K023 (from clone D	0.154	0.144	4 consistent probesets
U38847	Human TAR RNA loop binding protein (TRP-185) mRNA, c	0.154	0.145	4 consistent probesets
AF029669	Homo sapiens Rad51C (RAD51C) mRNA, complete cds /co	0.154	0.146	4 consistent probesets
U41060	HSU41060 Human breast cancer, estrogen regula	0.154	0.149	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
X12654	Human mRNA for cell cycle gene RCC1 /cds=(182,1447) /g	0.154	0.149	4 consistent probesets
X70340	HSTGFAA H.sapiens mRNA for transforming growt	0.154	0.154	4 consistent probesets
AL050269	Homo sapiens mRNA; cDNA DKFZp564C103 (from clone D	0.154	0.158	4 consistent probesets
AF082557	Homo sapiens TRF1-interacting ankyrin-related ADP-ribo	0.154	0.201	4 consistent probesets
AB014539	Homo sapiens mRNA for KIAA0639 protein, partial cds /c	0.154	0.215	4 consistent probesets
U02020	Human pre-B cell enhancing factor (PBEF) mRNA, complet	0.154	0.216	4 consistent probesets
AL031177	dJ889N15.2.1 (26S Proteasome subunit p28 (Ankyrin repe	0.154	0.225	4 consistent probesets
J04605	Human prolidase (imidodipeptidase) mRNA, complete cds	0.154	0.255	4 consistent probesets
X91906	H.sapiens voltage-gated chloride ion channel CLCN5 /cd	0.154	0.350	4 consistent probesets
D64142	D64142 Human mRNA for histone H1x, complete c	0.152	0.075	8 consistent probesets
U40705	HSU40705 Homo sapiens telomeric repeat bindin	0.151	0.176	13 consistent probesets
X13794	H.sapiens lactate dehydrogenase B gene exon 1 and 2 (E	0.150	0.053	4 inconsistent probesets
X12794	HSEAR2 Human v-erbA related ear-2 gene	0.150	0.059	7 inconsistent probesets
AL031178	Human DNA sequence from clone 341E18 on chromosome	0.150	0.070	4 consistent probesets
U82938	Human CD27BP (Siva) mRNA, complete cds /cds=(252,821	0.150	0.072	4 consistent probesets
M15796	HUMCYL Human cyclin protein gene, complete cd	0.150	0.075	4 consistent probesets
AI920820	wn82e10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.150	0.078	4 consistent probesets
X91148	H.sapiens mRNA for microsomal triglyceride transfer pr	0.150	0.081	4 consistent probesets
D84110	D84110 Homo sapiens mRNA for RBP-MS/type 4, c	0.150	0.083	8 consistent probesets
J05428	Human 3,4-catechol estrogen UDP-glucuronosyltransferas	0.150	0.086	8 consistent probesets
U78525	Homo sapiens eukaryotic translation initiation factor	0.150	0.087	4 consistent probesets
X79888	H.sapiens AUH mRNA /cds=(4,1023) /gb=X79888 /gi=7802	0.150	0.087	4 consistent probesets
AA192359	zp91c12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-627	0.150	0.090	3 consistent probesets
U39317	HSU39317 Human E2 ubiquitin conjugating enzym	0.150	0.095	4 consistent probesets
Y00371	Human hsc70 gene for 71 kd heat shock cognate protein	0.150	0.096	4 consistent probesets
U46570	Human tetratricopeptide repeat protein (tpr1) mRNA, co	0.150	0.098	4 consistent probesets
AL050172	Homo sapiens mRNA; cDNA DKFZp586F1322 (from clone	0.150	0.098	4 consistent probesets
X76228	H.sapiens mRNA for vacuolar H+ ATPase E subunit /cds=(	0.150	0.098	4 consistent probesets
U65928	HSU65928 Human Jun activation domain binding	0.150	0.100	4 consistent probesets
AF069735	Homo sapiens PCAF associated factor 65 alpha mRNA, cor	0.150	0.101	1 consistent probesets
M96233	HUMGSTM4A Human glutathione transferase class	0.150	0.101	1 consistent probesets
X63422	H.sapiens mRNA for delta-subunit of mitochondrial F1F0	0.150	0.101	1 consistent probesets
U44755	Human PSE-binding factor PTF delta subunit mRNA, comp	0.150	0.101	1 consistent probesets
D50663	D50663 Human mRNA for TCTEL1 gene, complete c	0.150	0.102	4 consistent probesets
AL050060	Homo sapiens mRNA; cDNA DKFZp566H073 (from clone D	0.150	0.106	4 consistent probesets
AF038186	Homo sapiens clone 23914 mRNA sequence /cds=UNKNO	0.150	0.108	4 consistent probesets
D55716	HUMP1CDC47 Human mRNA for P1cdc47, complete c	0.150	0.111	4 consistent probesets
AF055030	Homo sapiens clone 24538 mRNA sequence /cds=UNKNO	0.150	0.112	4 consistent probesets
AB018274	Homo sapiens mRNA for KIAA0731 protein, partial cds /c	0.150	0.116	4 consistent probesets
D11094	Human mRNA for MSS1, complete cds /cds=(66,1367) /gb=	0.150	0.119	4 consistent probesets
X06272	Human mRNA for docking protein (signal recognition par	0.150	0.122	4 consistent probesets
D83778	Human mRNA for KIAA0194 gene, partial cds /cds=(0,4309	0.150	0.123	3 inconsistent probesets
U18291	Human CDC16Hs mRNA, complete cds /cds=(24,1883) /gb	0.150	0.136	4 consistent probesets
AI540318	tq34f03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-221	0.150	0.142	4 consistent probesets
X15525	H.sapiens lysosomal acid phosphatase gene (EC 3.1.3.2)	0.150	0.143	2 consistent probesets
M74905	Human 3-alkyladenine DNA glycosylase (HAAG) mRNA, co	0.150	0.143	2 consistent probesets
AF016371	Homo sapiens U-snRNP-associated cyclophilin (USA-CyP)	0.150	0.145	3 consistent probesets
AF064243	Homo sapiens intersectin short form mRNA, complete cds	0.150	0.151	4 consistent probesets
Z96932	Homo sapiens mRNA for NA14 protein /cds=(46,405) /gb=Z	0.150	0.151	1 consistent probesets
D86331	D86331 Human MT2-MMP gene for matrix metallop	0.150	0.151	1 consistent probesets
D87682	Human mRNA for KIAA0241 gene, partial cds /cds=(0,1568	0.150	0.151	1 consistent probesets
W22655	71B9 Homo sapiens cDNA /clone=(not-directional) /gb=W2	0.150	0.151	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AL050100	Homo sapiens mRNA; cDNA DKFZp586D0919 (from clone	0.150	0.154	4 consistent probesets
D00723	Homo sapiens mRNA for hydrogen carrier protein, a comp	0.150	0.158	4 consistent probesets
AF026086	Homo sapiens peroxisome biogenesis disorder protein 1	0.150	0.161	2 consistent probesets
D25248	Homo sapiens mRNA, clone-RES4-4 /cds=UNKNOWN /gb=	0.150	0.166	3 consistent probesets
L20826	Human I-plastin mRNA, complete cds /cds=(97,1986) /gb=	0.150	0.167	4 consistent probesets
AL050147	Homo sapiens mRNA; cDNA DKFZp586E0820 (from clone	0.150	0.176	4 consistent probesets
AB007929	Homo sapiens mRNA for KIAA0460 protein, partial cds /c	0.150	0.176	3 consistent probesets
AB002390	Human mRNA for KIAA0392 gene, partial cds /cds=(0,1652	0.150	0.178	2 consistent probesets
L24559	HUMDNSPOLA Homo sapiens DNA polymerase alpha	0.150	0.182	2 consistent probesets
AF031166	Homo sapiens SRp46 splicing factor retropseudogene mRN	0.150	0.183	4 consistent probesets
U49082	Human transporter protein (g17) mRNA, complete cds /cd	0.150	0.186	3 consistent probesets
AJ006268	Homo sapiens mRNA for putative ATPase, partial /cds=(0	0.150	0.186	3 consistent probesets
U60325	HSU60325 Human DNA polymerase gamma mRNA, nuc	0.150	0.193	2 consistent probesets
U46461	HSU46461 Human dishevelled homolog (DVL) mRNA	0.150	0.202	1 consistent probesets
X63368	H.sapiens HSJ1 mRNA /cds=(25,1080) /gb=X63368 /gi=324	0.150	0.202	1 consistent probesets
AF055917	Homo sapiens protease-activated receptor 4 mRNA, compl	0.150	0.202	1 consistent probesets
AF026548	Homo sapiens branched chain alpha-ketoacid dehydrogena	0.150	0.202	1 consistent probesets
AL050345	Novel human gene mapping to chomosome 22 /cds=(115,4	0.150	0.202	1 consistent probesets
X83378	H.sapiens mRNA for putative chloride channel /cds=(26,	0.150	0.202	1 consistent probesets
AL022325	Homo sapiens DNA sequence from Fosmid 27C3 on chrom	0.150	0.203	2 consistent probesets
M23234	Human membrane glycoprotein P (mdr3) mRNA, complete	0.150	0.214	2 consistent probesets
AA195301	zr36g11.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-665	0.150	0.214	2 consistent probesets
AA877795	nr10g08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-116	0.150	0.215	4 consistent probesets
Y13835	Homo sapiens mRNA for farnesylated-proteins converting	0.150	0.221	3 consistent probesets
U63127	Human SEC7 homolog Tic (TIC) mRNA, complete cds /cds	0.150	0.222	3 consistent probesets
U48959	Homo sapiens myosin light chain kinase (MLCK) mRNA, co	0.150	0.227	3 consistent probesets
W27601	35a3 Homo sapiens cDNA /gb=W27601 /gi=1307549 /	0.150	0.231	4 consistent probesets
U76421	Human dsRNA adenosine deaminase DRADA2b (DRADA2	0.150	0.241	3 consistent probesets
AI768188	wg82b12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.150	0.243	3 consistent probesets
AL045811	DKFZp434H166_r1 Homo sapiens cDNA, 5 end /clone=DK	0.150	0.249	2 consistent probesets
AL080116	Homo sapiens mRNA; cDNA DKFZp564A0723 (from clone	0.150	0.252	1 consistent probesets
AB014551	Homo sapiens mRNA for KIAA0651 protein, complete cds /	0.150	0.252	1 consistent probesets
AL050110	Homo sapiens mRNA; cDNA DKFZp586J0619 (from clone	0.150	0.252	1 consistent probesets
D87465	Human mRNA for KIAA0275 gene, complete cds /cds=(316	0.150	0.252	1 consistent probesets
AJ223349	Homo sapiens mRNA for HIRIP3 protein, clone pH4-31 /cd	0.150	0.252	1 consistent probesets
AF077953	Homo sapiens protein inhibitor of activated STAT prote	0.150	0.252	1 consistent probesets
U57646	Homo sapiens cysteine and glycine-rich protein 2 (CSRP	0.150	0.266	3 consistent probesets
AF017060	untitled /cds=(298,4314) /gb=AF017060 /gi=2343154 /	0.150	0.273	2 consistent probesets
X57303	H.sapiens REC1L mRNA /cds=(150,2039) /gb=X57303 /gi=	0.150	0.280	3 consistent probesets
AB006624	Homo sapiens mRNA for KIAA0286 gene, partial cds /cds=	0.150	0.282	3 consistent probesets
M62402	HUMIGFBP4 Human insulin-like growth factor bi	0.150	0.282	2 consistent probesets
U34044	Human selenium donor protein (selD) mRNA, complete cds	0.150	0.298	3 consistent probesets
M96684	H.sapiens Pur (pur-alpha) mRNA, complete cds /cds=(59,	0.150	0.301	2 consistent probesets
U97502	Homo sapiens butyrophilin (BT3.3) gene /cds=(73,2268)	0.150	0.302	1 consistent probesets
AI701156	we10f09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-234	0.150	0.302	1 consistent probesets
AB020643	Homo sapiens mRNA for KIAA0836 protein, partial cds /c	0.150	0.302	1 consistent probesets
U03494	Human transcription factor LSF mRNA, complete cds /cds	0.150	0.302	1 consistent probesets
AA311181	EST181940 Homo sapiens cDNA, 5 end /clone=ATCC-157	0.150	0.302	1 consistent probesets
AI972631	wr41c07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-249	0.150	0.302	1 consistent probesets
AF035594	Homo sapiens protein kinase C-alpha mRNA, partial 3 UT	0.150	0.314	3 consistent probesets
AF000986	Homo sapiens Drosophila fat facets related Y protein (	0.150	0.324	3 consistent probesets
U04735	Human microsomal stress 70 protein ATPase core (stch)	0.150	0.350	2 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
H18080	ym38h10.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-50	0.150	0.353	1 consistent probesets
AW006742	wr28g10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.150	0.353	1 consistent probesets
AL049332	Homo sapiens mRNA; cDNA DKFZp564L176 (from clone D	0.150	0.353	1 consistent probesets
U70728	Human cytohesin-2 mRNA, complete cds /cds=(158,1360) /	0.150	0.375	2 consistent probesets
U17743	HSU17743 Human JNK activating kinase (JNKK1)	0.150	0.375	2 consistent probesets
AF045583	Homo sapiens tubby like protein 3 (TULP3) mRNA, comple	0.150	0.384	2 consistent probesets
AI743654	wg41d01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.150	0.385	3 consistent probesets
AB020634	Homo sapiens mRNA for KIAA0827 protein, complete cds /	0.150	0.403	1 consistent probesets
L16782	HUMMPP1X Human putative M phase phosphoprotei	0.150	0.403	1 consistent probesets
AF052514	Homo sapiens thymus specific serine peptidase mRNA, co	0.150	0.403	1 consistent probesets
Y00272	Human cell cycle control gene CDC2 /cds=(126,1019) /gb	0.150	0.434	2 consistent probesets
AB012162	Homo sapiens mRNA for APCL protein, complete cds /cds=	0.150	0.439	1 consistent probesets
AB020650	Homo sapiens mRNA for KIAA0843 protein, complete cds /	0.150	0.454	1 consistent probesets
W30677	zb75h10.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-309	0.150	0.454	1 consistent probesets
AI829701	wf09d11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.150	0.454	1 consistent probesets
Z15005	H.sapiens CENP-E mRNA /cds=(90,8081) /gb=Z15005 /gi=	0.150	0.454	1 consistent probesets
AF038170	Homo sapiens clone 23817 mRNA sequence /cds=UNKNO	0.150	0.496	2 consistent probesets
K02566	Human alpha-2-thiol proteinase inhibitor mRNA, complet	0.150	0.504	1 consistent probesets
U29607	Human methionine aminopeptidase mRNA, complete cds /c	0.150	0.510	2 consistent probesets
AB023159	Homo sapiens mRNA for KIAA0942 protein, partial cds /c	0.150	0.535	2 consistent probesets
U55766	Human Rev interacting protein Rip-1 mRNA, complete cds	0.150	0.553	2 consistent probesets
AF084535	Homo sapiens laforin (EPM2A) mRNA, partial cds /cds=(0	0.150	0.554	1 consistent probesets
M25809	Human endomembrane proton pump subunit mRNA, comp	0.150	0.554	1 consistent probesets
AF047432	Homo sapiens ADP-ribosylation factor mRNA, complete cd	0.150	0.555	2 consistent probesets
AF091433	Homo sapiens cyclin E2 mRNA, complete cds /cds=(37,125	0.150	0.605	1 consistent probesets
U66035	Human X-linked deafness dystonia protein (DDP) mRNA, c	0.150	0.605	1 consistent probesets
AF039397	untitled /cds=(30,974) /gb=AF039397 /gi=2746761 /	0.150	0.655	1 consistent probesets
AL035305	H.sapiens gene from PAC 102G20 /cds=(117,803) /gb=AL0	0.150	0.655	1 consistent probesets
W25874	14e9 Homo sapiens cDNA /gb=W25874 /gi=1306015 /	0.150	0.655	1 consistent probesets
AL080173	Homo sapiens mRNA; cDNA DKFZp434H071 (from clone D	0.150	0.655	1 consistent probesets
AI668626	yo63b04.x5 Homo sapiens cDNA, 3 end /clone=IMAGE-18	0.150	0.756	1 consistent probesets
AF035121	AF035121 Homo sapiens KDR/flk-1 protein mRNA,	0.150	0.806	1 consistent probesets
AF035296	Homo sapiens clone 23837 mRNA sequence /cds=UNKNO	0.150	0.958	1 consistent probesets
AB018340	Homo sapiens mRNA for KIAA0797 protein, partial cds /c	0.150	1.260	1 consistent probesets
L22075	HUMG13A Human guanine nucleotide regulatory p	0.148	0.224	7 inconsistent probesets
L21936	Human succinate dehydrogenase flavoprotein subunit (SD	0.146	0.056	4 consistent probesets
U41654	Human adenovirus protein E3-14.7k interacting protein	0.146	0.062	4 consistent probesets
U37689	HSU37689 Human RNA polymerase II subunit (hsR	0.146	0.070	8 consistent probesets
M11567	HUMAGG Human angiogenin gene, complete cds, a	0.146	0.077	4 consistent probesets
X53002	Human mRNA for integrin beta-5 subunit /cds=(336,2735)	0.146	0.078	4 consistent probesets
AF084481	Homo sapiens transmembrane protein (WFS1) mRNA, com	0.146	0.078	4 inconsistent probesets
AF053641	Homo sapiens brain cellular apoptosis susceptibility p	0.146	0.086	4 consistent probesets
D45333	HUMHG7879 Homo sapiens cDNA /gb=D45333 /gi=113673	0.146	0.088	4 consistent probesets
L38935	Homo sapiens GT212 mRNA /cds=UNKNOWN /gb=L38935	0.146	0.089	4 consistent probesets
U61232	Human tubulin-folding cofactor E mRNA, complete cds /c	0.146	0.091	4 consistent probesets
AF034176	AF034176 Homo sapiens cDNA /clone=ntcon5-contig /gb=A	0.146	0.091	4 consistent probesets
X00734	HSREP10 Human beta-tubulin gene (5-beta) with	0.146	0.092	4 consistent probesets
AI203737	qf76b12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-175	0.146	0.096	4 inconsistent probesets
D87437	Human mRNA for KIAA0250 gene, complete cds /cds=(424	0.146	0.098	4 inconsistent probesets
AF009225	Homo sapiens IkB kinase alpha subunit (IKK alpha) mRNA	0.146	0.098	4 consistent probesets
L25876	HUMPTPB Homo sapiens protein tyrosine phospha	0.146	0.107	4 consistent probesets
M64349	HUMCYCD1 Human cyclin D (cyclin D1) mRNA, com	0.146	0.108	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U34252	Human gamma-aminobutyraldehyde dehydrogenase mRNA	0.146	0.108	4 consistent probesets
D67025	Homo sapiens mRNA for proteasome subunit p58, complet	0.146	0.109	4 consistent probesets
AF062529	Homo sapiens clone 486790 diphosphoinositol polyphosph	0.146	0.111	4 consistent probesets
AB018330	Homo sapiens mRNA for KIAA0787 protein, partial cds /c	0.146	0.113	4 consistent probesets
M74002	Human arginine-rich nuclear protein mRNA, complete cds	0.146	0.113	4 consistent probesets
X02160	Human mRNA for insulin receptor precursor /cds=(48,416	0.146	0.113	4 consistent probesets
X13916	Human mRNA for LDL-receptor related protein /cds=(466,	0.146	0.113	4 consistent probesets
X76061	H.sapiens p130 mRNA for 130K protein /cds=(69,3488) /g	0.146	0.115	4 consistent probesets
AF029893	Homo sapiens i-beta-1,3-N-acetylglucosaminyltransferas	0.146	0.122	4 consistent probesets
W27594	34h4 Homo sapiens cDNA /gb=W27594 /gi=1307542 /	0.146	0.128	4 consistent probesets
AJ238094	Homo sapiens mRNA for Lsm1 protein /cds=(188,589) /gb=	0.146	0.130	4 consistent probesets
D29641	Human mRNA for KIAA0052 gene, partial cds /cds=(0,2510	0.146	0.131	4 consistent probesets
X03674	Human mRNA for glucose-6-phosphate dehydrogenase (G6	0.146	0.133	4 consistent probesets
U91985	Human DNA fragmentation factor-45 mRNA, complete cds	0.146	0.135	4 inconsistent probesets
AB012124	Homo sapiens TCFL5 mRNA for transcription factor-like	0.146	0.139	4 consistent probesets
U67319	Human Lice2 beta cysteine protease mRNA, complete cds	0.146	0.141	4 consistent probesets
AF030234	Homo sapiens splicing factor Sip1 mRNA, complete cds /	0.146	0.141	4 consistent probesets
AB007920	Homo sapiens mRNA for KIAA0451 protein, complete cds /	0.146	0.145	4 consistent probesets
D87116	D87116 Human mRNA for MAP kinase kinase 3b ,c	0.146	0.146	4 consistent probesets
AB014578	Homo sapiens mRNA for KIAA0678 protein, partial cds /c	0.146	0.150	4 consistent probesets
AB002348	Human mRNA for KIAA0350 gene, partial cds /cds=(0,2754	0.146	0.152	4 consistent probesets
AI660963	wf20e04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.146	0.160	4 consistent probesets
D26361	Human mRNA for KIAA0042 gene, complete cds /cds=(439	0.146	0.160	4 consistent probesets
AF012024	AF012024 Homo sapiens integrin cytoplasmic do	0.146	0.163	4 consistent probesets
M74524	HUMHHR6A Human HHR6A (yeast RAD 6 homologue)	0.146	0.164	4 consistent probesets
AF020761	Homo sapiens stimulator of Fe transport mRNA, complete	0.146	0.166	4 consistent probesets
AF052151	Homo sapiens clone 24574 mRNA sequence /cds=UNKNO	0.146	0.169	4 consistent probesets
AF049884	Homo sapiens Arg/Abl-interacting protein ArgBP2a (ArgB	0.146	0.176	4 consistent probesets
D67029	Human SEC14L mRNA, complete cds /cds=(303,2450) /gb=	0.146	0.178	4 consistent probesets
AF053305	Homo sapiens mitotic checkpoint kinase Bub1 (BUB1) mRN	0.146	0.186	4 consistent probesets
M24283	Human major group rhinovirus receptor (HRV) mRNA, com	0.144	0.142	3 inconsistent probesets
R48209	yj63d12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-153	0.144	0.188	3 consistent probesets
X97548	H.sapiens mRNA for TIF1beta zinc finger protein /cds=(	0.142	0.064	4 inconsistent probesets
W26056	18e1 Homo sapiens cDNA /gb=W26056 /gi=1306323 /	0.142	0.067	4 inconsistent probesets
U50939	Human amyloid precursor protein-binding protein 1 mRNA	0.142	0.076	4 consistent probesets
AJ000534	Homo sapiens mRNA for epsilon-sarcoglycan /cds=(68,138	0.142	0.078	4 consistent probesets
D49490	Homo sapiens mRNA for protein disulfide isomerase-rela	0.142	0.078	4 consistent probesets
X70944	H.sapiens mRNA for PTB-associated splicing factor /cds	0.142	0.082	4 consistent probesets
U72514	Human C2f mRNA, complete cds /cds=(0,720) /gb=U72514	0.142	0.085	4 consistent probesets
L35249	Homo sapiens vacuolar H+-ATPase Mr 56,000 subunit (HO	0.142	0.087	4 consistent probesets
X14850	Human H2A.X mRNA encoding histone H2A.X /cds=(73,504	0.142	0.088	4 consistent probesets
M62403	HUMIGFBP5 Human insulin-like growth factor bi	0.142	0.088	4 consistent probesets
AF054996	Homo sapiens clone 23783 mRNA sequence /cds=UNKNO	0.142	0.088	4 consistent probesets
AI864120	wg64a06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.142	0.089	4 consistent probesets
X87237	H.sapiens mRNA for processing a-glucosidase I /cds=(13	0.142	0.089	4 consistent probesets
X69141	H.sapiens mRNA for squalene synthase /cds=(91,1344) /g	0.142	0.091	4 consistent probesets
AB023173	Homo sapiens mRNA for KIAA0956 protein, partial cds /c	0.142	0.091	4 consistent probesets
D84109	Homo sapiens mRNA for RBP-MS/type 3, complete cds /cd	0.142	0.104	4 consistent probesets
AF074723	Homo sapiens RNA polymerase transcriptional regulation	0.142	0.108	4 consistent probesets
M80254	H.sapiens cyclophilin isoform (hCyP3) mRNA, complete c	0.142	0.112	4 consistent probesets
AF016369	Homo sapiens U4/U6 small nuclear ribonucleoprotein hPr	0.142	0.113	4 inconsistent probesets
AF083190	Homo sapiens SPF31 (SPF31) mRNA, complete cds /cds=(	0.142	0.114	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
L22475	HUMBAXG Human Bax gamma mRNA, complete cds"	0.142	0.117	4 inconsistent probesets
U14193	HSU14193 Human TFIIA gamma subunit mRNA, comp	0.142	0.125	4 consistent probesets
D50919	Human mRNA for KIAA0129 gene, complete cds /cds=(10,1	0.142	0.125	4 consistent probesets
U68063	HSU68063 Human transformer-2 beta (htra-2 bet	0.142	0.127	4 consistent probesets
D78335	Human mRNA for 5-terminal region of UMK, complete cds	0.142	0.134	4 consistent probesets
U53225	Human sorting nexin 1 (SNX1) mRNA, complete cds /cds=(	0.142	0.139	4 consistent probesets
AB007915	Homo sapiens mRNA for KIAA0446 protein, complete cds /	0.142	0.140	4 consistent probesets
D86963	Human mRNA for KIAA0208 gene, complete cds /cds=(140	0.142	0.143	4 inconsistent probesets
M88468	Homo sapiens mevalonate kinase mRNA, complete cds /cd	0.142	0.146	4 consistent probesets
U07804	HSU07804 Human DNA topoisomerase I mRNA, part	0.142	0.148	4 consistent probesets
AB007944	Homo sapiens mRNA for KIAA0475 protein, complete cds /	0.142	0.149	4 consistent probesets
AI095508	qb29a06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.142	0.168	4 consistent probesets
AF042386	Homo sapiens cyclophilin-33B (CYP-33) mRNA, complete c	0.142	0.171	4 consistent probesets
U77942	Human syntaxin 7 mRNA, complete cds /cds=(79,864) /gb=	0.142	0.173	4 consistent probesets
W26226	22e3 Homo sapiens cDNA /gb=W26226 /gi=1306637 /	0.142	0.178	4 consistent probesets
U49184	Human occludin mRNA, complete cds /cds=(167,1735) /gb=	0.142	0.194	4 consistent probesets
Y08136	H.sapiens mRNA for ASM-like phosphodiesterase 3a /cds=	0.142	0.202	4 consistent probesets
AB023154	Homo sapiens mRNA for KIAA0937 protein, partial cds /c	0.142	0.206	4 consistent probesets
AF001307	Homo sapiens aryl hydrocarbon receptor nuclear translo	0.142	0.225	4 consistent probesets
U05227	HSU05227 Human Rar protein mRNA, complete cds	0.142	0.231	4 consistent probesets
AF038202	Homo sapiens clone 23570 mRNA sequence /cds=UNKNO	0.142	0.234	4 consistent probesets
AF007130	Homo sapiens clone 23750 unknown mRNA, partial cds /cd	0.142	0.237	4 consistent probesets
AI925946	wh12f04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.142	0.239	4 consistent probesets
D86970	Human mRNA for KIAA0216 gene, complete cds /cds=(484	0.142	0.245	4 consistent probesets
AB018308	Homo sapiens mRNA for KIAA0765 protein, partial cds /c	0.142	0.255	4 consistent probesets
D26155	Human mRNA for transcriptional activator hSNF2a, compl	0.142	0.281	4 consistent probesets
AF022853	untitled /cds=(0,4202) /gb=AF022853 /gi=2585771 /	0.140	0.136	7 inconsistent probesets
X02344	Homo sapiens beta 2 gene /cds=(0,1337) /gb=X02344 /gi=	0.140	0.033	8 inconsistent probesets
AF004828	Homo sapiens rab3-GAP regulatory domain mRNA, comple	0.139	0.346	3 consistent probesets
U46692	Human cystatin B gene, complete cds /cds=(96,392) /gb=	0.138	0.058	4 inconsistent probesets
AB024704	Homo sapiens mRNA for fls353, complete cds /cds=(471,2	0.138	0.058	4 consistent probesets
AB002368	Human mRNA for KIAA0370 gene, partial cds /cds=(0,2406	0.138	0.067	4 consistent probesets
J04088	HUMTOPII Human DNA topoisomerase II (top2) mR	0.138	0.070	4 inconsistent probesets
J05614	HUMPCNAPRM Human proliferating cell nuclear a	0.138	0.073	4 inconsistent probesets
AL050128	Homo sapiens mRNA; cDNA DKFZp586G051 (from clone D	0.138	0.075	4 consistent probesets
X79448	H.sapiens IFI-4 mRNA for type I protein /cds=(1165,396	0.138	0.077	4 consistent probesets
D64154	Human mRNA for Mr 110,000 antigen, complete cds /cds=(	0.138	0.080	4 inconsistent probesets
U47634	HSU47634 Human beta-tubulin class III isotype	0.138	0.085	4 inconsistent probesets
AJ131244	Homo sapiens mRNA for Sec24 protein (Sec24A isoform),	0.138	0.085	4 inconsistent probesets
AF077820	Homo sapiens LDL receptor member LR3 mRNA, complete	0.138	0.087	4 consistent probesets
U66306	Human retinoid X receptor alpha mRNA, 3 UTR, partial s	0.138	0.106	4 consistent probesets
AL050107	Homo sapiens mRNA; cDNA DKFZp586I1419 (from clone D	0.138	0.111	4 consistent probesets
D87969	Homo sapiens mRNA for CMP-sialic acid transporter, com	0.138	0.112	4 consistent probesets
AI032612	ow17e07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.138	0.117	4 consistent probesets
U33635	Human colon carcinoma kinase-4 (CCK4) mRNA, complete	0.138	0.119	4 inconsistent probesets
X80230	HSSTPKC2K H.sapiens mRNA (clone C-2k) mRNA fo	0.138	0.127	4 consistent probesets
AB020687	Homo sapiens mRNA for KIAA0880 protein, complete cds /	0.138	0.138	4 consistent probesets
D16611	Human mRNA for coproporphyrinogen oxidase, complete c	0.138	0.141	4 consistent probesets
AB011091	Homo sapiens mRNA for KIAA0519 protein, complete cds /	0.138	0.144	4 consistent probesets
D64110	Homo sapiens mRNA for ANA, complete cds /cds=(94,852)	0.138	0.152	4 consistent probesets
U57627	Human fetal brain oculocerebrorenal syndrome (OCRL1) m	0.138	0.152	4 consistent probesets
X94232	H.sapiens mRNA for novel T-cell activation protein /cd	0.138	0.190	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AI760932	wi70d01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.138	0.190	4 consistent probesets
M16967	Human coagulation factor V mRNA, complete cds /cds=(90	0.138	0.199	4 consistent probesets
U94362	Homo sapiens glycogenin-2 alpha (glycogenin-2) mRNA, c	0.138	0.227	4 consistent probesets
C18655	C18655 Homo sapiens cDNA, 5 end /clone=GEN-565G08 /	0.138	0.247	4 consistent probesets
D87438	Human mRNA for KIAA0251 gene, partial cds /cds=(0,2463	0.137	0.066	4 consistent probesets
AB011159	Homo sapiens mRNA for KIAA0587 protein, complete cds /	0.137	0.072	4 consistent probesets
L34587	HUMRPIE Homo sapiens RNA polymerase II elonga	0.137	0.078	4 inconsistent probesets
AJ132583	Homo sapiens mRNA for puromycin sensitive aminopeptida	0.137	0.078	4 consistent probesets
AF044195	Homo sapiens IkappaB kinase complex associated protein	0.137	0.093	4 consistent probesets
AF029750	Homo sapiens tapasin (NGS-17) mRNA, complete cds /cds=	0.137	0.094	4 consistent probesets
AJ132637	Homo sapiens mRNA for ATP-dependent metalloprotease Y	0.137	0.101	4 consistent probesets
X66899	HSEWS H.sapiens EWS mRNA	0.137	0.101	4 consistent probesets
AF037339	Homo sapiens cleft lip and palate transmembrane protei	0.137	0.110	4 inconsistent probesets
X53586	Human mRNA for integrin alpha 6 /cds=UNKNOWN /gb=X5	0.137	0.111	4 consistent probesets
AA922934	oh10g06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-14	0.137	0.112	4 consistent probesets
AB011156	Homo sapiens mRNA for KIAA0584 protein, partial cds /c	0.137	0.125	4 consistent probesets
U36787	Human putative holocytochrome c-type synthetase mRNA,	0.137	0.147	4 consistent probesets
L34600	Human nuclear-encoded mitochondrial initiation factor	0.137	0.158	4 consistent probesets
AB020716	Homo sapiens mRNA for KIAA0909 protein, partial cds /c	0.137	0.164	4 consistent probesets
AB011087	Homo sapiens mRNA for KIAA0515 protein, partial cds /c	0.137	0.167	4 consistent probesets
D63506	Homo sapiens mRNA for unc-18homologue, complete cds /	0.137	0.168	4 consistent probesets
L07541	Human replication factor C, 38-kDa subunit mRNA, compl	0.137	0.173	4 consistent probesets
D83781	Human mRNA for KIAA0197 gene, partial cds /cds=(0,3945	0.137	0.173	4 consistent probesets
AI739308	wi30c12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.137	0.182	4 consistent probesets
D87957	D87957 Homo sapiens gene for protein involved	0.137	0.205	4 consistent probesets
U19775	HSU19775 Human MAP kinase Mxi2 (MXI2) mRNA, c	0.137	0.214	4 consistent probesets
X02812	Human mRNA for transforming growth factor-beta (TGF-be	0.137	0.248	4 consistent probesets
D25547	Homo sapiens mRNA for PIMT isozyme I, complete cds /cd	0.135	0.052	8 inconsistent probesets
U45328	HSU45328 Human ubiquitin-conjugating enzyme (	0.133	0.060	4 inconsistent probesets
AB015631	Homo sapiens mRNA for type II membrane protein, comple	0.133	0.061	4 inconsistent probesets
M73554	HUMBCL1 Human bcl-1 mRNA, complete CDS"	0.133	0.067	4 consistent probesets
M35011	HUMIBSUB Human integrin beta-5 subunit mRNA,	0.133	0.067	4 consistent probesets
AL023553	dJ347H13.3 (phosphomannomutase 1 (PMMH-22, yeast SE	0.133	0.075	4 consistent probesets
X02596	HSBCRR Human mRNA for bcr (breakpoint cluster	0.133	0.077	5 consistent probesets
D14878	Human mRNA for protein D123, complete cds /cds=(280,12	0.133	0.078	4 consistent probesets
L19711	Human dystroglycan (DAG1) mRNA, complete cds /cds=(39	0.133	0.080	4 consistent probesets
AL080234	Homo sapiens mRNA; cDNA DKFZp586L081 (from clone D	0.133	0.083	4 consistent probesets
U32680	HSU32680 Human CLN3 mRNA, complete cds"	0.133	0.084	4 consistent probesets
U17886	Human succinate dehydrogenase iron-protein subunit (sd	0.133	0.086	4 consistent probesets
X81625	H.sapiens mRNA for Cl1 protein /cds=(135,1448) /gb=X81	0.133	0.088	4 consistent probesets
AI961929	wt39g02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.133	0.089	4 consistent probesets
X77548	H. sapiens cDNA for RFG /cds=(76,1920) /gb=X77548 /gi=	0.133	0.090	4 consistent probesets
M77142	Human polyadenylate binding protein (TIA-1) mRNA, comp	0.133	0.090	4 inconsistent probesets
L33801	HUMGLSYKIN Human protein kinase mRNA, complet	0.133	0.091	8 consistent probesets
M86752	HUMIEF Human transformation-sensitive protein	0.133	0.092	4 consistent probesets
X15804	Human mRNA for alpha-actinin /cds=(198,2876) /gb=X158	0.133	0.092	4 consistent probesets
D78577	D78576S2 Human DNA for 14-3-3 protein eta cha	0.133	0.094	4 consistent probesets
U49248	Human canalicular multispecific organic anion transpor	0.133	0.096	4 consistent probesets
M97856	Homo sapiens histone-binding protein mRNA, complete cd	0.133	0.097	4 consistent probesets
AF039555	Homo sapiens visinin-like protein 1 (VSNL1) mRNA, comp	0.133	0.101	4 consistent probesets
AJ011001	Homo sapiens mRNA for TM7XN1 protein /cds=(316,2379)	0.133	0.104	4 consistent probesets
AF038954	Homo sapiens vacuolar H(+)-ATPase subunit mRNA, comp	0.133	0.108	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
X77494	H.sapiens MSSP-2 mRNA /cds=(231,1400) /gb=X77494 /gi	0.133	0.108	4 consistent probesets
AB018284	Homo sapiens mRNA for KIAA0741 protein, complete cds /	0.133	0.112	4 consistent probesets
AB016247	Homo sapiens mRNA for sterol-C5-desaturase, complete c	0.133	0.116	4 inconsistent probesets
X01703	Human gene for alpha-tubulin (b alpha 1) /cds=(213,156	0.133	0.116	4 consistent probesets
M83088	Human phosphoglucomutase 1 (PGM1) mRNA, complete c	0.133	0.116	4 consistent probesets
M13452	Human lamin A mRNA, 3end /cds=(0,1547) /gb=M13452 /g	0.133	0.117	7 consistent probesets
M12267	Human ornithine aminotransferase mRNA, complete cds /c	0.133	0.127	4 consistent probesets
AL050223	Homo sapiens mRNA; cDNA DKFZp586L1323 (from clone	0.133	0.127	4 consistent probesets
W26496	30d2 Homo sapiens cDNA /gb=W26496 /gi=1307195 /	0.133	0.127	4 consistent probesets
AB011174	Homo sapiens mRNA for KIAA0602 protein, partial cds /c	0.133	0.131	3 consistent probesets
U51205	Human COP9 homolog (HCOP9) mRNA, complete cds /cds	0.133	0.133	4 consistent probesets
X83857	H.sapiens mRNA for prostaglandin E receptor (EP3a1) /c	0.133	0.134	1 consistent probesets
AL080070	Homo sapiens mRNA; cDNA DKFZp564M112 (from clone D	0.133	0.142	4 consistent probesets
AF001294	Homo sapiens IPL (IPL) mRNA, complete cds /cds=(56,514	0.133	0.149	4 inconsistent probesets
U40763	Human Clk-associated RS cyclophilin CARS-Cyp mRNA, c	0.133	0.150	4 consistent probesets
D38553	Human mRNA for KIAA0074 gene, partial cds /cds=(0,2244	0.133	0.163	3 consistent probesets
D42047	Human mRNA for KIAA0089 gene, partial cds /cds=(0,1236	0.133	0.170	4 consistent probesets
U64805	Homo sapiens Brca1-delta11b (Brca1) mRNA, complete cds	0.133	0.174	4 consistent probesets
D83703	D83703 Homo sapiens mRNA for peroxisome assem	0.133	0.185	3 consistent probesets
D38255	Homo sapiens mRNA for CAB1, complete cds /cds=(121,14	0.133	0.186	3 consistent probesets
D29954	Human mRNA for KIAA0056 gene, partial cds /cds=(0,4524	0.133	0.186	3 consistent probesets
AA191426	zp83g09.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-62	0.133	0.189	4 consistent probesets
AI004207	ot94g05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-162	0.133	0.192	3 consistent probesets
AF063308	Homo sapiens coiled-coil related protein DEEPEST (DEEP	0.133	0.201	4 consistent probesets
D79985	Human mRNA for KIAA0163 gene, complete cds /cds=(185	0.133	0.209	3 consistent probesets
D88208	Homo sapiens hSGT1 mRNA for hSgt1p, complete cds /cds	0.133	0.234	4 consistent probesets
X64044	H.sapiens mmRNA for large subunit of splicing factor U	0.133	0.234	3 consistent probesets
AB002445	Homo sapiens mRNA from chromosome 5q21-22, clone-FB	0.133	0.248	3 consistent probesets
AF052123	Homo sapiens clone 23770 mRNA sequence /cds=UNKNO	0.133	0.254	4 consistent probesets
AL049669	Human gene from PAC 612B18, chromosome 1 /cds=(272,	0.133	0.265	3 consistent probesets
N23137	yx67h12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-26	0.133	0.270	3 consistent probesets
D79988	Human mRNA for KIAA0166 gene, complete cds /cds=(163	0.133	0.276	3 consistent probesets
M55905	Human mitochondrial NAD(P)+ dependent malic enzyme m	0.133	0.278	3 consistent probesets
X86098	H.sapiens mRNA for BS69 protein /cds=(244,1932) /gb=X8	0.133	0.280	3 consistent probesets
AB007860	Homo sapiens KIAA0400 mRNA, complete cds /cds=(340,3	0.133	0.290	4 consistent probesets
AF035621	Homo sapiens kinesin-related protein (KIF3C) mRNA, com	0.133	0.302	3 consistent probesets
AI827730	wf11d12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.133	0.304	4 consistent probesets
W74442	zd75e09.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-34	0.133	0.311	3 consistent probesets
L35546	Homo sapiens gamma-glutamylcysteine synthetase light s	0.133	0.314	3 consistent probesets
AF098638	Homo sapiens rabaptin-4 mRNA, complete cds /cds=(0,248	0.133	0.368	3 consistent probesets
D42055	Human mRNA for KIAA0093 gene, partial cds /cds=(0,2784	0.133	0.382	3 consistent probesets
AI984786	wr85c06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.133	0.470	1 consistent probesets
D26600	HUMPSH3 Human mRNA for proteasome subunit HsN	0.131	0.055	8 consistent probesets
AA121509	zk88c10.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-48	0.131	0.082	8 inconsistent probesets
AI984234	wz57e04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.131	0.158	8 consistent probesets
L15388	HUMGRK5A Human G protein-coupled receptor kin	0.131	0.096	6 inconsistent probesets
D64015	Homo sapiens mRNA for T-cluster binding protein, compl	0.131	0.145	6 inconsistent probesets
AA402332	zu48h12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-74	0.130	0.288	5 consistent probesets
U52960	Human RNA polymerase II complex component SRB7 mRN	0.130	0.294	5 consistent probesets
X62822	H.sapiens gene encoding beta-galactoside alpha-2,6-sia	0.129	0.070	4 consistent probesets
U66867	Human ubiquitin conjugating enzyme 9 (hUBC9) mRNA, co	0.129	0.073	4 inconsistent probesets
X66397	HSTPRM H.sapiens tpr mRNA	0.129	0.078	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D44466	D44466 Homo sapiens mRNA for proteasome subun	0.129	0.082	4 consistent probesets
D67031	Homo sapiens ADDL mRNA for adducin-like protein, compl	0.129	0.082	4 consistent probesets
AF052134	Homo sapiens clone 23585 mRNA sequence /cds=UNKNO	0.129	0.084	4 consistent probesets
AB028990	Homo sapiens mRNA for KIAA1067 protein, partial cds /c	0.129	0.089	4 consistent probesets
AF047437	Homo sapiens sperm acrosomal protein mRNA, complete c	0.129	0.091	4 consistent probesets
M55153	HUMTGASE Human transglutaminase (TGase) mRNA,	0.129	0.095	4 inconsistent probesets
AF052131	Homo sapiens clone 23930 mRNA sequence /cds=UNKNO	0.129	0.097	4 consistent probesets
AL034428	Human DNA sequence from clone 705D16 on chromosome	0.129	0.100	4 inconsistent probesets
AB007510	Homo sapiens mRNA for PRP8 protein, complete cds /cds=	0.129	0.102	4 consistent probesets
U03985	Human N-ethylmaleimide-sensitive factor mRNA, partial	0.129	0.104	4 consistent probesets
L27706	Human chaperonin protein (Tcp20) gene complete cds /cd	0.129	0.105	4 consistent probesets
L24123	Homo sapiens NRF1 protein (NRF1) mRNA /cds=UNKNOW	0.129	0.105	4 consistent probesets
M69013	HUMGTPBRPA Human guanine nucleotide-binding r	0.129	0.110	8 inconsistent probesets
U70671	Human ataxin-2 related protein mRNA, partial cds /cds=	0.129	0.111	4 consistent probesets
AL050366	Homo sapiens mRNA; cDNA DKFZp564A126 (from clone D	0.129	0.112	4 consistent probesets
AB014530	Homo sapiens mRNA for KIAA0630 protein, partial cds /c	0.129	0.118	4 consistent probesets
Z11584	H.sapiens mRNA for NuMA protein /cds=(258,6563) /gb=Z1	0.129	0.122	4 consistent probesets
D87127	D87127 Homo sapiens mRNA for translocation pr	0.129	0.124	8 consistent probesets
AB018325	Homo sapiens mRNA for KIAA0782 protein, partial cds /c	0.129	0.125	4 consistent probesets
AB020636	Homo sapiens mRNA for KIAA0829 protein, partial cds /c	0.129	0.128	4 consistent probesets
M32334	HUMICAM4 Homo sapiens intercellular adhesion	0.129	0.133	4 consistent probesets
AB002322	Human mRNA for KIAA0324 gene, partial cds /cds=(0,3868	0.129	0.138	4 consistent probesets
AA428150	zw57c05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-77	0.129	0.148	4 consistent probesets
AB020657	Homo sapiens mRNA for KIAA0850 protein, complete cds /	0.129	0.166	4 consistent probesets
AB011108	Homo sapiens mRNA for KIAA0536 protein, partial cds /c	0.129	0.167	4 consistent probesets
AF053977	Homo sapiens cell division cycle protein 23 (CDC23) mR	0.129	0.180	4 consistent probesets
AF038564	Homo sapiens atrophin-1 interacting protein 4 (AIP4) m	0.129	0.187	4 consistent probesets
AA933984	on95f04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-156	0.129	0.193	4 consistent probesets
H82458	yv80b07.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-249	0.129	0.204	4 consistent probesets
D79983	Human mRNA for KIAA0161 gene, complete cds /cds=(348	0.129	0.204	4 consistent probesets
L19297	Human nuclear-encoded mitochondrial carbonic anhydrase	0.129	0.205	4 consistent probesets
W25932	15b1 Homo sapiens cDNA /gb=W25932 /gi=1306055 /	0.129	0.224	4 consistent probesets
U60521	HSU60521 Human protease proMch6 (Mch6) mRNA,	0.129	0.227	4 consistent probesets
U83857	Human Aac11 (aac11) mRNA, complete cds /cds=(77,1663	0.129	0.255	4 consistent probesets
X12534	Human rap2 mRNA for ras-related protein /cds=(3,554) /	0.128	0.171	6 consistent probesets
AF048730	Homo sapiens cyclin T1 mRNA, complete cds /cds=(44,222	0.128	0.233	3 consistent probesets
M34309	HUMHER3A Human epidermal growth factor recept	0.127	0.062	8 consistent probesets
W72239	zd62h08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-345	0.127	0.082	8 consistent probesets
Z48054	H.sapiens mRNA for peroxisomal targeting signal 1 (SKL	0.127	0.148	5 consistent probesets
X82877	H.sapiens Na+-D-glucose cotransport regulator gene /cd	0.126	0.131	7 consistent probesets
AJ245434	Homo sapiens mRNA for G3a protein (G3a gene, located i	0.125	0.039	4 consistent probesets
AI986201	wr81a01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.125	0.068	4 inconsistent probesets
AF034373	Homo sapiens ataxin-2-like protein A2LP (A2LG) mRNA, c	0.125	0.069	4 inconsistent probesets
X91504	H.sapiens mRNA for ARP1 protein /cds=(11,616) /gb=X915	0.125	0.071	2 consistent probesets
X80692	H.sapiens ERK3 mRNA /cds=(478,2643) /gb=X80692 /gi=7	0.125	0.074	4 consistent probesets
AB003102	AB003102 Homo sapiens mRNA for 26S proteasome	0.125	0.076	4 consistent probesets
U96915	Homo sapiens sin3 associated polypeptide p18 (SAP18) m	0.125	0.079	4 inconsistent probesets
AJ005866	Homo sapiens mRNA for putative Sqv-7-like protein, par	0.125	0.083	4 consistent probesets
D89052	Homo sapiens mRNA for proton-ATPase-like protein, comp	0.125	0.084	4 consistent probesets
X92106	HSBLEO H.sapiens mRNA for bleomycin hydrolase	0.125	0.085	8 inconsistent probesets
X85750	H.sapiens mRNA for transcript associated with monocyte	0.125	0.086	4 consistent probesets
X05299	Human mRNA (~95%) for major centromere autoantigen C	0.125	0.088	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D87343	Homo sapiens mRNA for DCRA, complete cds /cds=(239,1	0.125	0.094	4 consistent probesets
U94317	Homo sapiens ribonuclease P protein subunit p40 (RPP40	0.125	0.106	4 consistent probesets
AB006537	Homo sapiens mRNA for interleukin 1 receptor accessory	0.125	0.107	4 consistent probesets
X61100	Human mRNA for mitochondrial 75 kDa iron sulphur prote	0.125	0.108	4 inconsistent probesets
AA487755	ab13f01.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-840	0.125	0.111	4 inconsistent probesets
AB014595	Homo sapiens mRNA for KIAA0695 protein, complete cds /	0.125	0.112	4 consistent probesets
D84145	Human WS-3 mRNA, complete cds /cds=(87,659) /gb=D84	0.125	0.112	4 consistent probesets
U23803	Human heterogeneous ribonucleoprotein A0 mRNA, comple	0.125	0.112	4 consistent probesets
X79535	H.sapiens mRNA for beta tubulin, clone nuk_278 /cds=(6	0.125	0.116	4 consistent probesets
X77723	H.sapiens mRNA for unknown protein of uterine endometr	0.125	0.118	4 consistent probesets
U30872	Human mitosin mRNA, complete cds /cds=(72,9413) /gb=U	0.125	0.119	4 inconsistent probesets
AB007963	Homo sapiens mRNA for KIAA0494 protein, complete cds /	0.125	0.120	4 consistent probesets
U18934	Human receptor tyrosine kinase (DTK) mRNA, complete cd	0.125	0.121	4 consistent probesets
AF009242	Homo sapiens proline-rich Gla protein 1 (PRGP1) mRNA,	0.125	0.122	4 consistent probesets
U29656	HSU29656 Human DR-nm23 mRNA, complete cds"	0.125	0.125	2 consistent probesets
U28014	HSU28014 Human cysteine protease (ICErel-II)	0.125	0.125	4 consistent probesets
X95808	H.sapiens mRNA for protein encoded by a candidate gene	0.125	0.127	4 consistent probesets
U84007	Human glycogen debranching enzyme isoform 1 (AGL) mR	0.125	0.128	4 consistent probesets
L16991	HUMCDC8X Human thymidylate kinase (CDC8) mRNA	0.125	0.130	4 consistent probesets
AA725102	ai08h05.s1 Homo sapiens cDNA, 3 end /clone=1342233 /c	0.125	0.132	4 inconsistent probesets
U90426	HSU90426 Human nuclear RNA helicase, complete	0.125	0.133	4 consistent probesets
U38276	Human semaphorin III family homolog mRNA, complete cd	0.125	0.135	4 consistent probesets
AL050281	Homo sapiens mRNA; cDNA DKFZp586G1219 (from clone	0.125	0.136	4 consistent probesets
AL036744	DKFZp564I1663_r1 Homo sapiens cDNA, 5 end /clone=DK	0.125	0.136	4 consistent probesets
M84820	HUMRXRB Human retinoid X receptor beta (RXR-b	0.125	0.149	4 consistent probesets
AB014522	Homo sapiens mRNA for KIAA0622 protein, partial cds /c	0.125	0.151	4 consistent probesets
AL049998	Homo sapiens mRNA; cDNA DKFZp564L222 (from clone D	0.125	0.161	4 inconsistent probesets
W80358	zh49a07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-41	0.125	0.161	2 consistent probesets
X96586	H.sapiens mRNA for FAN protein /cds=(12,2765) /gb=X965	0.125	0.162	4 consistent probesets
D63477	Human mRNA for KIAA0143 gene, partial cds /cds=(0,2658	0.125	0.163	4 consistent probesets
U83993	Human P2X4 purinoreceptor mRNA, complete cds /cds=(30	0.125	0.171	4 consistent probesets
AI557497	Pt2.1_16_A04.r Homo sapiens cDNA, 3 end /clone_end=3	0.125	0.172	4 consistent probesets
U38545	HSU38545 Human ARF-activated phosphatidylchol	0.125	0.173	4 consistent probesets
AA811338	ob81g05.s1 Homo sapiens cDNA /clone=IMAGE-1337816 /	0.125	0.182	2 consistent probesets
U23028	Human eukaryotic initiation factor 2B-epsilon mRNA, pa	0.125	0.186	4 consistent probesets
AF054284	Homo sapiens spliceosomal protein SAP 155 mRNA, comp	0.125	0.191	4 consistent probesets
AB010414	Homo sapiens mRNA for G-protein gamma 7, complete cds	0.125	0.197	2 consistent probesets
U39064	HSU39064 Human MAP kinase kinase 6 mRNA, comp	0.125	0.198	4 consistent probesets
U88153	Homo sapiens p160 mRNA, partial cds /cds=(0,3755) /gb=	0.125	0.199	4 consistent probesets
AF029670	Homo sapiens Rad51C truncated protein (RAD51C) mRNA.	0.125	0.199	4 consistent probesets
AB016194	Homo sapiens elk1 oncogene, complete cds /cds=(309,159	0.125	0.203	2 consistent probesets
AL080071	Homo sapiens mRNA; cDNA DKFZp564M082 (from clone D	0.125	0.203	2 consistent probesets
M84443	H.sapiens galactokinase (GK2) mRNA, complete cds /cds=	0.125	0.205	4 consistent probesets
D28588	Human mRNA for KIAA0048 gene, complete cds /cds=(334	0.125	0.217	2 consistent probesets
L04510	Human nucleotide binding protein mRNA, complete cds /c	0.125	0.232	2 consistent probesets
AB018315	Homo sapiens mRNA for KIAA0772 protein, complete cds /	0.125	0.238	2 consistent probesets
AB030654	Homo sapiens mRNA for AP-4 clathrin adaptor-related co	0.125	0.252	2 consistent probesets
M16038	HUMLYN Human lyn mRNA encoding a tyrosine kin	0.125	0.263	4 consistent probesets
U69645	Human zinc finger protein mRNA, complete cds /cds=(122	0.125	0.276	4 consistent probesets
AA916905	oh85a09.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-14	0.125	0.294	2 consistent probesets
X87871	H.sapiens mRNA for hepatocyte nuclear factor 4b /cds=(	0.125	0.303	2 consistent probesets
U62961	Human succinyl CoA-3-oxoacid CoA transferase precursor	0.125	0.419	2 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
J03798	Human autoantigen small nuclear ribonucleoprotein Sm-D	0.125	0.449	2 consistent probesets
AA923149	on18f09.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-155	0.125	0.665	2 consistent probesets
AF067656	Homo sapiens ZW10 interactor Zwint mRNA, complete cds	0.125	0.058	4 inconsistent probesets
M29870	HUMRACA Human ras-related C3 botulinum toxin	0.125	0.072	4 consistent probesets
L12168	HUMADCY Homo sapiens adenylyl cyclase-associa	0.125	0.079	4 inconsistent probesets
AL049969	Homo sapiens mRNA; cDNA DKFZp564A072 (from clone D	0.125	0.083	4 consistent probesets
U07418	HSHMLHI Human DNA mismatch repair (hmlh1) mRN	0.125	0.091	4 consistent probesets
L47276	HUMTOPATR Homo sapiens (cell line HL-60) alph	0.125	0.098	4 inconsistent probesets
AB010882	Homo sapiens mRNA for hSNF2H, complete cds /cds=(202	0.125	0.102	4 consistent probesets
AF031647	Homo sapiens JAB1-containing signalosome subunit 3 (SG	0.125	0.107	4 consistent probesets
D86957	Human mRNA for KIAA0202 gene, partial cds /cds=(0,1527	0.125	0.107	4 consistent probesets
AL050373	Homo sapiens mRNA; cDNA DKFZp586F1318 (from clone	0.125	0.116	4 inconsistent probesets
AF012072	Homo sapiens eIF4GII mRNA, complete cds /cds=(256,501	0.125	0.163	4 consistent probesets
U71087	HSU71087 Human MAP kinase kinase MEK5b mRNA,	0.125	0.191	4 consistent probesets
D86981	Human mRNA for KIAA0228 gene, partial cds /cds=(0,2045	0.124	0.206	7 consistent probesets
M14333	HUMCSYNA Homo sapiens c-syn protooncogene mRN	0.123	0.089	8 inconsistent probesets
X15998	H.sapiens mRNA for the chondroitin sulphate proteoglyc	0.123	0.102	8 consistent probesets
X95632	HSARGBPIA H.sapiens mRNA for Arg protein tyro	0.123	0.145	8 inconsistent probesets
D10495	HUMPKSCD Homo sapiens mRNA for protein kinase	0.122	0.164	6 consistent probesets
X80062	H.sapiens SA mRNA /cds=(160,1452) /gb=X80062 /gi=6632	0.122	0.198	6 consistent probesets
U19599	HSU19599 Human (BAX delta) mRNA, complete cds	0.121	0.077	7 inconsistent probesets
D00017	HUMLIC Homo sapiens mRNA for lipocortin II, c	0.121	0.041	4 inconsistent probesets
M96982	Homo sapiens U2 snRNP auxiliary factor small subunit,	0.121	0.055	4 inconsistent probesets
AB018288	Homo sapiens mRNA for KIAA0745 protein, partial cds /c	0.121	0.062	4 consistent probesets
U78027	Homo sapiens Brutons tyrosine kinase (BTK), alpha-D-ga	0.121	0.062	4 consistent probesets
D90086	Human pyruvate dehydrogenase (EC 1.2.4.1) beta subunit	0.121	0.064	4 inconsistent probesets
U29344	Human breast carcinoma fatty acid synthase mRNA, compl	0.121	0.067	4 consistent probesets
Z29505	H.sapiens mRNA for nucleic acid binding protein sub2.3	0.121	0.068	4 consistent probesets
U14603	HSU14603 Human protein-tyrosine phosphatase (	0.121	0.070	8 consistent probesets
M63959	Human alpha-2-macroglobulin receptor-associated protei	0.121	0.072	4 consistent probesets
M55210	HUMLB2A26 Human laminin B2 chain gene, exon 2	0.121	0.072	4 consistent probesets
AA160708	zo72c02.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-592	0.121	0.074	4 consistent probesets
Z97074	HSRAB9P40 Homo sapiens mRNA for Rab9 effector	0.121	0.076	4 consistent probesets
AF040958	Homo sapiens lysosomal neuraminidase precursor, mRNA,	0.121	0.079	4 consistent probesets
U48734	Human non-muscle alpha-actinin mRNA, complete cds /cds	0.121	0.080	4 consistent probesets
AB007898	Homo sapiens KIAA0438 mRNA, complete cds /cds=(117,2	0.121	0.080	4 consistent probesets
X06815	Human mRNA for hU1-70K small nuclear RNP protein (RNI	0.121	0.081	4 inconsistent probesets
AI743745	wg53d06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.121	0.082	4 consistent probesets
U05340	Human p55CDC mRNA, complete cds /cds=(110,1609) /gb=	0.121	0.082	4 consistent probesets
AA056747	zk81f02.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-489	0.121	0.084	4 consistent probesets
D86987	Homo sapiens mRNA for KIAA0214 protein, complete cds /	0.121	0.086	4 inconsistent probesets
U53588	Homo sapiens MHC class 1 region /cds=(199,579) /gb=U53	0.121	0.087	4 consistent probesets
AA152202	zl06a03.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-491	0.121	0.088	4 inconsistent probesets
Y15227	Homo sapiens mRNA for leukemia associated gene 1 /cds=	0.121	0.090	4 inconsistent probesets
AL050280	Homo sapiens mRNA; cDNA DKFZp586B0519 (from clone	0.121	0.092	4 consistent probesets
U66689	HSU66689 Homo sapiens cDNA /gb=U66689 /gi=1906574 /	0.121	0.093	4 inconsistent probesets
D83174	Human mRNA for collagen binding protein 2, complete cd	0.121	0.094	8 consistent probesets
AB028950	Homo sapiens mRNA for KIAA1027 protein, partial cds /c	0.121	0.094	4 consistent probesets
AJ007398	Homo sapiens mRNA for PBK1 protein /cds=(5,1558) /gb=A	0.121	0.095	4 consistent probesets
AF035280	Homo sapiens clone 23689 mRNA, complete cds /cds=(46,	0.121	0.095	4 consistent probesets
L22473	HUMBAXA Human Bax alpha mRNA, complete cds"	0.121	0.098	4 inconsistent probesets
U60061	Human FEZ2 mRNA, partial cds /cds=(0,461) /gb=U60061 /	0.121	0.100	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AL080061	Homo sapiens mRNA; cDNA DKFZp564H182 (from clone D	0.121	0.102	4 consistent probesets
U37012	Human cleavage and polyadenylation specificity factor	0.121	0.105	4 consistent probesets
U17986	Human GABA/noradrenaline transporter mRNA, complete c	0.121	0.105	4 inconsistent probesets
AB011116	Homo sapiens mRNA for KIAA0544 protein, partial cds /c	0.121	0.106	4 consistent probesets
AB023181	Homo sapiens mRNA for KIAA0964 protein, complete cds /	0.121	0.107	4 consistent probesets
Y15521	Homo sapiens ASMTL gene /cds=(0,1889) /gb=Y15521 /gi=	0.121	0.108	4 consistent probesets
D80000	Human mRNA for KIAA0178 gene, partial cds /cds=(0,3679	0.121	0.108	4 consistent probesets
AF046001	Homo sapiens zinc finger transcription factor (ZNF207)	0.121	0.108	4 consistent probesets
AW024812	wu69c05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-99	0.121	0.127	4 consistent probesets
U17566	Human 65 kDa hydrophobic protein mRNA, complete cds /c	0.121	0.133	4 consistent probesets
AB020645	Homo sapiens mRNA for KIAA0838 protein, complete cds /	0.121	0.133	4 consistent probesets
AF046059	Homo sapiens cytokine receptor related protein 4 (CYTO	0.121	0.155	4 consistent probesets
AL050297	Homo sapiens mRNA; cDNA DKFZp564N123 (from clone D	0.121	0.156	4 consistent probesets
Y17169	Homo sapiens mRNA for A6 related protein /cds=(104,115	0.121	0.158	4 consistent probesets
AB011536	Homo sapiens mRNA for MEGF2, partial cds /cds=(0,4095)	0.121	0.158	4 consistent probesets
M85085	Human cleavage stimulation factor, complete cds /cds=(	0.121	0.164	4 consistent probesets
X56807	Human DSC2 mRNA for desmocollins type 2a and 2b /cds=	0.121	0.173	4 inconsistent probesets
AJ224979	Homo sapiens mRNA for MTMR1 protein /cds=(0,1990) /gb	0.121	0.174	4 consistent probesets
D42053	Human mRNA for KIAA0091 gene, complete cds /cds=(496	0.121	0.175	4 consistent probesets
D14720	Homo sapiens gene for peripheral myelin protein zero (	0.121	0.183	4 consistent probesets
AF038172	Homo sapiens clone 23923 mRNA sequence /cds=UNKNO	0.121	0.186	4 consistent probesets
L41351	HUMPROS Homo sapiens prostasin mRNA, complete	0.121	0.194	4 consistent probesets
AI749193	at40e04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-237	0.121	0.207	4 consistent probesets
U58970	Human putative outer mitochondrial membrane 34 kDa tra	0.119	0.058	7 consistent probesets
Y08200	HSRABGTRA Homo sapiens mRNA for rab geranylge	0.119	0.077	7 consistent probesets
D78586	HUMMUPCAD Human CAD mRNA for multifunctional	0.119	0.066	8 inconsistent probesets
N24355	yx14b01.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-261	0.117	0.057	4 inconsistent probesets
D38555	Human mRNA for KIAA0079 gene, complete cds /cds=(114	0.117	0.067	4 inconsistent probesets
M34423	Human beta-galactosidase (GLB1) mRNA, complete cds /c	0.117	0.067	4 consistent probesets
X70476	H.sapiens subunit of coatomer complex /cds=(68,2788) /	0.117	0.070	4 consistent probesets
U56833	HSU56833 Human VHL binding protein-1 (VBP-1)	0.117	0.074	4 inconsistent probesets
D21063	Human mRNA for KIAA0030 gene, partial cds /cds=(0,2745	0.117	0.076	4 consistent probesets
U43747	Human frataxin (FRDA) mRNA, complete cds /cds=(525,11	0.117	0.080	4 consistent probesets
AL049955	Homo sapiens mRNA; cDNA DKFZp564J0123 (from clone	0.117	0.082	4 inconsistent probesets
AF026031	Homo sapiens putative mitochondrial outer membrane pro	0.117	0.084	4 inconsistent probesets
X95592	H.sapiens mRNA for C1D protein /cds=(117,542) /gb=X955	0.117	0.085	4 consistent probesets
X86018	H.sapiens mRNA for MUF1 protein /cds=(0,1853) /gb=X860	0.117	0.087	3 inconsistent probesets
AL080097	Homo sapiens mRNA; cDNA DKFZp564P0462 (from clone	0.117	0.089	4 consistent probesets
D87684	Human mRNA for KIAA0242 gene, partial cds /cds=(0,1590	0.117	0.090	4 inconsistent probesets
X69838	H.sapiens mRNA for G9a /cds=(47,3052) /gb=X69838 /gi=2	0.117	0.091	4 consistent probesets
U22897	Homo sapiens nuclear domain 10 protein (ndp52) mRNA, c	0.117	0.093	4 consistent probesets
AF047473	Homo sapiens testis mitotic checkpoint BUB3 (BUB3) mRN	0.117	0.093	4 inconsistent probesets
X58141	Human mRNA for erythrocyte adducin alpha subunit /cds=	0.117	0.093	4 consistent probesets
AA165701	zo75g08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-59	0.117	0.095	4 consistent probesets
L25444	Homo sapiens (TAFII70-alpha) mRNA, complete cds /cds=(	0.117	0.098	4 consistent probesets
AF038662	Homo sapiens chromosome 3q13 beta-1,4-galactosyltransf	0.117	0.098	4 consistent probesets
U84011	Human glycogen debranching enzyme isoform 6 (AGL) mR	0.117	0.100	4 inconsistent probesets
AI553878	tn30a05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-216	0.117	0.105	4 consistent probesets
U75370	Human mitochondrial RNA polymerase mRNA, nuclear gen	0.117	0.105	4 consistent probesets
U95822	Human putative transmembrane GTPase mRNA, partial cds	0.117	0.108	4 consistent probesets
X79683	H.sapiens LAMB2 mRNA for beta2 laminin /cds=(165,5561)	0.117	0.109	4 inconsistent probesets
X65784	H.sapiens CAR gene /cds=(0,428) /gb=X65784 /gi=452752	0.117	0.111	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U78082	Human RNA polymerase transcriptional regulation mediat	0.117	0.112	4 consistent probesets
L76191	HUMI1R Homo sapiens interleukin-1 receptor-as	0.117	0.113	4 consistent probesets
AB011179	Homo sapiens mRNA for KIAA0607 protein, partial cds /c	0.117	0.113	4 consistent probesets
AJ133769	Homo sapiens mRNA for nuclear transport receptor /cds=	0.117	0.119	3 consistent probesets
AB002310	Human mRNA for KIAA0312 gene, partial cds /cds=(0,5721	0.117	0.121	4 consistent probesets
AF055022	Homo sapiens clone 24684 mRNA sequence /cds=UNKNO	0.117	0.122	4 consistent probesets
U30826	Human splicing factor SRp40-1 (SRp40) mRNA, complete c	0.117	0.122	4 consistent probesets
X70649	Homo sapiens DDX1 gene, complete CDS /cds=(288,2510)	0.117	0.124	4 consistent probesets
AF035284	Homo sapiens clone 23716 mRNA sequence /cds=UNKNO	0.117	0.124	4 consistent probesets
AA621555	af53a04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-103	0.117	0.125	4 consistent probesets
U59878	Human low-Mr GTP-binding protein (RAB32) mRNA, partial	0.117	0.126	4 consistent probesets
AB014529	Homo sapiens mRNA for KIAA0629 protein, partial cds /c	0.117	0.127	4 consistent probesets
U22431	HSU22431 Human hypoxia-inducible factor 1 alp	0.117	0.131	4 consistent probesets
AB015344	Homo sapiens HRIHFB2157 mRNA, partial cds /cds=(0,122	0.117	0.135	4 consistent probesets
L40393	Homo sapiens (clone S171) mRNA, complete cds /cds=(538	0.117	0.135	4 consistent probesets
AB018347	Homo sapiens mRNA for KIAA0804 protein, partial cds /c	0.117	0.136	3 consistent probesets
AL049943	Homo sapiens mRNA; cDNA DKFZp564F0522 (from clone	0.117	0.137	4 consistent probesets
U78190	Human GTP cyclohydrolase I feedback regulatory protein	0.117	0.138	4 consistent probesets
U88966	Human protein rapamycin associated protein (FRAP2) gen	0.117	0.145	4 consistent probesets
AF102265	Homo sapiens N-acetylglucosamine-phosphate mutase mR	0.117	0.145	4 consistent probesets
AF052432	Homo sapiens katanin p80 subunit mRNA, complete cds /c	0.117	0.156	3 consistent probesets
U81607	Homo sapiens gravin mRNA, complete cds /cds=(191,5536	0.117	0.158	4 consistent probesets
M37712	Human p58/GTA (galactosyltransferase associated protei	0.117	0.167	4 consistent probesets
AB008515	Homo sapiens mRNA for RanBPM, complete cds /cds=(435	0.117	0.168	4 consistent probesets
AF001433	Human requiem (HREQ) mRNA, complete cds /cds=(41,12	0.117	0.173	3 consistent probesets
M60091	Homo sapiens galactose-1-phosphate uridyl transferase	0.117	0.175	3 consistent probesets
X80497	H.sapiens PHKLA mRNA /cds=(126,3833) /gb=X80497 /gi=	0.117	0.182	4 consistent probesets
AB029012	Homo sapiens mRNA for KIAA1089 protein, partial cds /c	0.117	0.185	3 consistent probesets
L78132	HUMPCTA1A Human prostate carcinoma tumor anti	0.117	0.191	3 consistent probesets
AJ011972	Homo sapiens mRNA for histone deacetylase-like protein	0.117	0.203	4 consistent probesets
D87466	Human mRNA for KIAA0276 gene, partial cds /cds=(0,931)	0.117	0.205	4 consistent probesets
M15400	HUMRBS Human retinoblastoma susceptibility mR	0.117	0.223	4 consistent probesets
AF057164	Homo sapiens organic cation transporter OCTN2 (OCTN2)	0.117	0.225	3 consistent probesets
Z29066	HSNEK2R H.sapiens nek2 mRNA for protein kinas	0.117	0.232	4 consistent probesets
AA917945	ol68g03.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-153	0.117	0.237	4 consistent probesets
AF054182	Homo sapiens mitochondrial processing peptidase beta-s	0.117	0.242	4 consistent probesets
D87435	Human mRNA for KIAA0248 gene, partial cds /cds=(0,5077	0.117	0.252	3 consistent probesets
U79256	Human clone 23719 mRNA sequence /cds=UNKNOWN /gb	0.117	0.253	3 consistent probesets
M69238	Human aryl hydrocarbon receptor nuclear translocator (	0.117	0.263	3 consistent probesets
X84194	H.sapiens mRNA for acylphosphatase, erythrocyte (CT) i	0.117	0.265	3 consistent probesets
AL049974	Homo sapiens mRNA; cDNA DKFZp564B222 (from clone D	0.117	0.279	4 consistent probesets
U50527	HSU50527 Human BRCA2 region, mRNA sequence CG	0.117	0.281	3 consistent probesets
AL079697	DKFZp434E1930_r1 Homo sapiens cDNA, 5 end /clone=D	0.117	0.289	4 consistent probesets
AL079283	Homo sapiens mRNA full length insert cDNA clone EUROIN	0.117	0.324	3 consistent probesets
AB011120	Homo sapiens mRNA for KIAA0548 protein, partial cds /c	0.117	0.363	4 consistent probesets
X95677	H.sapiens mRNA for ArgBPIB protein /cds=(134,1033) /gb	0.117	0.365	3 consistent probesets
U75309	HSU75309 Human TBP-associated factor (hTAFII1	0.117	0.486	3 consistent probesets
M96995	HUMEGFGRBA Homo sapiens epidermal growth fact	0.115	0.082	8 inconsistent probesets
U22662	HSU22662 Human nuclear orphan receptor LXR-al	0.114	0.107	6 consistent probesets
X87949	H.sapiens mRNA for BiP protein /cds=(222,2183) /gb=X87	0.113	0.050	4 inconsistent probesets
AF026816	Homo sapiens putative oncogene protein mRNA, partial c	0.113	0.058	4 consistent probesets
AF064603	Homo sapiens GA17 protein mRNA, complete cds /cds=(51	0.113	0.061	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D14812	Human mRNA for KIAA0026 gene, complete cds /cds=(305	0.113	0.061	4 consistent probesets
AI557912	pro3.2-2.E07.r Homo sapiens cDNA, 5 end /clone_end=5	0.113	0.061	4 consistent probesets
AB002323	Human mRNA for KIAA0325 gene, partial cds /cds=(0,6265	0.113	0.064	4 inconsistent probesets
AF015254	Homo sapiens serine/threonine kinase (STK-1) mRNA, com	0.113	0.064	4 consistent probesets
AB018331	Homo sapiens mRNA for KIAA0788 protein, partial cds /c	0.113	0.068	4 consistent probesets
AL096723	Homo sapiens mRNA; cDNA DKFZp564H2023 (from clone	0.113	0.072	4 consistent probesets
X64229	H.sapiens dek mRNA /cds=(33,1160) /gb=X64229 /gi=3050	0.113	0.072	4 consistent probesets
X87212	HSCATHCGE H.sapiens mRNA for cathepsin C	0.113	0.072	4 consistent probesets
Y00503	Human mRNA for keratin 19 /cds=(32,1234) /gb=Y00503 /g	0.113	0.072	4 consistent probesets
L03840	HUMFGFR4X Human fibroblast growth factor rece	0.113	0.073	4 consistent probesets
D13900	Homo sapiens mRNA for mitochondrial short-chain enoyl-	0.113	0.075	4 consistent probesets
U03056	HSU03056 Homo sapiens putative tumor suppress	0.113	0.076	4 consistent probesets
X68060	H.sapiens topIIb mRNA for topoisomerase IIb /cds=(0,48	0.113	0.079	4 consistent probesets
X05409	Human RNA for mitochondrial aldehyde dehydrogenase I A	0.113	0.081	4 consistent probesets
AI521453	th60h07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-212	0.113	0.081	4 consistent probesets
U90878	Homo sapiens carboxyl terminal LIM domain protein (CLI	0.113	0.082	4 consistent probesets
AF024636	Homo sapiens STE20-like kinase 3 (mst-3) mRNA, complet	0.113	0.082	4 consistent probesets
D29958	Human mRNA for KIAA0116 gene, partial cds /cds=(0,873)	0.113	0.085	4 inconsistent probesets
X54304	Human mRNA for myosin regulatory light chain /cds=(114	0.113	0.085	4 consistent probesets
AA760866	nz14h07.s1 Homo sapiens cDNA /clone=IMAGE-1287805 /(	0.113	0.085	4 inconsistent probesets
AI262789	qk35e02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-18	0.113	0.085	4 consistent probesets
AF052111	Homo sapiens clone 23953 mRNA sequence /cds=UNKNO	0.113	0.088	4 consistent probesets
U09877	Human helicase-like protein (HLP) mRNA, complete cds /	0.113	0.090	4 consistent probesets
AF035313	Homo sapiens clone 23851 mRNA sequence /cds=UNKNO	0.113	0.090	4 consistent probesets
AI680675	tx40a08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-227	0.113	0.091	4 consistent probesets
AL050109	Homo sapiens mRNA; cDNA DKFZp586J0119 (from clone	0.113	0.091	4 consistent probesets
M63180	Human threonyl-tRNA synthetase mRNA, complete cds /cds	0.113	0.092	4 consistent probesets
U77948	HSU77948 Human Bruton s tyrosine kinase-assoc	0.113	0.093	4 consistent probesets
Y13936	Homo sapiens mRNA for protein phosphatase 2C gamma /(	0.113	0.094	4 consistent probesets
U09564	HSU09564 Human serine kinase mRNA, complete c	0.113	0.101	4 consistent probesets
AB018257	Homo sapiens mRNA for KIAA0714 protein, partial cds /c	0.113	0.104	4 consistent probesets
U46751	Human phosphotyrosine independent ligand p62 for the L	0.113	0.104	4 consistent probesets
U54778	HSU54778 Human 14-3-3 epsilon mRNA, complete	0.113	0.106	4 consistent probesets
AL049954	Homo sapiens mRNA; cDNA DKFZp564A1523 (from clone	0.113	0.109	4 consistent probesets
M55621	Human N-acetylglucosaminyltransferase I (GlcNAc-TI) mR	0.113	0.109	4 consistent probesets
U31814	Human transcriptional regulator homolog RPD3 mRNA, con	0.113	0.112	4 consistent probesets
D21090	HUMHHR23B Human mRNA for XP-C repair compleme	0.113	0.116	4 consistent probesets
H06628	H06628 yl82g03.r1 Soares infant brain 1NIB Ho	0.113	0.121	4 consistent probesets
AF023158	Homo sapiens tyrosine phosphatase (cdc14B) mRNA, comp	0.113	0.122	4 consistent probesets
AB002359	Human mRNA for KIAA0361 gene, KIAA0361 protein /cds=	0.113	0.123	4 inconsistent probesets
D63484	Human mRNA for KIAA0150 gene, partial cds /cds=(0,2835	0.113	0.129	4 consistent probesets
AF038661	Homo sapiens chromosome 1q21-1q23 beta-1,4-galactosylt	0.113	0.133	4 consistent probesets
D42087	Human mRNA for KIAA0118 gene, partial cds /cds=(0,485)	0.113	0.135	4 inconsistent probesets
U11700	Human copper transporting ATPase mRNA, complete cds /(	0.113	0.135	4 consistent probesets
AF002668	Homo sapiens putative fatty acid desaturase MLD mRNA,	0.113	0.148	4 consistent probesets
AB012911	Homo sapiens mRNA for Frizzled-6, complete cds /cds=(2	0.113	0.151	4 consistent probesets
AF091395	Homo sapiens Trio isoform mRNA, complete cds /cds=(66,	0.113	0.154	4 consistent probesets
D30756	Human mRNA for KIAA0049 gene, complete cds /cds=(140	0.113	0.155	4 consistent probesets
AB028944	Homo sapiens mRNA for KIAA1021 protein, partial cds /c	0.113	0.169	4 consistent probesets
L25270	Human XE169 mRNA, complete cds /cds=(531,5213) /gb=L	0.113	0.172	4 consistent probesets
AF020760	Homo sapiens serine protease (Omi) mRNA, complete cds	0.113	0.192	4 consistent probesets
U73338	Human methionine synthase mRNA, complete cds /cds=(39	0.113	0.192	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U79282	Human clone 23801 mRNA sequence /cds=UNKNOWN /gb	0.113	0.196	4 consistent probesets
AI952267	wx50d11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.113	0.199	4 consistent probesets
X65873	H.sapiens mRNA for kinesin (heavy chain) /cds=(313,320	0.113	0.207	4 consistent probesets
AJ007669	Homo sapiens mRNA for Fanconi anemia group G /cds=(49	0.113	0.234	4 consistent probesets
AB011100	Homo sapiens mRNA for KIAA0528 protein, complete cds /	0.113	0.245	4 consistent probesets
AL031282	dJ283E3.6.1 (PUTATIVE novel protein similar to many (a	0.112	0.095	12 consistent probesets
U66042	Human clone 191B7 placenta expressed mRNA from chrom	0.112	0.136	7 inconsistent probesets
M64571	HUMMAP4 Human microtubule-associated protein	0.111	0.063	12 inconsistent probesets
X57348	H.sapiens mRNA (clone 9112) /cds=(165,911) /gb=X57348	0.110	0.058	8 inconsistent probesets
X73478	H.sapiens hPTPA mRNA /cds=(189,1160) /gb=X73478 /gi=	0.110	0.074	8 consistent probesets
Z25821	H.sapiens gene for mitochondrial dodecenoyl-CoA delta-	0.108	0.045	4 inconsistent probesets
D00762	HUMPSC8 Human mRNA for proteasome subunit HC8	0.108	0.059	4 inconsistent probesets
AL050084	Homo sapiens mRNA; cDNA DKFZp566O1646 (from clone	0.108	0.064	4 consistent probesets
AF072902	Homo sapiens gp130 associated protein GAM mRNA, comp	0.108	0.064	4 inconsistent probesets
M65028	Human hnRNP type A/B protein mRNA, complete cds /cds=	0.108	0.068	4 consistent probesets
AB020671	Homo sapiens mRNA for KIAA0864 protein, partial cds /c	0.108	0.068	4 consistent probesets
D87292	Homo sapiens mRNA for rhodanese, complete cds /cds=(48	0.108	0.072	4 inconsistent probesets
U78310	Homo sapiens pescadillo mRNA, complete cds /cds=(58,18	0.108	0.072	4 consistent probesets
AF000982	Homo sapiens dead box, X isoform (DBX) mRNA, alternati	0.108	0.075	4 consistent probesets
AF070616	Homo sapiens clone 24772 BDP-1 protein mRNA, partial c	0.108	0.076	4 consistent probesets
M22324	Human aminopeptidase N/CD13 mRNA encoding aminoper	0.108	0.078	4 consistent probesets
AF070638	Homo sapiens clone 24448 unknown mRNA, partial cds /cd	0.108	0.078	4 consistent probesets
AJ007509	Homo sapiens mRNA for E1B-55kDa-associated protein /co	0.108	0.078	4 consistent probesets
D84557	Homo sapiens mRNA for HsMcm6, complete cds /cds=(61,	0.108	0.080	4 consistent probesets
L37043	Homo sapiens casein kinase I epsilon mRNA, complete cd	0.108	0.081	4 consistent probesets
U70660	Human copper transport protein HAH1 (HAH1) mRNA, com	0.108	0.081	4 consistent probesets
D17517	HUMSKY Human sky mRNA for Sky, complete cds"	0.108	0.081	4 inconsistent probesets
AF002697	Homo sapiens E1B 19K/Bcl-2-binding protein Nip3 mRNA,	0.108	0.083	4 consistent probesets
AF070546	Homo sapiens clone 24607 mRNA sequence /cds=UNKNO	0.108	0.084	4 consistent probesets
U59309	Human fumarase precursor (FH) mRNA, nuclear gene enco	0.108	0.084	4 consistent probesets
X98743	HSRNAHELC H.sapiens mRNA for RNA helicase (My	0.108	0.085	4 inconsistent probesets
X58079	Human mRNA for S100 alpha protein /cds=(113,397) /gb=X	0.108	0.085	4 consistent probesets
M86707	Homo sapiens myristoyl CoA-protein N-myristoyltransfer	0.108	0.087	4 consistent probesets
U12779	HSU12779 Human MAP kinase activated protein k	0.108	0.088	4 consistent probesets
D28124	Human mRNA for unknown product, complete cds /cds=(61	0.108	0.091	4 consistent probesets
M34641	HUMFGF1A Human fibroblast growth factor (FGF)	0.108	0.091	8 consistent probesets
D21260	Human mRNA for KIAA0034 gene, complete cds /cds=(172	0.108	0.093	4 consistent probesets
Y16521	Homo sapiens mRNA for CDS2 protein /cds=(258,1595) /gb	0.108	0.095	4 consistent probesets
AA158243	zo76c01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-592	0.108	0.098	4 inconsistent probesets
M16424	Human beta-hexosaminidase alpha chain (HEXA) gene /cds	0.108	0.099	4 consistent probesets
AJ005698	Homo sapiens mRNA for poly(A)-specific ribonuclease /c	0.108	0.100	4 consistent probesets
D30037	HUMPITPB Human mRNA for phosphatidylinositol	0.108	0.104	4 inconsistent probesets
AB001740	Homo sapiens mRNA for p27, complete cds /cds=(20,619)	0.108	0.105	4 consistent probesets
AJ243310	Homo sapiens mRNA for C14orf3 protein /cds=(131,1147)	0.108	0.108	4 consistent probesets
M93311	HUMMETIII Human metallothionein-III gene, com	0.108	0.111	4 consistent probesets
X51956	Human ENO2 gene for neuron specific (gamma) enolase /c	0.108	0.113	4 consistent probesets
AI828210	wk81e09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.108	0.114	4 consistent probesets
D14663	Human mRNA for KIAA0107 gene, complete cds /cds=(25,1	0.108	0.117	4 consistent probesets
M67468	Human Fragile X mental retardation 1 FMR-1 gene, 3 en	0.108	0.118	4 consistent probesets
U03851	Human capping protein alpha mRNA, partial cds /cds=(16	0.108	0.122	4 consistent probesets
U07231	Homo sapiens G-rich sequence factor-1 (GRSF-1) mRNA, c	0.108	0.124	4 consistent probesets
U52426	Homo sapiens GOK (STIM1) mRNA, complete cds /cds=(56	0.108	0.126	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D26067	Human mRNA for KIAA0033 gene, partial cds /cds=(0,1008	0.108	0.128	4 consistent probesets
AF089814	Homo sapiens growth suppressor related (DOC-1R) mRNA,	0.108	0.131	4 consistent probesets
U07358	HSU07358 Human protein kinase (zpk) mRNA, com	0.108	0.136	4 consistent probesets
U69559	U69559 Homo sapiens cDNA /clone=26077 /gb=U69559 /gi	0.108	0.148	4 consistent probesets
AL049442	Homo sapiens mRNA; cDNA DKFZp586N1720 (from clone	0.108	0.153	4 consistent probesets
AL079314	Homo sapiens mRNA full length insert cDNA clone EUROIM	0.108	0.154	4 consistent probesets
X54232	Human mRNA for heparan sulfate proteaglycan (glypican)	0.108	0.156	4 consistent probesets
M60527	HUMDCKATPB Human deoxycytidine kinase mRNA, c	0.108	0.156	4 consistent probesets
X69962	H.sapiens FMR-1 mRNA /cds=(219,2117) /gb=X69962 /gi=2	0.108	0.161	4 consistent probesets
Z46606	H.sapiens HLTF gene for helicase-like transcription fa	0.108	0.173	4 consistent probesets
AB018281	Homo sapiens mRNA for KIAA0738 protein, complete cds /	0.108	0.177	4 consistent probesets
M20470	Human lymphocyte clathrin light-chain B mRNA, complete	0.108	0.184	4 consistent probesets
U69611	HSU69611 Human TNF-alpha converting enzyme mR	0.108	0.213	4 inconsistent probesets
D13633	Human mRNA for KIAA0008 gene, complete cds /cds=(121	0.108	0.220	4 consistent probesets
AB002386	Human mRNA for KIAA0388 gene, complete cds /cds=(100	0.108	0.223	4 consistent probesets
U50535	HSU50535 Human BRCA2 region, mRNA sequence CG	0.108	0.238	4 consistent probesets
AB028945	Homo sapiens mRNA for KIAA1022 protein, partial cds /c	0.108	0.246	4 consistent probesets
U22233	Human methylthioadenosine phosphorylase (MTAP) mRNA	0.108	0.295	4 consistent probesets
U64820	Homo sapiens josephin MJD1 mRNA, complete cds /cds=(5	0.108	0.413	2 consistent probesets
L39874	HUMDODDA Homo sapiens deoxycytidylate deamina	0.106	0.042	8 inconsistent probesets
M55265	HUMACKII Human casein kinase II alpha subunit	0.106	0.062	8 consistent probesets
U66617	HSU66617 Human SWI/SNF complex 60 KDa subunit	0.106	0.087	8 inconsistent probesets
AL031781	dJ51J12.1.3 (human ortholog of mouse KH Domain RNA B	0.106	0.121	8 inconsistent probesets
U84971	Homo sapiens fetal unknown mRNA, complete cds /cds=(31	0.106	0.382	3 consistent probesets
U72209	HSU72209 Human YY1-associated factor 2 (YAF2)	0.105	0.204	7 consistent probesets
D13969	Human mRNA for Mel-18 protein, complete cds /cds=(201,	0.104	0.053	4 consistent probesets
X57351	HS18D Human 1-8D gene from interferon-inducib	0.104	0.053	4 consistent probesets
D38551	Human mRNA for KIAA0078 gene, complete cds /cds=(184	0.104	0.055	4 consistent probesets
X71973	H.sapiens GPx-4 mRNA for phospholipid hydroperoxide gl	0.104	0.058	4 inconsistent probesets
X59408	H.sapiens, gene for Membrane cofactor protein /cds=UNK	0.104	0.063	4 inconsistent probesets
L15189	Homo sapiens mitochondrial HSP75 mRNA, complete cds /	0.104	0.064	4 inconsistent probesets
N30151	yx81f01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-268	0.104	0.064	4 consistent probesets
U86782	Human 26S proteasome-associated pad1 homolog (POH1)	0.104	0.064	4 inconsistent probesets
D50063	HUMP40MOV Human mRNA for proteasome subunit p	0.104	0.066	8 consistent probesets
U97188	Homo sapiens putative RNA binding protein KOC (koc) mR	0.104	0.068	4 inconsistent probesets
M94362	Human lamin B2 (LAMB2) mRNA, partial cds /cds=(0,1547)	0.104	0.075	4 consistent probesets
AA156987	zl19b05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-502	0.104	0.080	4 inconsistent probesets
U77604	HSU77604 Homo sapiens microsomal glutathione	0.104	0.080	4 consistent probesets
M94046	Human zinc finger protein (MAZ) mRNA /cds=UNKNOWN /	0.104	0.082	4 consistent probesets
U70322	Human transportin (TRN) mRNA, complete cds /cds=(93,27	0.104	0.082	8 inconsistent probesets
X76105	H.sapiens DAP-1 mRNA /cds=(159,467) /gb=X76105 /gi=43	0.104	0.084	4 consistent probesets
AF007875	Homo sapiens dolichol monophosphate mannose synthase	0.104	0.085	4 consistent probesets
X82260	H.sapiens mRNA for RanGTPase activating protein 1 /cds	0.104	0.085	4 consistent probesets
AF051782	Homo sapiens diaphanous 1 (HDIA1) mRNA, complete cds	0.104	0.088	4 consistent probesets
D43949	Human mRNA for KIAA0082 gene, partial cds /cds=(0,1824	0.104	0.089	4 consistent probesets
AI683748	tw53e07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-22(	0.104	0.089	4 inconsistent probesets
AL049934	Homo sapiens mRNA; cDNA DKFZp564M1416 (from clone	0.104	0.089	4 consistent probesets
AF110377	Homo sapiens PCAF-associated factor 400 (PAF400) mRN	0.104	0.093	4 consistent probesets
Z69030	H.sapiens mRNA for gamma 1 isoform of 61kDa regulatory	0.104	0.095	8 consistent probesets
AF026947	Homo sapiens aflatoxin aldehyde reductase AFAR mRNA, (	0.104	0.095	4 consistent probesets
X79865	H.sapiens Mrp17 mRNA /cds=(137,733) /gb=X79865 /gi=13	0.104	0.100	4 consistent probesets
X14487	Human gene for acidic (type I) cytokeratin 10 /cds=(25	0.104	0.100	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D84064	Homo sapiens mRNA for Hrs, complete cds /cds=(60,2393)	0.104	0.102	4 consistent probesets
X63692	H.sapiens mRNA for DNA (cytosin-5)-methyltransferase /	0.104	0.102	4 consistent probesets
AF070548	Homo sapiens clone 24408 2-oxoglutarate carrier protei	0.104	0.103	4 consistent probesets
U26162	Human myosin regulatory light chain mRNA, complete cds	0.104	0.105	4 consistent probesets
R92331	yq03h03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-195	0.104	0.109	4 inconsistent probesets
Z50115	H.sapiens mRNA for thimet oligopeptidase (metalloprote	0.104	0.118	4 consistent probesets
U63810	Homo sapiens WD40 protein Ciao 1 mRNA, complete cds /	0.104	0.119	4 consistent probesets
AL031058	Human DNA sequence from clone 512B11 on chromosome	0.104	0.120	4 consistent probesets
AF015767	Homo sapiens brain and reproductive organ-expressed pr	0.104	0.123	4 consistent probesets
M96577	HUME2F Homo sapiens (E2F-1) pRB-binding prote	0.104	0.125	4 consistent probesets
AF026291	Homo sapiens chaperonin containing t-complex polypepti	0.104	0.128	4 consistent probesets
Y18004	Homo sapiens mRNA for SCML2 protein /cds=(91,2193) /gb	0.104	0.129	4 consistent probesets
W28360	46f9 Homo sapiens cDNA /gb=W28360 /gi=1308371 /	0.104	0.136	4 consistent probesets
AB011004	Homo sapiens HuUAP1 mRNA for UDP-N-acetylglucosamin	0.104	0.144	4 consistent probesets
AB023179	Homo sapiens mRNA for KIAA0962 protein, partial cds /c	0.104	0.150	4 consistent probesets
L19783	Human GPI-H mRNA, complete cds /cds=(60,626) /gb=L19	0.104	0.154	4 consistent probesets
AB007960	chromosome 1 specific transcript KIAA0491 /cds=UNKNOW	0.104	0.159	4 consistent probesets
AF073362	Homo sapiens endo/exonuclease Mre11 (MRE11A) mRNA,	0.104	0.175	8 consistent probesets
AI525379	PT1.1_06_H01.r Homo sapiens cDNA, 5 end /clone_end=5	0.104	0.175	4 consistent probesets
AI201243	qf70f09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-175	0.104	0.183	4 consistent probesets
Y12226	H.sapiens mRNA for gamma-adaptin /cds=(28,2505) /gb=Y	0.104	0.183	4 consistent probesets
AI828880	wj37b02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-240	0.104	0.199	4 consistent probesets
AF098162	Homo sapiens timeless homolog mRNA, complete cds /cds	0.104	0.204	4 consistent probesets
AF058718	Homo sapiens putative 13 S Golgi transport complex 90k	0.104	0.233	4 consistent probesets
U50534	HSU50534 Human BRCA2 region, mRNA sequence CG	0.104	0.234	4 consistent probesets
AL050006	Homo sapiens mRNA; cDNA DKFZp564A033 (from clone D	0.104	0.259	4 consistent probesets
AJ007590	Homo sapiens mRNA for XRP2 protein /cds=(172,1224) /gb	0.104	0.280	4 consistent probesets
AB020684	Homo sapiens mRNA for KIAA0877 protein, partial cds /c	0.104	0.289	4 consistent probesets
U18321	HSU18321 Human ionizing radiation resistance	0.103	0.125	5 consistent probesets
M23410	HUMPLAKO Human plakoglobin (PLAK) mRNA, compl	0.100	0.050	4 consistent probesets
U15979	Human (dlk) mRNA, complete cds /cds=(173,1321) /gb=U15	0.100	0.057	4 consistent probesets
AL080209	Homo sapiens mRNA; cDNA DKFZp586F2423 (from clone	0.100	0.068	4 consistent probesets
X02750	Human liver mRNA for protein C /cds=(97,1482) /gb=X027	0.100	0.072	4 consistent probesets
L37368	Human (clone E5.1) RNA-binding protein mRNA, complete	0.100	0.075	4 consistent probesets
AL041124	DKFZp434D0316_s1 Homo sapiens cDNA, 3 end /clone=D	0.100	0.080	4 consistent probesets
AF040707	Homo sapiens candidate tumor suppressor gene 21 protei	0.100	0.084	8 consistent probesets
AJ132258	Homo sapiens mRNA for staufen protein, partial /cds=(3	0.100	0.084	4 consistent probesets
D79991	Human mRNA for KIAA0169 gene, partial cds /cds=(0,5238	0.100	0.085	4 consistent probesets
X65550	HSMKI67 H.sapiens mki67a mRNA (long type) for	0.100	0.089	8 consistent probesets
X75346	HSMAPKAP H.sapiens mRNA for MAP kinase activa	0.100	0.089	7 inconsistent probesets
W80399	zh49e04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-415	0.100	0.094	4 consistent probesets
U43572	Human alpha-N-acetylglucosaminidase (NAGLU) gene, com	0.100	0.094	4 consistent probesets
D82351	Human retropseudogene MSSP-1 DNA, complete cds /cds=	0.100	0.095	6 inconsistent probesets
U65785	Human 150 kDa oxygen-regulated protein ORP150 mRNA,	0.100	0.097	4 consistent probesets
M31523	HUMTFAA Human transcription factor (E2A) mRNA	0.100	0.099	4 consistent probesets
X96381	H.sapiens erm gene, exon 2,3,4,5 (and joined CDS) /cds	0.100	0.100	4 consistent probesets
AF074382	Homo sapiens IkB kinase gamma subunit (IKK-gamma) mR	0.100	0.101	1 consistent probesets
AF004230	Homo sapiens monocyte/macrophage Ig-related receptor M	0.100	0.101	4 inconsistent probesets
L32976	HUMMLK3A Human protein kinase (MLK-3) mRNA, c	0.100	0.103	6 inconsistent probesets
X69699	HSPAX8A H.sapiens Pax8 mRNA	0.100	0.110	4 consistent probesets
X80907	HSPHOSINK H.sapiens mRNA for p85 beta subunit	0.100	0.113	2 consistent probesets
AF057140	Homo sapiens cargo selection protein TIP47 (TIP47) mRN	0.100	0.116	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D63487	Human mRNA for KIAA0153 gene, partial cds /cds=(0,1918	0.100	0.119	3 consistent probesets
AB007882	Homo sapiens KIAA0422 mRNA, partial cds /cds=(0,2926)	0.100	0.119	3 consistent probesets
L25879	Homo sapiens p53/HEH epoxide hydrolase (EPHX) mRNA,	0.100	0.122	4 consistent probesets
D32143	Human mRNA for biliverdin-IXbeta reductase I /cds=(109	0.100	0.122	4 consistent probesets
S63912	D10S102=FBRNP [human, fetal brain, mRNA, 3043 nt] /cds	0.100	0.122	4 consistent probesets
AF060502	Homo sapiens peroxisome assembly protein PEX10 mRNA	0.100	0.123	4 consistent probesets
AI658639	tu06g05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-225	0.100	0.125	4 consistent probesets
AJ006973	Homo sapiens mRNA for TOM1 protein /cds=(61,1539) /gb=	0.100	0.130	4 consistent probesets
AF039652	Homo sapiens ribonuclease H type II mRNA, complete cds	0.100	0.135	4 consistent probesets
M64788	HUMRAP1GAP Human GTPase activating protein (r	0.100	0.135	3 consistent probesets
U41815	Human nucleoporin 98 (NUP98) mRNA, complete cds /cds=	0.100	0.136	3 consistent probesets
M60922	Human surface antigen mRNA, complete cds /cds=(126,126	0.100	0.141	3 consistent probesets
AI018098	ov65b11.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.100	0.143	2 consistent probesets
X86779	H.sapiens mRNA for FAST kinase /cds=(21,1670) /gb=X867	0.100	0.147	2 consistent probesets
X56841	H.sapiens HLA-E gene /cds=(0,363) /gb=X56841 /gi=43349	0.100	0.147	2 consistent probesets
AF020202	Homo sapiens Munc13 mRNA, complete cds /cds=(224,499	0.100	0.147	2 consistent probesets
Z48481	HSMMPM1 H.sapiens mRNA for membrane-type matr	0.100	0.150	3 consistent probesets
AF049910	Homo sapiens TACC1 (TACC1) mRNA, complete cds /cds=	0.100	0.151	3 consistent probesets
X08020	Human mRNA for glutathione S-transferase subunit 4 (EC	0.100	0.151	1 consistent probesets
D87449	Human mRNA for KIAA0260 gene, partial cds /cds=(0,1153	0.100	0.151	1 consistent probesets
AF096870	Homo sapiens estrogen-responsive B box protein (EBBP)	0.100	0.151	1 consistent probesets
AL080177	Homo sapiens mRNA; cDNA DKFZp434K151 (from clone D	0.100	0.157	3 consistent probesets
U90546	Human butyrophilin (BTF4) mRNA, complete cds /cds=(188	0.100	0.157	3 consistent probesets
U78521	Homo sapiens immunophilin homolog ARA9 mRNA, compl	0.100	0.161	4 consistent probesets
AJ006266	Homo sapiens mRNA for AND-1 protein /cds=(39,3428) /gb	0.100	0.169	3 consistent probesets
AB007867	Homo sapiens KIAA0407 mRNA, complete cds /cds=(270,6	0.100	0.169	2 consistent probesets
J04501	Human muscle glycogen synthase mRNA, complete cds /co	0.100	0.176	3 consistent probesets
AL049305	Homo sapiens mRNA; cDNA DKFZp564A186 (from clone D	0.100	0.176	4 consistent probesets
U43195	Human Rho-associated, coiled-coil containing protein k	0.100	0.177	4 consistent probesets
AF091084	Homo sapiens clone 638 unknown mRNA, complete sequer	0.100	0.182	2 consistent probesets
X95190	H.sapiens mRNA for Branched chain Acyl-CoA Oxidase /cd	0.100	0.185	4 consistent probesets
AB014602	Homo sapiens mRNA for KIAA0702 protein, complete cds /	0.100	0.186	3 consistent probesets
AF093265	Homo sapiens homer-3 mRNA, complete cds /cds=(90,1160	0.100	0.188	4 consistent probesets
U90547	Human Ro/SSA ribonucleoprotein homolog (RoRet) mRNA,	0.100	0.197	2 consistent probesets
U84570	Human A2 mRNA, complete cds /cds=(239,883) /gb=U8457	0.100	0.202	1 consistent probesets
L36463	HUMA Homo sapiens ras interactor (RIN1) mRNA,	0.100	0.202	1 consistent probesets
AB017430	AB017430 Homo sapiens mRNA for kinesin-like D	0.100	0.202	1 consistent probesets
U56816	HSU56816 Human kinase Myt1 (Myt1) mRNA, compl	0.100	0.204	3 consistent probesets
AL050135	Homo sapiens mRNA; cDNA DKFZp586K091 (from clone D	0.100	0.208	3 consistent probesets
M35252	Human CO-029 /cds=(137,850) /gb=M35252 /gi=180925 /	0.100	0.216	3 consistent probesets
AL049378	Homo sapiens mRNA; cDNA DKFZp586I1518 (from clone D	0.100	0.217	2 consistent probesets
X81882	H.sapiens mRNA for for vasopressin activated calcium m	0.100	0.220	3 consistent probesets
X84740	HSDNALIG3 H.sapiens mRNA for DNA ligase III	0.100	0.222	6 consistent probesets
M58297	Human zinc finger protein 42 (MZF-1) mRNA, complete cd	0.100	0.229	3 consistent probesets
L12579	Human alternatively spliced CUTL1 mRNA, complete cds /	0.100	0.231	3 consistent probesets
U66879	HSU66879 Human Bcl-2 binding component 6 (bbc	0.100	0.234	3 consistent probesets
U85267	Homo sapiens down syndrome candidate region 1 (DSCR1)	0.100	0.237	4 consistent probesets
X89066	H.sapiens mRNA for TRPC1 protein /cds=(137,2416) /gb=X	0.100	0.237	5 consistent probesets
AF072468	Homo sapiens (JH8) mRNA, partial cds /cds=(0,1251) /gb	0.100	0.248	2 consistent probesets
AB007863	Homo sapiens KIAA0403 mRNA, partial cds /cds=(0,1250)	0.100	0.252	1 consistent probesets
M83363	Human plasma membrane calcium-pumping ATPase (PMC	0.100	0.252	1 consistent probesets
AF064084	Homo sapiens prenylcysteine carboxyl methyltransferase	0.100	0.252	1 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AL080218	Homo sapiens mRNA; cDNA DKFZp586N1323 (from clone	0.100	0.252	1 consistent probesets
AB015982	Homo sapiens EPK2 mRNA for serine/threonine kinase, co	0.100	0.252	2 consistent probesets
D90359	Human CCG1 mRNA /cds=(51,5669) /gb=D90359 /gi=5593	0.100	0.274	3 consistent probesets
M73047	Homo sapiens tripeptidyl peptidase II mRNA, complete c	0.100	0.278	3 consistent probesets
Y18643	Homo sapiens mRNA for methyltransferase-like protein 1	0.100	0.291	2 consistent probesets
AF040628	Homo sapiens ectodysplasin-A (EDA) mRNA, complete cds	0.100	0.293	1 consistent probesets
AI991631	wr12h09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.100	0.302	1 consistent probesets
AF016052	Homo sapiens zinc finger protein ZNF191 (ZNF191) gene,	0.100	0.302	1 consistent probesets
X92762	H.sapiens mRNA for tafazzins protein /cds=(288,1166) /	0.100	0.302	1 consistent probesets
U20428	Human SNC19 mRNA sequence /cds=UNKNOWN /gb=U20	0.100	0.302	1 consistent probesets
X97198	H.sapiens mRNA for receptor phosphate PCP-2 /cds=(132,	0.100	0.302	1 consistent probesets
U52155	Human ATP-dependent inwardly rectifying potassium chan	0.100	0.302	1 consistent probesets
M93718	HUMNIOXSYN Human nitric oxide synthase mRNA,	0.100	0.303	2 consistent probesets
U74324	Human guanine nucleotide exchange factor mss4 mRNA, c	0.100	0.308	2 consistent probesets
U44754	Human PSE-binding factor PTF gamma subunit mRNA, cor	0.100	0.319	2 consistent probesets
M61906	HUMP13KIN Human P13-kinase associated p85 mRN	0.100	0.339	4 consistent probesets
U19142	Human GAGE-1 protein mRNA, complete cds /cds=(48,464	0.100	0.343	2 consistent probesets
AB018293	Homo sapiens mRNA for KIAA0750 protein, complete cds /	0.100	0.343	2 consistent probesets
AI738463	wi32b08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.100	0.353	1 consistent probesets
L25665	Human GTP-binding protein (HSR1) mRNA, complete cds /	0.100	0.353	1 consistent probesets
U66468	HSU66468 Human cell growth regulator CGR11 mR	0.100	0.403	1 consistent probesets
D10523	Human mRNA for 2-oxoglutarate dehydrogenase, complete	0.100	0.403	1 consistent probesets
W26641	34b7 Homo sapiens cDNA /gb=W26641 /gi=1307484 /	0.100	0.403	1 consistent probesets
AB011420	Homo sapiens mRNA for DRAK1, complete cds /cds=(117,	0.100	0.403	1 consistent probesets
AF052941	Homo sapiens DAP-kinase related protein 1 mRNA, comple	0.100	0.413	2 consistent probesets
AI961220	wt15b04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.100	0.439	1 consistent probesets
Z18956	H.sapiens mRNA for taurine transporter /cds=(19,1878)	0.100	0.439	1 consistent probesets
AF068227	Homo sapiens putative transmembrane protein (CLN5) mR	0.100	0.454	1 consistent probesets
AI547308	PN001_AH_B03.r Homo sapiens cDNA, 5 end /clone_end=	0.100	0.504	1 consistent probesets
AF086904	Homo sapiens protein kinase Chk2 (CHK2) mRNA, complet	0.100	0.504	1 consistent probesets
U60269	Human endogenous retrovirus HERV-K(HML6) proviral clon	0.100	0.609	2 consistent probesets
D83780	Human mRNA for KIAA0196 gene, complete cds /cds=(273	0.100	0.622	2 consistent probesets
U13045	Human nuclear respiratory factor-2 subunit beta 1 mRNA	0.100	0.655	1 consistent probesets
AI189287	qd05c04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-17	0.100	0.706	1 consistent probesets
W28892	53c11 Homo sapiens cDNA /gb=W28892 /gi=1308840 /	0.100	0.907	1 consistent probesets
AF042273	Homo sapiens signal transducing adaptor molecule 2A (S	0.100	0.958	1 consistent probesets
X76534	H.sapiens NMB mRNA /cds=(91,1773) /gb=X76534 /gi=666	0.100	1.008	1 consistent probesets
J04164	HUM927A Human interferon-inducible protein 9-	0.100	0.052	4 inconsistent probesets
D26599	HUMPSH2 Human mRNA for proteasome subunit HsC	0.100	0.057	4 inconsistent probesets
X83467	H.sapiens PXMP1 gene, exon 1 (and joined CDS) /cds=(0,	0.100	0.058	4 inconsistent probesets
X00129	Human mRNA for retinol binding protein (RBP) /cds=(51,	0.100	0.060	4 consistent probesets
AF061016	Homo sapiens UDP-glucose dehydrogenase (UGDH) mRNA	0.100	0.061	4 consistent probesets
L19779	HUMH2A2A Homo sapiens histone H2A.2 mRNA, com	0.100	0.064	4 consistent probesets
U51920	Human signal recognition particle (SRP54) mRNA, comple	0.100	0.075	4 consistent probesets
D32129	Human mRNA for HLA class-I (HLA-A26) heavy chain, com	0.100	0.083	4 consistent probesets
AI597616	tn15f08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-216	0.100	0.085	4 consistent probesets
S81003	S81003 L-UBC=ubiquitin conjugating enzyme [hu	0.100	0.090	4 consistent probesets
U09585	Homo sapiens putative interferon-related protein (SM15	0.100	0.093	4 consistent probesets
M83751	Human arginine-rich protein (ARP) gene, complete cds /	0.100	0.093	4 consistent probesets
AA149486	zl27g01.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-503	0.100	0.095	4 inconsistent probesets
M15205	HUMTKRA Human thymidine kinase gene, complete	0.100	0.097	4 consistent probesets
D14660	Human mRNA for KIAA0104 gene, complete cds /cds=(34,8	0.100	0.098	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
J02645	HUMEIF2A Human translational initiation facto	0.100	0.098	4 consistent probesets
X03363	HSERB2R Human c-erb-B-2 mRNA	0.100	0.098	4 inconsistent probesets
X68985	HSHEPLF H.sapiens mRNA for hepatic leukemia f	0.100	0.101	4 inconsistent probesets
AL039831	DKFZp434D1112_s1 Homo sapiens cDNA, 3 end /clone=D	0.100	0.105	4 consistent probesets
AA890010	aj89h08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-140	0.100	0.110	4 inconsistent probesets
W02490	za48b02.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-295	0.100	0.127	4 consistent probesets
X78678	H.sapiens KHK mRNA for ketohexokinase, clone pHKHK3a	0.100	0.134	1 consistent probesets
U84371	Human adenylate kinase 2A (AK2A) mRNA, complete cds /	0.100	0.135	4 consistent probesets
AF047469	Homo sapiens arsenite translocating ATPase (ASNA1) mRN	0.100	0.141	4 consistent probesets
J05448	HUMRPOLAA Human RNA polymerase subunit hRPB 3	0.100	0.169	4 consistent probesets
X12949	HSRETPON Human ret proto-oncogene mRNA for ty	0.100	0.202	1 consistent probesets
AF035582	Homo sapiens CASK mRNA, complete cds /cds=(15,2708)	0.100	0.207	4 consistent probesets
AI924382	wn60d01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.100	0.235	1 consistent probesets
M25393	HUMPTPASE Human protein tyrosine phosphatase	0.100	0.473	2 consistent probesets
AL049650	dJ734P14.2.2 (snRNP (small nuclear ribonucleoprotein p	0.098	0.034	8 inconsistent probesets
M64174	HUMPTKJAK1 Human protein-tyrosine kinase (JAK	0.098	0.091	8 consistent probesets
M14758	HUMMDR1 Homo sapiens P-glycoprotein (PGY1) mR	0.097	0.131	6 consistent probesets
U46920	Human metaxin (MTX) gene, complete cds /cds=(0,953) /g	0.096	0.055	4 consistent probesets
X59798	Human PRAD1 mRNA for cyclin /cds=(147,1034) /gb=X597	0.096	0.058	4 inconsistent probesets
X16396	Human mRNA for NAD-dependent methylene tetrahydrofola	0.096	0.061	4 consistent probesets
AF015926	Homo sapiens ezrin-radixin-moesin binding phosphoprote	0.096	0.062	4 consistent probesets
U18919	Human chromosome 17q12-21 mRNA, clone pOV-2, partia	0.096	0.062	8 consistent probesets
X86691	H.sapiens mRNA for 218kD Mi-2 protein /cds=(89,5827) /	0.096	0.064	4 consistent probesets
AL049981	Homo sapiens mRNA; cDNA DKFZp564D012 (from clone D	0.096	0.066	4 inconsistent probesets
D32050	Human mRNA for alanyl-tRNA synthetase, complete cds /c	0.096	0.068	4 consistent probesets
D50405	Human mRNA for RPD3 protein, complete cds /cds=(63,15	0.096	0.068	4 consistent probesets
AF055008	Homo sapiens clone 24720 epithelin 1 and 2 mRNA, compl	0.096	0.068	4 consistent probesets
M11433	Human cellular retinol-binding protein mRNA, complete	0.096	0.070	4 inconsistent probesets
U94831	Homo sapiens multispanning membrane protein mRNA, co	0.096	0.072	4 inconsistent probesets
AF047185	Homo sapiens NADH-ubiquinone oxidoreductase subunit C	0.096	0.078	4 consistent probesets
X51405	Human mRNA for carboxypeptidase E (EC 3.4.17.10) /cds=	0.096	0.081	4 consistent probesets
X56253	Human MPR46 gene for 46kd mannose 6-phosphate recept	0.096	0.082	4 consistent probesets
W28807	52a3 Homo sapiens cDNA /gb=W28807 /gi=1308755 /	0.096	0.085	4 consistent probesets
M17754	Human BN51 mRNA, complete cds /cds=(51,1238) /gb=M1	0.096	0.085	4 consistent probesets
D13146	HUM3CNP3 Homo sapiens gene for 2 ,3 -cyclic-n	0.096	0.086	4 consistent probesets
M13485	HUMMT1B2 Human metallothionein I-B gene, exon	0.096	0.087	4 consistent probesets
U41371	Human spliceosome associated protein (SAP 145) mRNA, c	0.096	0.088	4 consistent probesets
U59435	Human cell cycle protein p38-2G4 homolog (hG4-1) mRNA	0.096	0.088	4 consistent probesets
U18420	Human ras-related small GTP binding protein Rab5 (rab5	0.096	0.088	4 consistent probesets
D25278	Human mRNA for KIAA0036 gene, complete cds /cds=(156	0.096	0.091	4 inconsistent probesets
U90916	Human clone 23815 mRNA sequence /cds=UNKNOWN /gb	0.096	0.091	4 consistent probesets
R59606	yh02e06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-41	0.096	0.092	4 consistent probesets
U04953	Human isoleucyl-tRNA synthetase mRNA, complete cds /co	0.096	0.093	4 consistent probesets
U43923	HSU43923 Human transcription factor SUPT4H mR	0.096	0.096	4 consistent probesets
D14694	Human mRNA for KIAA0024 gene, complete cds /cds=(102	0.096	0.098	4 consistent probesets
AI223047	qg70a09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-18	0.096	0.098	4 inconsistent probesets
X94910	Homo sapiens mRNA for ERp28 protein /cds=(11,796) /gb=	0.096	0.098	4 consistent probesets
Z29064	H.sapiens AF-1p mRNA /cds=(92,2782) /gb=Z29064 /gi=47	0.096	0.101	4 consistent probesets
AF062006	Homo sapiens orphan G protein-coupled receptor HG38 mR	0.096	0.102	4 consistent probesets
AB014566	Homo sapiens mRNA for KIAA0666 protein, partial cds /c	0.096	0.105	4 inconsistent probesets
AL031983	dJ271M21.6 (Diubiquitin) /cds=(18,515) /gb=AL031983 /g	0.096	0.105	4 consistent probesets
D31766	Human mRNA for KIAA0060 gene, complete cds /cds=(401	0.096	0.110	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AI652978	wb42a05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.096	0.111	4 consistent probesets
U29175	Human transcriptional activator (BRG1) mRNA, complete	0.096	0.111	4 consistent probesets
U57721	Human L-kynurenine hydrolase mRNA, complete cds /cds=	0.096	0.112	4 consistent probesets
AI800499	tc11f11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-206	0.096	0.113	4 consistent probesets
L04658	Homo sapiens gene sequence /cds=UNKNOWN /gb=L0465	0.096	0.116	4 consistent probesets
AF057160	Homo sapiens putative poly(ADP-ribosyl) transferase (P	0.096	0.122	4 consistent probesets
D38435	Homo sapiens hPMS3 mRNA, partial cds /cds=(0,772) /gb=	0.096	0.125	4 consistent probesets
AB023219	Homo sapiens mRNA for KIAA1002 protein, complete cds /	0.096	0.128	4 inconsistent probesets
X13482	Human mRNA for U2 snRNP-specific A protein /cds=(56,82	0.096	0.128	4 consistent probesets
L40802	Homo sapiens 17-beta-hydroxysteroid dehydrogenase (17-	0.096	0.131	4 consistent probesets
U52427	Human RNA polymerase II seventh subunit (rpb-7) gene,	0.096	0.132	4 consistent probesets
M96803	Human general beta-spectrin (SPTBN1) mRNA, complete c	0.096	0.133	4 consistent probesets
U46689	Human microsomal aldehyde dehydrogenase (ALD10) mRN	0.096	0.136	4 consistent probesets
X14789	H.sapiens alpha-A crystallin gene exon 1,2 and pseudoe	0.096	0.137	4 inconsistent probesets
Z14000	H.sapiens RING1 gene /cds=(75,1208) /gb=Z14000 /gi=296	0.096	0.141	4 consistent probesets
X14830	Human mRNA for muscle acetylcholine receptor beta-subu	0.096	0.148	4 inconsistent probesets
L33075	HUMIQGA Homo sapiens ras GTPase-activating-li	0.096	0.155	4 consistent probesets
S52028	cystathionine gamma-lyase {clone HCL-1} [human, liver,	0.096	0.170	4 consistent probesets
X57766	Human stromelysin-3 mRNA /cds=(9,1475) /gb=X57766 /gi=	0.096	0.174	4 consistent probesets
AF043324	Homo sapiens N-myristoyltransferase 1 mRNA, complete c	0.096	0.177	4 consistent probesets
AL050157	Homo sapiens mRNA; cDNA DKFZp586O0120 (from clone	0.096	0.182	4 consistent probesets
AB028962	Homo sapiens mRNA for KIAA1039 protein, partial cds /c	0.096	0.186	4 consistent probesets
AL049923	Homo sapiens mRNA; cDNA DKFZp547E2210 (from clone	0.096	0.214	4 consistent probesets
AB029004	Homo sapiens mRNA for KIAA1081 protein, partial cds /c	0.096	0.226	4 consistent probesets
S67334	phosphatidylinositol 3-kinase p110 beta isoform=110 kd	0.096	0.234	4 consistent probesets
X16901	Human mRNA for RAP30 subunit of transcription initiati	0.096	0.238	4 consistent probesets
AB011161	Homo sapiens mRNA for KIAA0589 protein, partial cds /c	0.094	0.079	3 inconsistent probesets
M84562	Human formyl peptide receptor-like receptor (FPRL1) mR	0.094	0.169	3 consistent probesets
U66618	HSU66618 Human SWI/SNF complex 60 KDa subunit	0.094	0.070	8 inconsistent probesets
L11672	HUMKRUPZN Human Kruppel related zinc finger p	0.094	0.105	8 consistent probesets
U09759	HSU09759 Human protein kinase (JNK2) mRNA, co	0.094	0.120	8 consistent probesets
D82348	Homo sapiens mRNA for 5-aminoimidazole-4-carboxamide	0.092	0.055	4 consistent probesets
AF059293	Homo sapiens cytokine-like factor-1 precursor (CLF-1)	0.092	0.055	4 consistent probesets
X57398	Human mRNA for pM5 protein /cds=(0,3572) /gb=X57398 /(	0.092	0.057	4 inconsistent probesets
X06994	Human mRNA for cytochrome c1 /cds=(8,985) /gb=X06994	0.092	0.058	4 consistent probesets
X01060	Human mRNA for transferrin receptor /cds=(263,2545) /g	0.092	0.058	4 consistent probesets
U37690	HSU37690 Human RNA polymerase II subunit (hsR	0.092	0.060	4 inconsistent probesets
W28483	47e11 Homo sapiens cDNA /gb=W28483 /gi=1308431 /	0.092	0.061	4 inconsistent probesets
L09708	Human complement component 2 (C2) gene allele b /cds=(	0.092	0.067	4 consistent probesets
AF067139	Homo sapiens NADH-ubiquinone oxidoreductase NDUFS3	0.092	0.068	4 consistent probesets
M63167	HUMRACPC Human rac protein kinase alpha mRNA,	0.092	0.068	4 inconsistent probesets
U02570	HSU02570 Human CDC42 GTPase-activating protei	0.092	0.068	8 consistent probesets
W28498	50e2 Homo sapiens cDNA /gb=W28498 /gi=1308653 /	0.092	0.070	4 consistent probesets
L18960	HUMEIF4C Human protein synthesis factor (eIF-	0.092	0.072	8 inconsistent probesets
U81800	Homo sapiens monocarboxylate transporter (MCT3) mRNA	0.092	0.072	4 consistent probesets
D00760	HUMPSC3 Human mRNA for proteasome subunit HC3	0.092	0.072	4 inconsistent probesets
D25274	Homo sapiens mRNA, clone-PO2ST9 /cds=UNKNOWN /gb	0.092	0.072	4 consistent probesets
D90239	Human mRNA for glycine decarboxylase /cds=(150,3212) /	0.092	0.077	4 inconsistent probesets
U51478	Human sodium/potassium-transporting ATPase beta-3 subu	0.092	0.080	4 consistent probesets
AF038195	Homo sapiens clone 23661 unknown protein mRNA, comple	0.092	0.081	4 consistent probesets
AL080223	Homo sapiens mRNA; cDNA DKFZp566H2446 (from clone	0.092	0.081	4 consistent probesets
D87075	Human mRNA for KIAA0238 gene, partial cds /cds=(0,992)	0.092	0.082	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
X62534	H.sapiens HMG-2 mRNA /cds=(214,843) /gb=X62534 /gi=3	0.092	0.084	4 inconsistent probesets
D83782	Human mRNA for KIAA0199 gene, partial cds /cds=(0,3834	0.092	0.084	4 inconsistent probesets
U30521	Human P311 HUM (3.1) mRNA, complete cds /cds=(202,40	0.092	0.084	4 consistent probesets
AB014520	Homo sapiens mRNA for KIAA0620 protein, partial cds /c	0.092	0.085	4 consistent probesets
D38037	HUMOTK4 Human mRNA for FK506-binding protein	0.092	0.086	4 consistent probesets
M88163	Human global transcription activator homologous sequen	0.092	0.086	4 consistent probesets
M23254	Human Ca2-activated neutral protease large subunit (CA	0.092	0.087	4 consistent probesets
AI620381	tu94d05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-225	0.092	0.089	4 consistent probesets
D13892	Human mRNA for carboxyl methyltransferase, complete cd	0.092	0.091	4 consistent probesets
Z50749	H.sapiens sds22-like mRNA /cds=(15,1097) /gb=Z50749 /g	0.092	0.094	4 consistent probesets
M98343	Homo sapiens amplaxin (EMS1) mRNA, complete cds /cds=	0.092	0.094	4 consistent probesets
AF027299	Homo sapiens protein 4.1-G mRNA, complete cds /cds=(44	0.092	0.095	4 consistent probesets
D31767	Human mRNA for KIAA0058 gene, complete cds /cds=(69,5	0.092	0.098	4 consistent probesets
X55733	H.sapiens initiation factor 4B cDNA /cds=(0,1835) /gb=	0.092	0.100	4 consistent probesets
N90755	zb22c08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-302	0.092	0.104	4 consistent probesets
D85758	Homo sapiens mRNA for human protein homologous to DR	0.092	0.115	4 inconsistent probesets
M83822	Human beige-like protein (BGL) mRNA, partial cds /cds=	0.092	0.116	4 consistent probesets
M29877	Human alpha-L-fucosidase, complete cds /cds=UNKNOWN	0.092	0.116	4 consistent probesets
AF020543	Homo sapiens palmitoyl-protein thioesterase-2 (PPT2) m	0.092	0.122	4 consistent probesets
U48296	HSU48296 Homo sapiens protein tyrosine phosph	0.092	0.127	4 consistent probesets
AB019435	Homo sapiens mRNA for putative phospholipase, complete	0.092	0.132	4 consistent probesets
U41804	Human putative T1/ST2 receptor binding protein precurs	0.092	0.136	4 consistent probesets
D88308	Homo sapiens mRNA for very-long-chain acyl-CoA synthet	0.092	0.138	4 consistent probesets
L35035	Homo sapiens ribose 5-phosphate isomerase (RPI) mRNA /	0.092	0.141	4 consistent probesets
AF035940	Homo sapiens MAGOH mRNA, complete cds /cds=(65,505	0.092	0.150	4 consistent probesets
L20321	HUMSTK2A Human protein serine/threonine kinas	0.092	0.152	4 consistent probesets
L38933	Homo sapiens GT198 mRNA, complete ORF /cds=(133,744	0.092	0.158	4 consistent probesets
D80006	Human mRNA for KIAA0184 gene, partial cds /cds=(0,2591	0.092	0.158	4 consistent probesets
Y07867	H.sapiens mRNA for Pirin, isolate 1 /cds=(204,1076) /g	0.092	0.160	4 consistent probesets
AB018249	Homo sapiens gene for CC chemokine LEC, complete cds /	0.092	0.162	4 consistent probesets
Y11251	H.sapiens mRNA for novel member of serine-arginine dom	0.092	0.171	4 consistent probesets
D00860	Homo sapiens mRNA for phosphoribosyl pyrophosphate syn	0.092	0.172	4 consistent probesets
U67368	Human multiple exostosis 2 (EXT2) gene /cds=(30,2186)	0.092	0.179	4 consistent probesets
AF058922	Homo sapiens GLE1 (GLE1) mRNA, complete cds /cds=(87	0.092	0.180	4 consistent probesets
AB021663	Homo sapiens mRNA for leucine-zipper protein, complete	0.092	0.180	4 consistent probesets
AB023421	Homo sapiens mRNA for heat shock protein apg-1, comple	0.092	0.197	4 consistent probesets
AL120374	DKFZp761C088_r1 Homo sapiens cDNA, 5 end /clone=DK	0.092	0.200	4 consistent probesets
U31317	HSU31317 Human JAK family tyrosine protein ki	0.092	0.226	2 consistent probesets
AW052084	wy86f07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.092	0.237	4 consistent probesets
AF070557	Homo sapiens clone 24422 mRNA sequence /cds=UNKNO	0.092	0.298	4 consistent probesets
M93650	Human paired box gene (PAX6) homologue, complete cds /	0.092	0.316	4 consistent probesets
AF070595	Homo sapiens clone 24583 mRNA sequence /cds=UNKNO	0.092	0.320	4 consistent probesets
X17033	HSINAL2 Human mRNA for integrin alpha-2 subun	0.090	0.246	5 consistent probesets
X99906	Homo sapiens mRNA for alpha endosulfine /cds=(125,490)	0.090	0.069	8 inconsistent probesets
D88378	D88378 Homo sapiens mRNA for proteasome inhib	0.090	0.112	8 consistent probesets
AA846749	aj99c10.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-140	0.088	0.053	8 inconsistent probesets
U72511	Human B-cell receptor associated protein (hBAP) mRNA,	0.088	0.054	4 consistent probesets
D28423	HUMPSF82 Human mRNA for pre-mRNA splicing fac	0.088	0.058	4 inconsistent probesets
U53347	Human neutral amino acid transporter B mRNA, complete	0.088	0.058	4 consistent probesets
U29953	Human pigment epithelium-derived factor gene, complete	0.088	0.061	4 consistent probesets
M90657	HUML6A Human tumor antigen (L6) mRNA, complet	0.088	0.061	4 consistent probesets
U31875	Human Hep27 protein mRNA, complete cds /cds=(433,1275	0.088	0.061	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
L10678	Human profilin II mRNA, complete cds /cds=(13,435) /gb	0.088	0.061	4 consistent probesets
D00761	HUMPSC5 Human mRNA for proteasome subunit HC5	0.088	0.061	4 consistent probesets
D13748	HUM4AI Human mRNA for eukaryotic initiation f	0.088	0.068	4 consistent probesets
AF080561	Homo sapiens SYT interacting protein SIP mRNA, complet	0.088	0.068	4 consistent probesets
M91432	Human medium-chain acyl-CoA dehydrogenase (MCAD) ge	0.088	0.070	4 inconsistent probesets
AI819948	wj88e11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24(	0.088	0.072	4 inconsistent probesets
U51004	HSU51004 Homo sapiens protein kinase C inhibi	0.088	0.072	4 consistent probesets
X68733	H.sapiens gene for alpha1-antichymotrypsin, exon 1 /cd	0.088	0.072	4 inconsistent probesets
U27460	Human uridine diphosphoglucose pyrophosphorylase mRNA	0.088	0.075	4 inconsistent probesets
AB023208	Homo sapiens mRNA for KIAA0991 protein, complete cds /	0.088	0.075	4 consistent probesets
L12723	HUMHSP70H Human heat shock protein 70 (hsp70)	0.088	0.075	4 consistent probesets
M58525	Homo sapiens catechol-O-methyltransferase (COMT) mRN	0.088	0.075	4 inconsistent probesets
AI708983	at02f03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-235	0.088	0.078	4 inconsistent probesets
M24486	Human prolyl 4-hydroxylase alpha subunit mRNA, complet	0.088	0.081	4 inconsistent probesets
AL096719	Homo sapiens mRNA; cDNA DKFZp566N043 (from clone D	0.088	0.082	4 consistent probesets
D38073	Human mRNA for hRlf beta subunit (p102 protein), compl	0.088	0.082	4 consistent probesets
D83004	D83004 Human epidermoid carcinoma mRNA for ub	0.088	0.084	8 consistent probesets
AI249721	qj64d06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-186	0.088	0.085	4 consistent probesets
AF000364	Homo sapiens heterogeneous nuclear ribonucleoprotein R	0.088	0.086	4 consistent probesets
AF052158	Homo sapiens clone 24616 mRNA sequence /cds=UNKNO	0.088	0.089	4 consistent probesets
AB023224	Homo sapiens mRNA for KIAA1007 protein, partial cds /c	0.088	0.100	4 consistent probesets
D86967	Human mRNA for KIAA0212 gene, complete cds /cds=(58,2	0.088	0.101	4 consistent probesets
AI985272	ws06b05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.088	0.109	4 consistent probesets
AB004857	Homo sapiens mRNA for NRAMP2, complete cds /cds=(88,	0.088	0.109	4 consistent probesets
AF010309	Homo sapiens Pig3 (PIG3) mRNA, complete cds /cds=(527	0.088	0.112	4 consistent probesets
AF029777	Homo sapiens histone acetyltransferase (GCN5) mRNA, pa	0.088	0.116	4 consistent probesets
L48692	Homo sapiens (clone p5-23-3) mRNA /cds=UNKNOWN /gb	0.088	0.119	4 inconsistent probesets
L13738	HUMNRTYKIN Human activated p21cdc42Hs kinase	0.088	0.120	4 consistent probesets
AI341574	qq94h09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-19	0.088	0.124	4 consistent probesets
AJ002428	Homo sapiens VDAC1 pseudogene /cds=(0,853) /gb=AJ002	0.088	0.125	4 inconsistent probesets
M30818	HUMMXB Human interferon-induced cellular resi	0.088	0.127	4 consistent probesets
U67171	Human selenoprotein W (selW) mRNA, complete cds /cds=	0.088	0.140	4 consistent probesets
AF022912	Homo sapiens cGMP phosphodiesterase delta subunit mRN	0.088	0.142	4 consistent probesets
D79992	Human mRNA for KIAA0170 gene, complete cds /cds=(13,6	0.088	0.142	4 consistent probesets
L76159	Homo sapiens FRG1 mRNA, complete cds /cds=(191,967)	0.088	0.146	4 consistent probesets
AF043250	Homo sapiens mitochondrial outer membrane protein (TOM	0.088	0.151	4 consistent probesets
AL050259	Homo sapiens mRNA; cDNA DKFZp547D0710 (from clone	0.088	0.151	4 consistent probesets
AI888084	wm29g09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2	0.088	0.155	4 consistent probesets
U59919	Human Smg GDS-associated protein SMAP mRNA, comple	0.088	0.156	4 consistent probesets
AF006259	Homo sapiens Rad51-interacting protein mRNA, complete	0.088	0.193	4 consistent probesets
W28299	44h4 Homo sapiens cDNA /gb=W28299 /gi=1308247 /	0.088	0.194	4 consistent probesets
AB000359	Homo sapiens PIGCP1 pseudogene /cds=(0,416) /gb=AB00	0.088	0.230	4 consistent probesets
AF065482	Homo sapiens sorting nexin 2 (SNX2) mRNA, complete cds	0.088	0.230	4 consistent probesets
U15085	Human HLA-DMB mRNA, complete cds /cds=(233,1024) /g	0.088	0.240	4 consistent probesets
AF061836	Homo sapiens putative tumor suppressor protein (RDA32)	0.088	0.249	4 consistent probesets
AF005361	HUMIMPA6 Homo sapiens importin alpha 6 mRNA,	0.088	0.290	4 consistent probesets
Z23115	HSBCLXL H.sapiens bcl-xL mRNA	0.085	0.063	8 consistent probesets
D12485	HUMNPP Human mRNA for nucleotide pyrophosphat	0.085	0.096	8 consistent probesets
AF047042	Homo sapiens citrate synthase mRNA, complete cds /cds=	0.083	0.050	4 inconsistent probesets
AL022721	dJ109F14.1.1 (Transcriptional Enhancer Factor TEF-5)(i	0.083	0.056	8 consistent probesets
D00096	Homo sapiens mRNA for prealbumin, complete cds /cds=(2	0.083	0.058	4 inconsistent probesets
Y18007	Homo sapiens mRNA for putative seven transmembrane do	0.083	0.064	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
X73424	Homo sapiens gene for propionyl-CoA carboxylase a subu	0.083	0.064	4 inconsistent probesets
D16481	Homo sapiens mRNA for mitochondrial 3-ketoacyl-CoA thi	0.083	0.064	4 inconsistent probesets
AJ001340	Homo sapiens mRNA for U3 snoRNP associated 55 kDa pr	0.083	0.069	3 consistent probesets
AF002163	Homo sapiens delta-adaptin mRNA, complete cds /cds=(31	0.083	0.070	8 consistent probesets
M80482	Human subtilisin-like protein (PACE4) mRNA, complete c	0.083	0.075	4 consistent probesets
AF087135	Homo sapiens F1FO-type ATPase subunit d mRNA, nuclea	0.083	0.075	4 consistent probesets
S67247	smooth muscle myosin heavy chain isoform SMemb [huma	0.083	0.075	4 consistent probesets
M95787	Human 22kDa smooth muscle protein (SM22) mRNA, comp	0.083	0.076	4 consistent probesets
L23959	HUMDP1A Homo sapiens E2F-related transcriptio	0.083	0.077	6 inconsistent probesets
X66436	H.sapiens hsr1 mRNA (partial) /cds=UNKNOWN /gb=X664	0.083	0.078	5 consistent probesets
AF037261	Homo sapiens SH3-containing adaptor molecule-1 mRNA, c	0.083	0.078	4 consistent probesets
L07956	Homo sapiens 1,4-alpha-glucan branching enzyme (HGBE)	0.083	0.080	4 consistent probesets
L13435	Human chromosome 3p21.1 gene sequence /cds=UNKNO	0.083	0.085	4 consistent probesets
M57729	Human complement component C5 mRNA, complete cds /c	0.083	0.090	4 consistent probesets
AF025887	Homo sapiens glutathione S-transferase A4-4 (GSTA4) mR	0.083	0.093	4 inconsistent probesets
U40282	Homo sapiens integrin-linked kinase (ILK) mRNA, comple	0.083	0.093	4 consistent probesets
AA005018	zh96a09.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-429	0.083	0.096	4 consistent probesets
U89505	Human Hlark mRNA, complete cds /cds=(55,1155) /gb=U89	0.083	0.098	4 consistent probesets
D43947	Human mRNA for KIAA0100 gene, complete cds /cds=(329	0.083	0.098	4 consistent probesets
AF038187	Homo sapiens clone 23714 mRNA sequence /cds=UNKNO	0.083	0.098	4 inconsistent probesets
J04543	Human synexin mRNA, complete cds /cds=(60,1460) /gb=J	0.083	0.098	4 consistent probesets
D29805	Human mRNA for beta-1,4-galactosyltransferase, complet	0.083	0.103	4 consistent probesets
X95735	Homo sapiens mRNA for zyxin /cds=(71,1789) /gb=X95735	0.083	0.104	3 consistent probesets
AL080084	Homo sapiens mRNA; cDNA DKFZp564G2362 (from clone	0.083	0.104	4 consistent probesets
M61916	HUMLAM101 Human laminin B1 chain mRNA, comple	0.083	0.107	4 consistent probesets
M79463	HUMPML2 Human PML-2 mRNA, complete CDS"	0.083	0.108	3 inconsistent probesets
W26634	34b10 Homo sapiens cDNA /gb=W26634 /gi=1307477 /	0.083	0.112	4 consistent probesets
AF052182	Homo sapiens clone 24590 mRNA sequence /cds=UNKNO	0.083	0.113	4 consistent probesets
AF038250	AF038250 Homo sapiens cDNA /clone=ntcon9 /gb=AF0382	0.083	0.114	4 consistent probesets
AB023205	Homo sapiens mRNA for KIAA0988 protein, complete cds /	0.083	0.116	3 consistent probesets
AI432190	tg77f11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-211	0.083	0.121	4 consistent probesets
M28213	HUMRAB2A Homo sapiens GTP-binding protein (RA	0.083	0.124	4 consistent probesets
D50931	Human mRNA for KIAA0141 gene, complete cds /cds=(80,1	0.083	0.128	4 consistent probesets
AI201108	qf69g07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-175	0.083	0.129	4 consistent probesets
X54871	HSRAB5B H.sapiens mRNA for ras-related protei	0.083	0.130	4 consistent probesets
AI741756	wg22e12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.083	0.131	4 consistent probesets
AB020713	Homo sapiens mRNA for KIAA0906 protein, partial cds /c	0.083	0.137	4 consistent probesets
J03824	Human uroporphyrinogen III synthase mRNA, complete cds	0.083	0.138	5 consistent probesets
AB014561	Homo sapiens mRNA for KIAA0661 protein, complete cds /	0.083	0.139	3 consistent probesets
D21337	HUMCO Human mRNA for collagen	0.083	0.139	5 consistent probesets
AB029038	Homo sapiens mRNA for KIAA1115 protein, complete cds /	0.083	0.141	3 consistent probesets
U07158	Human syntaxin mRNA, complete cds /cds=(66,959) /gb=U	0.083	0.141	4 consistent probesets
L26339	Human autoantigen mRNA, complete cds /cds=(136,3783) ,	0.083	0.145	3 consistent probesets
X14787	HSTS Human mRNA for thrombospondin	0.083	0.146	3 consistent probesets
U90917	Human clone 23641 mRNA sequence /cds=UNKNOWN /gb	0.083	0.147	3 consistent probesets
Z81326	H.sapiens mRNA for protease inhibitor 12 (PI12; neuros	0.083	0.147	3 consistent probesets
AB023187	Homo sapiens mRNA for KIAA0970 protein, complete cds /	0.083	0.152	4 consistent probesets
U79241	Human clone 23759 mRNA, partial cds /cds=(0,1315) /gb=	0.083	0.156	3 consistent probesets
AB015718	Homo sapiens lok mRNA for protein kinase, complete cds	0.083	0.159	3 consistent probesets
U11292	Human Ki nuclear autoantigen mRNA, complete cds /cds=(	0.083	0.165	3 consistent probesets
U18271	Human thymopoietin (TMPO) gene /cds=(313,2397) /gb=U1	0.083	0.165	4 inconsistent probesets
L13434	Human chromosome 3p21.1 gene sequence, complete cds	0.083	0.169	3 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF091087	Homo sapiens clone 643 unknown mRNA, complete sequer	0.083	0.169	3 consistent probesets
W28504	48e7 Homo sapiens cDNA /gb=W28504 /gi=1308515 /	0.083	0.172	3 consistent probesets
U42408	Human ladinin (LAD) mRNA, complete cds /cds=(219,1772	0.083	0.179	3 consistent probesets
U77327	Human Ki-1/57 intracellular antigen mRNA, partial cds	0.083	0.179	3 consistent probesets
AI885852	wl62d08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.083	0.181	3 consistent probesets
S68134	CREM=cyclic AMP-responsive element modulator beta isof	0.083	0.181	4 consistent probesets
W26334	26b1 Homo sapiens cDNA /gb=W26334 /gi=1306889 /	0.083	0.182	4 consistent probesets
AB014548	Homo sapiens mRNA for KIAA0648 protein, partial cds /c	0.083	0.183	4 consistent probesets
H68340	yr82b10.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-211	0.083	0.186	3 consistent probesets
X58521	Human mRNA for p62 nucleoporin /cds=(151,1719) /gb=X5	0.083	0.191	3 consistent probesets
Y15906	Homo sapiens mRNA for XAGL protein /cds=(132,1646) /gb	0.083	0.194	3 consistent probesets
AF070552	Homo sapiens clone 24767 mRNA sequence /cds=UNKNO	0.083	0.196	3 consistent probesets
AI889718	wo17c04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.083	0.207	3 consistent probesets
U90551	Human histone 2A-like protein (H2A/l) mRNA, complete c	0.083	0.210	4 consistent probesets
D49354	HUMHSP70A Human mRNA for enhancer protein in	0.083	0.215	3 consistent probesets
AB007893	Homo sapiens KIAA0433 mRNA, partial cds /cds=(509,424	0.083	0.218	4 consistent probesets
AI744294	tr08h04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-221	0.083	0.252	2 consistent probesets
Z35094	H.sapiens mRNA for SURF-2 /cds=(17,787) /gb=Z35094 /gi	0.083	0.254	3 consistent probesets
U20536	Human cysteine protease Mch2 isoform alpha (Mch2) mRN	0.083	0.263	3 consistent probesets
D37965	HUMPRLTS Human mRNA for PDGF receptor beta-li	0.083	0.284	3 consistent probesets
AF040965	Homo sapiens unknown protein IT12 mRNA, partial cds /c	0.083	0.434	3 consistent probesets
U35113	HSU35113 Human metastasis-associated mta1 mRN	0.082	0.057	10 consistent probesets
X55885	Human mRNA for a presumptive KDEL receptor /cds=(146,	0.081	0.072	8 inconsistent probesets
AF052155	Homo sapiens clone 24761 mRNA sequence /cds=UNKNO	0.081	0.087	8 consistent probesets
D26350	HUMHT2I Human mRNA for type 2 inositol 1,4,5-	0.081	0.124	8 inconsistent probesets
M68520	HUMCDC2A Human cdc2-related protein kinase mR	0.081	0.083	7 inconsistent probesets
S75174	S75174 E2F-4=transcription factor [human, Nal	0.081	0.101	7 consistent probesets
L19686	HUMMIF Homo sapiens macrophage migration inhi	0.079	0.047	4 inconsistent probesets
M93651	Human set gene, complete cds /cds=(3,836) /gb=M93651 /	0.079	0.047	4 consistent probesets
S95936	transferrin [human, liver, mRNA, 2347 nt] /cds=(79,217	0.079	0.050	4 consistent probesets
Z84718	HS322B1 Human DNA sequence from clone 322B1 o	0.079	0.052	8 inconsistent probesets
AF091085	Homo sapiens clone 638 unknown mRNA, complete sequer	0.079	0.055	4 consistent probesets
U12595	HSU12595 Human tumor necrosis factor type 1 r	0.079	0.055	4 consistent probesets
T58471	yb61c11.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-756	0.079	0.055	4 consistent probesets
L26232	Human phospholipid transfer protein mRNA, complete cds	0.079	0.055	4 inconsistent probesets
T57872	yb19b12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-71	0.079	0.055	4 consistent probesets
K02581	Human thymidine kinase mRNA, complete cds /cds=(57,76	0.079	0.055	4 consistent probesets
AB019409	Homo sapiens mRNA, expressed in fibroblasts of periodo	0.079	0.058	4 consistent probesets
AB016492	Homo sapiens hJTB gene, complete cds /cds=(464,904) /g	0.079	0.058	8 consistent probesets
U47101	Human NifU-like protein (hNifU) mRNA, partial cds /cds	0.079	0.061	4 inconsistent probesets
D49738	Human cytoskeleton associated protein (CG22) mRNA, con	0.079	0.061	4 inconsistent probesets
X15940	Human mRNA for ribosomal protein L31 /cds=(7,384) /gb=	0.079	0.061	4 consistent probesets
D63878	Human mRNA for KIAA0158 gene, complete cds /cds=(258	0.079	0.063	4 inconsistent probesets
D42041	Human mRNA for KIAA0088 gene, partial cds /cds=(0,2832	0.079	0.064	4 consistent probesets
U62962	Human Int-6 mRNA, complete cds /cds=(22,1359) /gb=U62	0.079	0.064	4 consistent probesets
AF035606	Homo sapiens calcium binding protein (ALG-2) mRNA, com	0.079	0.064	4 inconsistent probesets
AI803447	tc39g04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-206	0.079	0.065	4 consistent probesets
AI541285	pec1.2-4.D10.r Homo sapiens cDNA, 5 end /clone_end=5	0.079	0.065	4 consistent probesets
X57346	H.sapiens mRNA for HS1 protein /cds=(372,1112) /gb=X57	0.079	0.066	4 consistent probesets
X69907	H.sapiens gene for mitochondrial ATP synthase c subuni	0.079	0.067	4 inconsistent probesets
X87342	H.sapiens mRNA for human giant larvae homolog /cds=(13	0.079	0.068	4 consistent probesets
AI052724	oz27a12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.079	0.068	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D13435	HUMPIGF Human mRNA for PIG-F (phosphatidyl-in	0.079	0.070	4 inconsistent probesets
D84111	Homo sapiens mRNA for RBP-MS/type 5, partial cds /cds=	0.079	0.072	8 consistent probesets
L20298	Homo sapiens transcription factor (CBFB) mRNA, 3 end	0.079	0.072	4 consistent probesets
U34804	Human thermostable phenol sulfotransferase (STP2) gene	0.079	0.072	4 consistent probesets
X06323	Human MRL3 mRNA for ribosomal protein L3 homologue (	0.079	0.074	4 consistent probesets
M11730	Human tyrosine kinase-type receptor (HER2) mRNA, compl	0.079	0.075	4 inconsistent probesets
D00265	Homo sapiens mRNA for cytochrome c, partial cds /cds=(	0.079	0.075	4 consistent probesets
AI760162	wg58e09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.079	0.077	8 inconsistent probesets
U97067	Homo sapiens alpha-catenin-like protein mRNA, complete	0.079	0.077	4 consistent probesets
W28944	54h12 Homo sapiens cDNA /gb=W28944 /gi=1308955 /	0.079	0.078	4 consistent probesets
Z49254	H.sapiens L23-related mRNA /cds=(54,515) /gb=Z49254 /g	0.079	0.080	4 consistent probesets
AF026029	Homo sapiens poly(A) binding protein II (PABP2) gene,	0.079	0.085	4 consistent probesets
AB003177	AB003177 Homo sapiens mRNA for proteasome sub	0.079	0.085	4 consistent probesets
J02966	Human mitochondrial ADP/ADT translocator mRNA, comple	0.079	0.085	4 inconsistent probesets
S69189	peroxisomal acyl-coenzyme A oxidase [human, liver, mRN	0.079	0.085	4 consistent probesets
U95006	Human D9 splice variant A mRNA, complete cds /cds=(3,1	0.079	0.088	4 consistent probesets
M20471	Human brain-type clathrin light-chain a mRNA, complete	0.079	0.088	4 consistent probesets
S90469	S90469 cytochrome P450 reductase [human, plac	0.079	0.090	4 consistent probesets
J03040	HUMSPARC Human SPARC/osteonectin mRNA, comple	0.079	0.091	4 consistent probesets
AL050224	Homo sapiens mRNA; cDNA DKFZp586L2123 (from clone	0.079	0.091	4 consistent probesets
U89606	Human pyridoxal kinase mRNA, complete cds /cds=(6,944)	0.079	0.095	4 consistent probesets
U17714	HSU17714 Homo sapiens putative tumor suppress	0.079	0.095	4 consistent probesets
AI565760	tn20b01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-216	0.079	0.098	4 consistent probesets
AF053003	Homo sapiens diphthamide biosynthesis protein-2 (DPH2)	0.079	0.101	4 consistent probesets
Y12478	H.sapiens mRNA for CHD5 protein /cds=(42,566) /gb=Y124	0.079	0.102	4 inconsistent probesets
AL050395	Homo sapiens mRNA; cDNA DKFZp586D1020 (from clone	0.079	0.103	4 consistent probesets
U89436	Human tyrosyl-tRNA synthetase mRNA, complete cds /cds=	0.079	0.103	4 consistent probesets
W28235	43h8 Homo sapiens cDNA /gb=W28235 /gi=1308183 /	0.079	0.104	4 consistent probesets
L10373	Human (clone CCG-B7) mRNA sequence /cds=UNKNOWN	0.079	0.107	4 consistent probesets
U88528	Human transcription factor LZIP mRNA, complete cds /cd	0.079	0.111	4 consistent probesets
U96131	Homo sapiens HPV16 E1 protein binding protein mRNA, co	0.079	0.111	4 consistent probesets
AA176780	zp32a10.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-61	0.079	0.113	8 consistent probesets
X99656	H.sapiens mRNA for protein containing SH3 domain, SH3G	0.079	0.113	4 consistent probesets
AF054179	Homo sapiens H beta 58 homolog mRNA, complete cds /cd	0.079	0.114	4 consistent probesets
AI347155	tc04c11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-206	0.079	0.123	4 consistent probesets
AF073308	Homo sapiens nonsyndromic hearing impairment protein (	0.079	0.132	4 consistent probesets
X93086	H.sapiens mRNA for biliverdin IX alpha reductase /cds=	0.079	0.135	4 consistent probesets
M84349	Human transmembrane protein (CD59) gene /cds=(18,404)	0.079	0.135	4 consistent probesets
AF027208	Homo sapiens AC133 antigen mRNA, complete cds /cds=(3	0.079	0.140	4 consistent probesets
X02308	Human mRNA for thymidylate synthase (EC 2.1.1.45) /cds	0.079	0.142	4 consistent probesets
AW026535	wv14f10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-99	0.079	0.145	4 inconsistent probesets
U16028	HSU16028 Human CRE-BP1 transcription factor m	0.079	0.146	4 inconsistent probesets
X80754	Homo sapiens mRNA for GTP-binding protein /cds=(47,114	0.079	0.155	4 consistent probesets
U34038	Human proteinase-activated receptor-2 mRNA, complete c	0.079	0.162	4 consistent probesets
AF052181	Homo sapiens clone 24790 mRNA sequence /cds=UNKNO	0.079	0.163	4 consistent probesets
U02390	Human adenylyl cyclase-associated protein homolog CAP2	0.079	0.164	4 consistent probesets
AI701164	we10g07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.079	0.164	4 consistent probesets
S78085	S78085 PDCD2=programmed cell death-2/Rp8 homo	0.079	0.166	4 consistent probesets
AB023213	Homo sapiens mRNA for KIAA0996 protein, complete cds /	0.079	0.167	4 consistent probesets
U69141	Human glutaryl-CoA dehydrogenase mRNA, complete cds /	0.079	0.168	4 consistent probesets
AA015605	ze20c12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-35	0.079	0.171	4 consistent probesets
U10117	Human endothelial-monocyte activating polypeptide II m	0.079	0.184	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF059531	Homo sapiens protein arginine N-methyltransferase 3 (P	0.079	0.184	4 consistent probesets
R90942	yp92b03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-194	0.079	0.185	4 consistent probesets
D45288	HUMHG2121 Homo sapiens cDNA /gb=D45288 /gi=113668	0.079	0.185	4 consistent probesets
L16895	Human lysyl oxidase (LOX) gene, exon 7 /cds=(0,6) /gb=	0.079	0.202	4 consistent probesets
U46194	HSU46194 Human renal cell carcinoma antigen R	0.079	0.230	4 consistent probesets
M86852	Homo sapiens peroxisome assembly factor-1 (PAF-1) mRN	0.079	0.253	4 consistent probesets
AB014579	Homo sapiens mRNA for KIAA0679 protein, partial cds /c	0.079	0.260	4 consistent probesets
AF007872	Homo sapiens torsinB (DQ1) mRNA, partial cds /cds=(0,8	0.079	0.294	4 consistent probesets
AB000509	Homo sapiens mRNA for TRAF5, complete cds /cds=(54,17	0.079	0.297	4 consistent probesets
AL079286	Homo sapiens mRNA full length insert cDNA clone EUROIN	0.079	0.335	4 consistent probesets
U09510	Human glycyl-tRNA synthetase mRNA, complete cds /cds=	0.077	0.039	8 inconsistent probesets
Y08302	HSMAPKP4 H.sapiens mRNA for MAP kinase phosph	0.077	0.048	8 inconsistent probesets
J03473	HUMRISDAD Human poly(ADP-ribose) synthetase m	0.077	0.062	8 consistent probesets
J04182	Homo sapiens lysosomal membrane glycoprotein-1 (LAMP1	0.075	0.044	4 inconsistent probesets
M14199	HUMLAMR Human laminin receptor (2H5 epitope)	0.075	0.048	4 inconsistent probesets
J00077	Human alpha-fetoprotein (AFP) mRNA, complete cds /cds=	0.075	0.053	4 consistent probesets
AI525665	PT1.3_04_D06.r Homo sapiens cDNA, 5 end /clone_end=5	0.075	0.055	4 inconsistent probesets
U32986	HSU32986 Human xeroderma pigmentosum group E	0.075	0.062	4 inconsistent probesets
U68566	Human HS1 binding protein HAX-1 mRNA, nuclear gene en	0.075	0.064	4 inconsistent probesets
AI651806	wb55f10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.075	0.064	4 inconsistent probesets
Z48950	H.sapiens hH3.3B gene for histone H3.3 /cds=(10,420) /	0.075	0.067	4 consistent probesets
M64982	Human fibrinogen alpha chain gene, complete mRNAs /cds	0.075	0.068	4 inconsistent probesets
AI526078	DU3.2-7.G08.r Homo sapiens cDNA, 5 end /clone_end=5	0.075	0.068	4 consistent probesets
D50916	Human mRNA for KIAA0126 gene, complete cds /cds=(72,3	0.075	0.069	4 consistent probesets
M19311	HUMCAM Human calmodulin mRNA, complete cds"	0.075	0.070	4 consistent probesets
AL050396	Homo sapiens mRNA; cDNA DKFZp586K1720 (from clone	0.075	0.070	4 inconsistent probesets
M27396	Human asparagine synthetase mRNA, complete cds /cds=(	0.075	0.072	4 inconsistent probesets
AL050361	Homo sapiens mRNA; cDNA DKFZp564H0223 (from clone	0.075	0.076	4 consistent probesets
AL041493	DKFZp434F2117_s1 Homo sapiens cDNA, 3 end /clone=D	0.075	0.080	4 inconsistent probesets
AJ005168	Homo sapiens KHK gene, exons 4-8 /cds=(0,552) /gb=AJ00	0.075	0.080	4 inconsistent probesets
AL050051	Homo sapiens mRNA; cDNA DKFZp566D193 (from clone D	0.075	0.081	4 consistent probesets
AF006084	Homo sapiens Arp2/3 protein complex subunit p41-Arc (A	0.075	0.083	4 consistent probesets
AB007857	Homo sapiens KIAA0397 mRNA, complete cds /cds=(54,15	0.075	0.084	4 consistent probesets
D80008	Human mRNA for KIAA0186 gene, complete cds /cds=(94,6	0.075	0.085	4 inconsistent probesets
AA648295	ns20e08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-11	0.075	0.085	4 consistent probesets
D89937	Homo sapiens mRNA for follistatin-related protein (FRP	0.075	0.086	4 consistent probesets
X54326	H.sapiens mRNA for glutaminyl-tRNA synthetase /cds=(58	0.075	0.087	4 consistent probesets
AB018335	Homo sapiens mRNA for KIAA0792 protein, complete cds /	0.075	0.091	4 consistent probesets
L07597	HUMS6KINA Homo sapiens ribosomal protein S6 k	0.075	0.091	2 consistent probesets
J04599	Human hPGI mRNA encoding bone small proteoglycan I (b	0.075	0.091	2 consistent probesets
AJ005895	Homo sapiens mRNA for (JM3) preprotein translocase, co	0.075	0.091	2 consistent probesets
AF007151	Homo sapiens clone 23967 unknown mRNA, partial cds /cd	0.075	0.093	4 consistent probesets
AL050091	Homo sapiens mRNA; cDNA DKFZp586F1918 (from clone	0.075	0.096	4 consistent probesets
AF051894	Homo sapiens 15 kDa selenoprotein mRNA, complete cds /	0.075	0.097	4 inconsistent probesets
L25931	HUMLBR Human lamin B receptor (LBR) mRNA, com	0.075	0.098	4 consistent probesets
Y15409	Homo sapiens mRNA for putative glucose 6-phosphate tra	0.075	0.098	4 consistent probesets
D83779	Human mRNA for KIAA0195 gene, complete cds /cds=(203	0.075	0.101	4 consistent probesets
M92439	Human leucine-rich protein mRNA, complete cds /cds=UNK	0.075	0.105	4 consistent probesets
X04325	Human liver mRNA for gap junction protein /cds=(62,913	0.075	0.107	4 consistent probesets
D13629	Human mRNA for KIAA0004 gene, complete cds /cds=(83,3	0.075	0.112	4 consistent probesets
X95073	H.sapiens mRNA for translin associated protein X /cds=	0.075	0.112	4 consistent probesets
Z22865	H.sapiens dermatopontin mRNA, complete CDS /cds=(12,6	0.075	0.113	2 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
Z83844	Human DNA sequence from clone 37E16 on chromosome 2	0.075	0.113	2 consistent probesets
AB014552	Homo sapiens mRNA for KIAA0652 protein, complete cds /	0.075	0.113	2 consistent probesets
U79716	Human reelin (RELN) mRNA, complete cds /cds=(175,1055	0.075	0.116	4 consistent probesets
L40402	Homo sapiens (clone Zap2) mRNA fragment /cds=UNKNOV	0.075	0.117	4 consistent probesets
M88279	Human immunophilin (FKBP52) mRNA, complete cds /cds=	0.075	0.118	4 consistent probesets
U47927	Human isopeptidase T (ISOT) mRNA, complete cds /cds=(5	0.075	0.124	4 consistent probesets
AB002315	Human mRNA for KIAA0317 gene, complete cds /cds=(471	0.075	0.127	4 consistent probesets
X75621	Homo sapiens TSC2 mRNA for tuberin /cds=(18,5441) /gb=	0.075	0.128	4 consistent probesets
AI246726	qk40a09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-18	0.075	0.129	4 consistent probesets
AI674208	wc07f02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.075	0.135	4 inconsistent probesets
AB014574	Homo sapiens mRNA for KIAA0674 protein, partial cds /c	0.075	0.136	4 consistent probesets
M30269	Human nidogen mRNA, complete cds /cds=(90,3833) /gb=M	0.075	0.141	4 consistent probesets
Z80345	H.sapiens SCAD gene, 5 UTR exon 1 and 2 (and joined CD	0.075	0.141	4 consistent probesets
AL050286	Homo sapiens mRNA; cDNA DKFZp586A011 (from clone D	0.075	0.142	4 consistent probesets
D21261	Human mRNA for KIAA0120 gene, complete cds /cds=(73,6	0.075	0.147	4 consistent probesets
W27522	32a12 Homo sapiens cDNA /gb=W27522 /gi=1307326 /	0.075	0.147	4 consistent probesets
AL080118	Homo sapiens mRNA; cDNA DKFZp564F1123 (from clone	0.075	0.150	4 inconsistent probesets
L11931	Human cytosolic serine hydroxymethyltransferase (SHMT)	0.075	0.151	4 consistent probesets
W26407	29b8 Homo sapiens cDNA /gb=W26407 /gi=1307106 /	0.075	0.153	4 consistent probesets
Y18418	Homo sapiens mRNA for erythrocyte cytosolic protein of	0.075	0.158	4 consistent probesets
U01147	Human guanine nucleotide regulatory protein (ABR) mRNA	0.075	0.169	2 consistent probesets
Y09616	H.sapiens mRNA for putative carboxylesterase /cds=(78,	0.075	0.182	2 consistent probesets
U87408	Human clone IMAGE-74593 unknown protein mRNA, partia	0.075	0.184	2 consistent probesets
AF039241	AF039241 Homo sapiens cDNA /clone=11-67js /gb=AF039	0.075	0.197	2 consistent probesets
X52213	H.sapiens ltk mRNA /cds=(258,1652) /gb=X52213 /gi=3442	0.075	0.197	2 consistent probesets
X86810	Homo sapiens EDMD gene /cds=(110,874) /gb=X86810 /gi=	0.075	0.203	2 consistent probesets
U66711	Human Ly-6-related protein (9804) gene, complete cds /	0.075	0.203	2 consistent probesets
AF030409	Homo sapiens sodium-hydrogen exchanger 6 (NHE-6) mRN	0.075	0.205	4 consistent probesets
U79115	HSU79115 Human death adaptor molecule RAIDD (	0.075	0.215	2 consistent probesets
U51336	Human inositol 1,3,4-trisphosphate 5/6-kinase mRNA, co	0.075	0.217	2 consistent probesets
U66669	Homo sapiens 3-hydroxyisobutyryl-coenzyme A hydrolase	0.075	0.224	4 consistent probesets
U61981	HSU61981 Human putative mismatch repair/bindi	0.075	0.237	4 consistent probesets
L35240	Human enigma gene, complete cds /cds=(0,1367) /gb=L352	0.075	0.238	2 consistent probesets
AB018258	Homo sapiens mRNA for KIAA0715 protein, partial cds /c	0.075	0.279	4 consistent probesets
AB011162	Homo sapiens mRNA for KIAA0590 protein, complete cds /	0.075	0.280	2 consistent probesets
D29810	Human mRNA for unknown product, partial cds /cds=(0,10	0.075	0.282	2 consistent probesets
U47926	Human unknown protein B mRNA, complete cds /cds=(88,1	0.075	0.287	2 consistent probesets
AA431822	zw79d02.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-78	0.075	0.294	2 consistent probesets
AB015020	Homo sapiens mRNA for BAP2-beta protein, complete cds	0.075	0.294	2 consistent probesets
D32039	Human pgH3 mRNA for proteoglycan PG-M(V3), complete	0.075	0.294	2 consistent probesets
Z29067	HSNEK3R H.sapiens nek3 mRNA for protein kinas	0.075	0.338	2 consistent probesets
AI557374	PT2.1_6_C10.r Homo sapiens cDNA, 3 end /clone_end=3	0.075	0.369	2 consistent probesets
AF062075	Homo sapiens leupaxin mRNA, complete cds /cds=(93,125	0.075	0.375	2 consistent probesets
AF038960	Homo sapiens SKD1 homolog mRNA, complete cds /cds=(	0.075	0.384	2 consistent probesets
U40622	HSU40622 Human XRCC4 mRNA, complete cds"	0.075	0.384	2 consistent probesets
AI094610	oy64f07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-167	0.075	0.446	2 consistent probesets
U71300	Human snRNA activating protein complex 50kD subunit (S	0.075	0.446	2 consistent probesets
U39318	HSU39318 Human E2 ubiquitin conjugating enzym	0.075	0.076	8 consistent probesets
S71326	BGPc=biliary glycoprotein adhesion molecule {alternati	0.075	0.108	4 inconsistent probesets
AL043108	DKFZp434C0823_r1 Homo sapiens cDNA, 5 end /clone=D	0.075	0.151	4 consistent probesets
U26648	Homo sapiens syntaxin 5 mRNA, complete cds /cds=(26,93	0.075	0.179	4 consistent probesets
U66615	HSU66615 Human SWI/SNF complex 155 KDa subuni	0.073	0.068	8 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AL031228	dJ1033B10.11 (Retinoid X receptor beta (RXRB)) /cds=(1	0.072	0.046	18 inconsistent probesets
M64936	HUMRIRT Homo sapiens retinoic acid-inducible	0.072	0.067	9 consistent probesets
L78833	HUMBRCA1 Human BRCA1, Rho7 and vatI genes, co	0.072	0.093	15 inconsistent probesets
M22995	HUMKREV1A Human ras-related protein (Krev-1)	0.072	0.155	6 consistent probesets
D63874	Human mRNA for HMG-1, complete cds /cds=(76,723) /gb=	0.072	0.167	3 consistent probesets
V00599	HSTUB2 Human mRNA fragment encoding beta-tubu	0.071	0.027	4 inconsistent probesets
X04803	HSYUBG1 Homo sapiens ubiquitin gene	0.071	0.039	4 inconsistent probesets
M12529	HUMAPOE Human apolipoprotein E mRNA, complete	0.071	0.039	4 inconsistent probesets
AF047472	Homo sapiens spleen mitotic checkpoint BUB3 (BUB3) mR	0.071	0.047	4 inconsistent probesets
L39945	Human cytochrome b5 (CYB5) gene /cds=(120,548) /gb=L3	0.071	0.053	8 inconsistent probesets
L05147	HUMDSPHS Human dual specificity phosphatase t	0.071	0.054	8 inconsistent probesets
N47307	yy87a10.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-28	0.071	0.055	4 inconsistent probesets
U49869	Human ubiquitin gene, complete cds /cds=(94,783) /gb=U	0.071	0.060	4 inconsistent probesets
AJ012008	Homo sapiens genes encoding RNCC protein, DDAH protei	0.071	0.061	8 inconsistent probesets
U49278	Homo sapiens UEV-1 (UBE2V) mRNA, partial cds /cds=(0,2	0.071	0.061	4 inconsistent probesets
AF040105	Homo sapiens RCL (Rcl) mRNA, complete cds /cds=(17,54	0.071	0.061	4 consistent probesets
X56468	HS1433 Human mRNA for 14.3.3 protein, a prote	0.071	0.063	8 inconsistent probesets
AA883870	am26f01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-14	0.071	0.064	4 inconsistent probesets
AL034374	Human DNA sequence from clone 483K16 on chromosome	0.071	0.067	4 consistent probesets
L25080	HUMRHOAA Homo sapiens GTP-binding protein (rh	0.071	0.068	4 inconsistent probesets
AB002533	Homo sapiens mRNA for Qip1, complete cds /cds=(9,1574)	0.071	0.070	4 consistent probesets
U76366	Human Treacher Collins syndrome (TCOF1) mRNA, compl	0.071	0.072	4 consistent probesets
L11566	Homo sapiens ribosomal protein L18 (RPL18) mRNA, comp	0.071	0.072	4 consistent probesets
AJ131182	Homo sapiens mRNA for Epsilon COP /cds=(42,968) /gb=A	0.071	0.072	4 consistent probesets
AA402538	zu48g06.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-741	0.071	0.075	4 consistent probesets
AB002362	Human mRNA for KIAA0364 gene, complete cds /cds=(114	0.071	0.078	4 consistent probesets
X68194	H.sapiens h-Sp1 mRNA /cds=(33,812) /gb=X68194 /gi=324	0.071	0.079	4 consistent probesets
U79260	Human clone 23745 mRNA, complete cds /cds=(609,1028)	0.071	0.082	4 inconsistent probesets
X92518	H.sapiens mRNA for HMGI-C protein /cds=UNKNOWN /gb=	0.071	0.082	4 consistent probesets
D31883	Human mRNA for KIAA0059 gene, complete cds /cds=(221	0.071	0.083	4 consistent probesets
U49260	Human mevalonate pyrophosphate decarboxylase (MPD) m	0.071	0.085	4 inconsistent probesets
Z11559	H.sapiens mRNA for iron regulatory factor /cds=(107,27	0.071	0.086	4 consistent probesets
U81375	Human placental equilibrative nucleoside transporter 1	0.071	0.087	4 consistent probesets
M27937	Human male-enhanced antigen mRNA (Mea), complete cds	0.071	0.088	4 consistent probesets
X69804	H.sapiens mRNA for La/SS-B protein /cds=UNKNOWN /gb=	0.071	0.088	4 consistent probesets
D50857	Human DOCK180 protein mRNA, complete cds /cds=(23,56	0.071	0.091	4 consistent probesets
J04046	HUMCAMA Human calmodulin mRNA, complete cds"	0.071	0.091	4 consistent probesets
AI130910	qb81g08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-17	0.071	0.093	4 consistent probesets
AJ010842	Homo sapiens mRNA for putative ATP(GTP)-binding protei	0.071	0.093	4 consistent probesets
X99325	HSSTE20 H.sapiens mRNA for Ste20-like kinase	0.071	0.094	4 consistent probesets
U10362	Human GP36b glycoprotein mRNA, complete cds /cds=(0,1	0.071	0.095	4 consistent probesets
U57877	Human integral membrane protein CII-3 mRNA, nuclear ge	0.071	0.098	4 inconsistent probesets
U59151	Human Cbf5p homolog (CBF5) mRNA, complete cds /cds=	0.071	0.098	4 consistent probesets
X86809	H.sapiens mRNA for major astrocytic phosphoprotein PEA	0.071	0.098	4 consistent probesets
AF029786	Homo sapiens GBAS (GBAS) mRNA, complete cds /cds=(8	0.071	0.098	4 consistent probesets
X59932	HSCSRCKIN Human mRNA for C-SRC-kinase	0.071	0.103	4 consistent probesets
X80695	H.sapiens OXA1Hs mRNA /cds=(6,1313) /gb=X80695 /gi=6	0.071	0.104	4 consistent probesets
X76057	H.sapiens PMI1 mRNA for phosphomannose isomerase /cd	0.071	0.104	4 consistent probesets
AJ005579	Homo sapiens mRNA for Prer protein /cds=(0,1673) /gb=A	0.071	0.107	4 consistent probesets
AL049957	Homo sapiens mRNA; cDNA DKFZp564J0323 (from clone	0.071	0.108	4 consistent probesets
X60708	Human pcHDP7 mRNA for liver dipeptidyl peptidase IV /c	0.071	0.111	4 consistent probesets
U41668	Human deoxyguanosine kinase mRNA, complete cds /cds=	0.071	0.112	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U30888	Human tRNA-guanine transglycosylase mRNA, complete cd	0.071	0.113	4 consistent probesets
AB018290	Homo sapiens mRNA for KIAA0747 protein, partial cds /c	0.071	0.113	4 consistent probesets
M73547	HUMPOLLA Human polyposis locus (DP1 gene) mRN	0.071	0.114	4 consistent probesets
AL080181	Homo sapiens mRNA; cDNA DKFZp434O111 (from clone D	0.071	0.115	4 consistent probesets
AF026030	Homo sapiens putative mitochondrial inner membrane pro	0.071	0.115	4 consistent probesets
N92548	zb29g04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-30	0.071	0.116	4 consistent probesets
W73046	zd54h09.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-344	0.071	0.117	8 inconsistent probesets
U65410	HSU65410 Human Mad2 (hsMAD2) mRNA, complete c	0.071	0.118	4 consistent probesets
X71874	HSPROSCHY H.sapiens genes for proteasome-like	0.071	0.120	4 consistent probesets
AI816724	wj43c06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-240	0.071	0.120	4 consistent probesets
D21852	Human mRNA for KIAA0029 gene, partial cds /cds=(382,32	0.071	0.121	4 consistent probesets
AF082657	Homo sapiens Era GTPase A protein (HERA-A) mRNA, par	0.071	0.122	4 consistent probesets
AJ000186	Homo sapiens mRNA for MAD2 protein /cds=(66,683) /gb=A	0.071	0.127	4 inconsistent probesets
X04434	HSIGFIRR Human mRNA for insulin-like growth f	0.071	0.128	4 inconsistent probesets
X14329	Human mRNA for carboxypeptidase N small subunit (EC 3.	0.071	0.139	4 consistent probesets
W28214	45f7 Homo sapiens cDNA /gb=W28214 /gi=1308297 /	0.071	0.140	4 consistent probesets
AF058953	Homo sapiens ATP-specific succinyl-CoA synthetase beta	0.071	0.141	4 consistent probesets
AI547258	PN001_AH_H08.r Homo sapiens cDNA, 5 end /clone_end=	0.071	0.145	4 inconsistent probesets
AB012130	Homo sapiens SBC2 mRNA for sodium bicarbonate cotrans	0.071	0.150	4 consistent probesets
AB015345	Homo sapiens HRIHFB2216 mRNA, partial cds /cds=(0,146	0.071	0.151	4 consistent probesets
U09196	Human 1.1 kb mRNA upregulated in retinoic acid treated	0.071	0.160	4 consistent probesets
X59960	H.sapiens mRNA for sphingomyelinase /cds=(122,2005) /g	0.071	0.179	4 consistent probesets
AF025441	Homo sapiens Opa-interacting protein OIP5 mRNA, partia	0.071	0.179	4 consistent probesets
M68864	Human ORF mRNA, complete cds /cds=(135,1031) /gb=M6	0.071	0.180	4 consistent probesets
D12620	HUMCYT1 Homo sapiens mRNA for cytochrome P-45	0.071	0.194	4 consistent probesets
AF045229	Homo sapiens regulator of G protein signaling 10 mRNA,	0.071	0.195	4 consistent probesets
U59913	HSU59913 Human chromosome 5 Mad homolog Smad5	0.071	0.204	8 consistent probesets
AL079292	Homo sapiens mRNA full length insert cDNA clone EUROIN	0.071	0.207	4 consistent probesets
U67092	HSU67092 Human ataxia-telangiectasia locus pr	0.071	0.233	4 consistent probesets
U33052	HSU33052 Human lipid-activated, protein kinas	0.069	0.121	7 consistent probesets
AL022723	dJ377H14.1 (major histocompatibility complex, class I,	0.069	0.061	8 consistent probesets
U51903	HSU51903 Human RasGAP-related protein (IQGAP2	0.069	0.096	8 consistent probesets
D14530	HUMRSPT Human homolog of yeast ribosomal prot	0.067	0.039	4 consistent probesets
D31887	Human mRNA for KIAA0062 gene, partial cds /cds=(0,1597	0.067	0.041	4 consistent probesets
U09953	Human ribosomal protein L9 mRNA, complete cds /cds=(29	0.067	0.047	4 inconsistent probesets
AJ002308	Homo sapiens mRNA for synaptogyrin 2 /cds=(29,703) /gb	0.067	0.050	4 consistent probesets
U14966	Human ribosomal protein L5 mRNA, complete cds /cds=(30	0.067	0.050	4 consistent probesets
M81757	H.sapiens S19 ribosomal protein mRNA, complete cds /cd	0.067	0.050	4 consistent probesets
L02426	HUM26SPSIV Human 26S protease (S4) regulatory	0.067	0.050	4 inconsistent probesets
M62895	Human lipocortin (LIP) 2 pseudogene mRNA, complete cds	0.067	0.050	4 inconsistent probesets
U54558	Homo sapiens translation initiation factor eIF3 p66 su	0.067	0.053	4 consistent probesets
AL033377	Human DNA sequence from clone 287G14 on chromosome	0.067	0.055	4 consistent probesets
J02783	HUMTHBP Human thyroid hormone binding protein	0.067	0.055	4 consistent probesets
S79522	ubiquitin carboxyl extension protein [human, mRNA, 540	0.067	0.055	4 consistent probesets
AF069733	Homo sapiens ADA3-like protein mRNA, complete cds /cds	0.067	0.055	4 inconsistent probesets
AF004876	Homo sapiens 54TMp (54tm) mRNA, complete cds /cds=(1	0.067	0.055	4 inconsistent probesets
U09813	Human mitochondrial ATP synthase subunit 9, P3 gene co	0.067	0.056	4 consistent probesets
M58028	HUMUBIQAA Human ubiquitin-activating enzyme E	0.067	0.056	4 consistent probesets
J03242	HUMGFIL2 Human insulin-lke growth factor II m	0.067	0.057	8 inconsistent probesets
D38251	Homo sapiens mRNA for RPB5 (XAP4), complete cds /cds=	0.067	0.058	4 consistent probesets
AI541050	pec1.2-1.E08.r Homo sapiens cDNA, 5 end /clone_end=5	0.067	0.058	4 consistent probesets
AI535946	vicpro2.D07.r Homo sapiens cDNA, 5 end /clone_end=5	0.067	0.058	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AA131149	zo16d05.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-587	0.067	0.059	4 inconsistent probesets
U41303	Human small nuclear ribonuleoprotein particle N (SNRPN	0.067	0.060	4 consistent probesets
U03057	Human actin bundling protein (HSN) mRNA, complete cds	0.067	0.061	4 consistent probesets
M55409	HUMPANCAN Homo sapiens pancreatic tumor-relat	0.067	0.061	4 consistent probesets
AF005046	Homo sapiens serine/threonine kinase mRNA, complete cd	0.067	0.061	4 consistent probesets
X99209	H.sapiens mRNA for arginine methyltransferase /cds=(17	0.067	0.062	4 inconsistent probesets
U30255	Human phosphogluconate dehydrogenase (hPGDH) gene, c	0.067	0.064	4 consistent probesets
D83198	Homo sapiens mRNA expressed in thyroid gland /cds=(341	0.067	0.064	4 consistent probesets
L08666	Homo sapiens porin (por) mRNA, complete cds and trunca	0.067	0.064	4 consistent probesets
J03459	Human leukotriene A-4 hydrolase mRNA, complete cds /cd	0.067	0.064	4 consistent probesets
W16505	zb05e12.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-301	0.067	0.064	4 inconsistent probesets
S62140	TLS=translocated in liposarcoma [human, mRNA, 1824 nt]	0.067	0.064	4 consistent probesets
U90313	HSU90313 Human glutathione-S-transferase homo	0.067	0.065	4 inconsistent probesets
AF026292	Homo sapiens chaperonin containing t-complex polypepti	0.067	0.065	4 consistent probesets
V00567	HSMGLO Human messenger RNA fragment for the b	0.067	0.068	4 consistent probesets
X66975	H.sapiens mRNA for heterogeneous nuclear ribonucleopro	0.067	0.068	4 consistent probesets
AI557064	PT2.1_13_A12.r Homo sapiens cDNA, 3 end /clone_end=3	0.067	0.068	4 inconsistent probesets
Y07969	H.sapiens mRNA for APRIL protein /cds=(230,979) /gb=Y0	0.067	0.068	4 consistent probesets
AI971724	wr07a04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.067	0.070	4 consistent probesets
U22526	Human 2,3-oxidosqualene-lanosterol cyclase mRNA, compl	0.067	0.070	4 consistent probesets
K03000	Human aldehyde dehydrogenase 1 mRNA /cds=(0,1022) /g	0.067	0.070	4 consistent probesets
S82297	S82297 beta 2-microglobulin {11bp deleted bet	0.067	0.070	4 inconsistent probesets
D14696	Human mRNA for KIAA0108 gene, complete cds /cds=(146	0.067	0.070	4 inconsistent probesets
AF038962	Homo sapiens voltage dependent anion channel protein m	0.067	0.072	4 consistent probesets
J03075	Human 80K-H protein (kinase C substrate) mRNA, complet	0.067	0.072	4 consistent probesets
X90872	H.sapiens mRNA for gp25L2 protein /cds=(91,735) /gb=X9	0.067	0.074	4 consistent probesets
D55696	D55696 Homo sapiens mRNA for cysteine proteas	0.067	0.074	4 consistent probesets
J02984	Human insulinoma rig-analog mRNA encoding DNA-binding	0.067	0.074	4 consistent probesets
L23805	HUMCATENIN Human alpha1(E)-catenin mRNA, comp	0.067	0.075	4 consistent probesets
AA418779	zv98d05.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-767	0.067	0.075	4 inconsistent probesets
Z78368	HSZ78368 Homo sapiens cDNA /clone=3.142-(CEPH) /gb=	0.067	0.075	4 consistent probesets
AI357653	qy15c11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-20	0.067	0.075	4 consistent probesets
X16277	Human gene for ornithine decarboxylase ODC (EC 4.1.1.1	0.067	0.076	4 consistent probesets
X74795	HSP1CDC46 H.sapiens P1-Cdc46 mRNA	0.067	0.077	4 consistent probesets
X85116	H.sapiens epb72 gene exon 1 /cds=(61,927) /gb=X85116 /	0.067	0.077	4 consistent probesets
U85611	HSU85611 Human DNA-PK interaction protein (KI	0.067	0.078	4 consistent probesets
L21990	Human spliceosomal protein (SAP 62) gene, complete cds	0.067	0.078	4 inconsistent probesets
J00129	Human fibrinogen beta-chain mRNA, partial cds /cds=UNK	0.067	0.079	4 consistent probesets
AL035252	Human DNA sequence from clone 738P15 on chromosome	0.067	0.080	4 consistent probesets
AA526497	ni96d07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-984	0.067	0.083	4 consistent probesets
U04847	Human Ini1 mRNA, complete cds /cds=(69,1226) /gb=U048	0.067	0.085	4 consistent probesets
L15702	Human complement factor B mRNA, complete cds /cds=(40	0.067	0.085	4 consistent probesets
AF042377	Homo sapiens GDP-mannose 4,6 dehydratase mRNA, com	0.067	0.085	4 inconsistent probesets
W72440	zd65e10.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-345	0.067	0.086	4 consistent probesets
D29011	Human mRNA for proteasome subunit X, complete cds /cds	0.067	0.086	4 consistent probesets
X62654	H.sapiens gene for Me491/CD63 antigen /cds=(69,785) /g	0.067	0.089	4 consistent probesets
AF005050	Homo sapiens aspartyl aminopeptidase mRNA, complete c	0.067	0.089	4 consistent probesets
Z69043	H.sapiens mRNA translocon-associated protein delta sub	0.067	0.091	4 consistent probesets
U73477	Human acidic nuclear phosphoprotein pp32 mRNA, comple	0.067	0.091	4 consistent probesets
S80562	acidic calponin [human, kidney, mRNA, 1607 nt] /cds=(8	0.067	0.093	4 consistent probesets
AI086057	oz44f07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-167	0.067	0.093	4 consistent probesets
U80184	Homo sapiens FLII gene, complete cds /cds=(35,3844) /g	0.067	0.094	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AA203476	zx55e01.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-446	0.067	0.095	4 inconsistent probesets
X12451	Human mRNA for pro-cathepsin L (major excreted protein	0.067	0.096	4 consistent probesets
X59812	HSVD3HYD H.sapiens CYP 27 mRNA for vitamin D3	0.067	0.096	4 consistent probesets
D10522	Homo sapiens mRNA for 80K-L protein, complete cds /cds	0.067	0.098	4 consistent probesets
AL050108	Homo sapiens mRNA; cDNA DKFZp586I2219 (from clone D	0.067	0.098	4 consistent probesets
U50383	Human retinoic acid-responsive protein (NN8-4AG) mRNA,	0.067	0.098	4 consistent probesets
AF030424	Homo sapiens histone acetyltransferase 1 mRNA, complet	0.067	0.102	4 consistent probesets
S61953	S61953 c-erbB3=receptor tyrosine kinase {alte	0.067	0.107	8 inconsistent probesets
L42542	Human RLIP76 protein mRNA, complete cds /cds=(223,219	0.067	0.107	4 consistent probesets
Z98265	Homo sapiens mRNA for plakophilin 3 /cds=(74,2467) /gb	0.067	0.108	3 consistent probesets
X62078	H.sapiens mRNA for GM2 activator protein /cds=UNKNOW	0.067	0.108	4 consistent probesets
D63481	Human mRNA for KIAA0147 gene, partial cds /cds=(0,4657	0.067	0.108	4 consistent probesets
U81504	Homo sapiens beta-3A-adaptin subunit of the AP-3 compl	0.067	0.108	4 consistent probesets
U06631	Human (H326) mRNA, complete cds /cds=(176,1969) /gb=U	0.067	0.108	4 consistent probesets
M28212	HUMRAB6A Homo sapiens GTP-binding protein (RA	0.067	0.110	4 inconsistent probesets
AF042083	Homo sapiens BH3 interacting domain death agonist (BID	0.067	0.113	3 inconsistent probesets
AB011093	Homo sapiens mRNA for KIAA0521 protein, partial cds /c	0.067	0.119	4 consistent probesets
D87432	Human mRNA for KIAA0245 gene, complete cds /cds=(261	0.067	0.122	4 consistent probesets
AI557062	PT2.1_13_A09.r Homo sapiens cDNA, 3 end /clone_end=3	0.067	0.128	4 consistent probesets
AB018334	Homo sapiens mRNA for KIAA0791 protein, complete cds /	0.067	0.129	3 consistent probesets
AB002135	Homo sapiens mRNA for glycosylphosphatidylinositol anc	0.067	0.129	4 consistent probesets
AI557295	PT2.1_16_D02.r Homo sapiens cDNA, 3 end /clone_end=3	0.067	0.136	2 consistent probesets
L13943	Human glycerol kinase (GK) mRNA exons 1-4, complete co	0.067	0.136	4 consistent probesets
L13761	Human dihydrolipoamide dehydrogenase gene /cds=(2077,	0.067	0.137	4 consistent probesets
AB001106	AB001106 Homo sapiens mRNA for glia maturatio	0.067	0.140	4 consistent probesets
Z67743	H.sapiens mRNA for CLC-7 chloride channel protein /cds	0.067	0.141	4 consistent probesets
L37747	Homo sapiens lamin B1 gene /cds=(340,2100) /gb=L37747	0.067	0.143	4 consistent probesets
L11669	Human tetracycline transporter-like protein mRNA, comp	0.067	0.143	4 consistent probesets
AB002318	Human mRNA for KIAA0320 gene, partial cds /cds=(0,2849	0.067	0.146	4 consistent probesets
AF016492	Homo sapiens UDP-glucuronosyltransferase 2B mRNA, cor	0.067	0.147	4 consistent probesets
X78711	H.sapiens mRNA for glycerol kinase testis specific 1 /	0.067	0.148	4 consistent probesets
U75968	Human clone C3 CHL1 protein (CHLR1) mRNA, alternative	0.067	0.150	3 consistent probesets
D86962	Human mRNA for KIAA0207 gene, complete cds /cds=(781	0.067	0.150	4 consistent probesets
R54564	yg81b12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-39	0.067	0.151	3 consistent probesets
X02994	Human mRNA for adenosine deaminase (adenosine aminol	0.067	0.152	4 consistent probesets
L28821	Homo sapiens alpha mannosidase II isozyme mRNA, comp	0.067	0.155	4 consistent probesets
W28281	47e7 Homo sapiens cDNA /gb=W28281 /gi=1308436 /	0.067	0.155	4 inconsistent probesets
AF042081	Homo sapiens SH3 domain binding glutamic acid-rich-lik	0.067	0.155	4 consistent probesets
AL050356	Homo sapiens mRNA; cDNA DKFZp564L2016 (from clone	0.067	0.156	4 consistent probesets
AB007896	Homo sapiens KIAA0436 mRNA, partial cds /cds=(0,2069)	0.067	0.156	4 consistent probesets
U02609	Human transducin-like protein mRNA, complete cds /cds=	0.067	0.158	3 consistent probesets
AF035318	Homo sapiens clone 23705 mRNA sequence /cds=UNKNO	0.067	0.163	3 consistent probesets
X60221	H.sapiens mRNA for H+-ATP synthase subunit b /cds=(32,	0.067	0.168	1 consistent probesets
H05692	yl76b12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-437	0.067	0.172	3 consistent probesets
M29551	Human calcineurin A2 mRNA, complete cds /cds=(116,169	0.067	0.176	3 consistent probesets
AF018081	Homo sapiens type XVIII collagen (COL18A1) mRNA, alter	0.067	0.180	3 consistent probesets
W28595	48h8 Homo sapiens cDNA /gb=W28595 /gi=1308543 /	0.067	0.184	3 consistent probesets
U63329	HSU63329 Human mutY homolog (hMYH) gene, comp	0.067	0.184	4 consistent probesets
U52191	Human SMCY (H-Y) mRNA, complete cds /cds=(275,4894)	0.067	0.186	4 consistent probesets
D86322	Homo sapiens mRNA for calmegin, complete cds /cds=(101	0.067	0.192	4 consistent probesets
AF002210	Homo sapiens copper chaperone for superoxide dismutase	0.067	0.193	3 consistent probesets
S54641	HZF-16=Kruppel-related zinc finger gene homolog {alter	0.067	0.202	1 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AL050078	Homo sapiens mRNA; cDNA DKFZp566G0746 (from clone	0.067	0.205	4 inconsistent probesets
AB014564	Homo sapiens mRNA for KIAA0664 protein, partial cds /c	0.067	0.209	3 consistent probesets
X57522	H.sapiens RING4 cDNA /cds=(30,2456) /gb=X57522 /gi=36	0.067	0.213	3 consistent probesets
W28193	43d12 Homo sapiens cDNA /gb=W28193 /gi=1308141 /	0.067	0.213	3 consistent probesets
AA151716	zo30d07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-58	0.067	0.213	4 consistent probesets
U33838	HSU33838 Human NF-kappa-B p65delta3 mRNA, spl	0.067	0.215	4 consistent probesets
AL050030	Homo sapiens mRNA; cDNA DKFZp566E0124 (from clone	0.067	0.215	4 inconsistent probesets
X96717	H.sapiens mRNA for transcription factor TFE3 /cds=(234	0.067	0.223	3 consistent probesets
U58522	Human huntingtin interacting protein (HIP2) mRNA, comp	0.067	0.232	4 consistent probesets
AJ130733	Homo sapiens mRNA 2-methylacyl-CoA racemase /cds=(88	0.067	0.235	3 consistent probesets
AF094481	Homo sapiens trinucleotide repeat DNA binding protein	0.067	0.237	3 consistent probesets
AF016582	Homo sapiens checkpoint kinase Chk1 (CHK1) mRNA, com	0.067	0.240	3 consistent probesets
M81637	Human grancalcin mRNA, complete cds /cds=(119,772) /gb	0.067	0.240	4 consistent probesets
M92287	HUMCYCD3A Homo sapiens cyclin D3 (CCND3) mRNA	0.067	0.241	3 consistent probesets
AB007917	Homo sapiens mRNA for KIAA0448 protein, complete cds /	0.067	0.247	4 consistent probesets
X55740	Human placental cDNA coding for 5nucleotidase (EC 3.1.	0.067	0.253	3 consistent probesets
S72904	APK1 antigen=MAb KI recognized [human, ovarian carcino	0.067	0.274	4 consistent probesets
AB008226	Homo sapiens FCMD mRNA for fukutin, complete cds /cds=	0.067	0.283	3 consistent probesets
AB011169	Homo sapiens mRNA for KIAA0597 protein, partial cds /c	0.067	0.300	4 consistent probesets
X84195	H.sapiens mRNA for acylphosphatase, muscle type (MT) i	0.067	0.364	3 consistent probesets
U88048	Human clone KiSS-16 unknown product mRNA, complete c	0.067	0.370	1 consistent probesets
AL049449	Homo sapiens mRNA; cDNA DKFZp586B1722 (from clone	0.067	0.383	2 consistent probesets
AI761567	wg66a05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.067	0.497	2 consistent probesets
U37518	HSU37518 Human TNF-related apoptosis inducing	0.067	0.526	3 consistent probesets
U03100	Human alpha2(E)-catenin mRNA, complete cds /cds=(4,279	0.065	0.053	8 inconsistent probesets
AW044624	wy78c04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.065	0.058	8 inconsistent probesets
AA628946	af28f05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-103	0.065	0.085	8 consistent probesets
L13848	HUMRNAHELA Human RNA helicase A mRNA, complet	0.065	0.085	8 consistent probesets
M31767	HUMDNAMET Human O6-methylguanine-DNA methyltr	0.065	0.105	8 consistent probesets
U37139	Human beta 3-endonexin mRNA, long form and short form,	0.064	0.122	7 consistent probesets
X66363	HSSTHPKD H.sapiens mRNA PCTAIRE-1 for serine/	0.064	0.062	12 consistent probesets
AL031432	Human DNA sequence from clone 465N24 on chromosome	0.064	0.124	6 consistent probesets
M64241	HUMQM Human Wilm s tumor-related protein (QM)	0.063	0.039	4 inconsistent probesets
M72709	Human alternative splicing factor mRNA, complete cds /	0.063	0.048	4 inconsistent probesets
J04988	HUMHSP90B Human 90 kD heat shock protein gene	0.063	0.053	4 consistent probesets
Z12962	H.sapiens mRNA for homologue to yeast ribosomal protei	0.063	0.054	4 consistent probesets
M58458	Human ribosomal protein S4 (RPS4X) isoform mRNA, com	0.063	0.055	4 consistent probesets
J00307	human prothrombin (phii-3 clone) mrna /cds=UNKNOWN /g	0.063	0.055	4 consistent probesets
M92383	Homo sapiens thymosin beta-10 gene, 3end /cds=(0,149)	0.063	0.055	4 inconsistent probesets
D38548	Human mRNA for KIAA0076 gene, complete cds /cds=(86,5	0.063	0.058	4 inconsistent probesets
X15187	HSTRA1 Human tra1 mRNA for human homologue of	0.063	0.061	4 inconsistent probesets
AB002363	Human mRNA for KIAA0365 gene, partial cds /cds=(0,2819	0.063	0.064	4 consistent probesets
Z48199	H.sapiens syndecan-1 gene (exons 2-5) /cds=(0,866) /gb	0.063	0.064	4 inconsistent probesets
AF016903	Homo sapiens agrin precursor mRNA, partial cds /cds=(0	0.063	0.064	4 consistent probesets
D32053	Homo sapiens mRNA for Lysyl tRNA Synthetase, complete	0.063	0.067	4 consistent probesets
AL041780	DKFZp434A0418_s1 Homo sapiens cDNA, 3 end /clone=D	0.063	0.068	4 consistent probesets
AA976838	oq35c12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-15	0.063	0.068	4 consistent probesets
S78187	S78187 CDC25Hu2=cdc25+ homolog [human, mRNA,	0.063	0.070	4 inconsistent probesets
M13560	Human Ia-associated invariant gamma-chain gene /cds=(7	0.063	0.070	4 consistent probesets
M84739	Human autoantigen calreticulin mRNA, complete cds /cds	0.063	0.070	4 consistent probesets
Z82215	Homo sapiens DNA sequence from PAC 68O2 on chromos	0.063	0.072	4 consistent probesets
W68046	zd42a12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-34	0.063	0.075	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U51007	Human 26S protease subunit S5a mRNA, complete cds /cd	0.063	0.076	4 consistent probesets
X83573	Homo sapiens ARSE gene, complete CDS /cds=(67,1836) /	0.063	0.076	4 consistent probesets
X99584	H.sapiens mRNA for SMT3A protein /cds=(94,405) /gb=X99	0.063	0.077	4 consistent probesets
Z78324	HSZ78324 Homo sapiens cDNA /clone=2.45-(CEPH) /gb=Z	0.063	0.080	4 consistent probesets
U21858	HSU21858 Human transcriptional activation fac	0.063	0.082	4 inconsistent probesets
AL050367	Homo sapiens mRNA; cDNA DKFZp564A026 (from clone D	0.063	0.084	4 consistent probesets
AI540957	PEC1.2_15_G03.r Homo sapiens cDNA, 5 end /clone_end=	0.063	0.085	4 consistent probesets
D11139	HUMTIMP Human gene for tissue inhibitor of me	0.063	0.085	4 inconsistent probesets
L48516	Homo sapiens paraoxonase 3 (PON3) mRNA, 3 end of cds	0.063	0.085	4 inconsistent probesets
M76231	Human sepiapterin reductase mRNA, complete cds /cds=(2	0.063	0.089	4 consistent probesets
U13991	HSU13991 Human TATA-binding protein associate	0.063	0.089	4 consistent probesets
L06419	Homo sapiens lysyl hydroxylase (PLOD) mRNA, complete c	0.063	0.091	4 consistent probesets
U29091	Human selenium-binding protein (hSBP) mRNA, complete c	0.063	0.091	4 consistent probesets
X14813	Human liver mRNA for 3-oxoacyl-CoA thiolase /cds=(36,1	0.063	0.093	4 consistent probesets
M68516	Human protein C inhibitor gene, complete cds /cds=(115	0.063	0.093	4 consistent probesets
U51586	Human siah binding protein 1 (SiahBP1) mRNA, partial c	0.063	0.093	4 consistent probesets
M36340	Human ADP-ribosylation factor 1 (ARF1) mRNA, complete	0.063	0.096	4 consistent probesets
Y09008	H.sapiens mRNA for uracil-DNA glycosylase /cds=(70,101	0.063	0.096	4 consistent probesets
AF003837	Homo sapiens Jagged1 (JAG1) mRNA, complete cds /cds=	0.063	0.098	4 consistent probesets
AF034091	Homo sapiens nuclear localization signal containing pr	0.063	0.098	4 consistent probesets
Y00097	Human mRNA for protein p68 /cds=(100,2121) /gb=Y00097	0.063	0.102	4 inconsistent probesets
M95178	Human non-muscle alpha-actinin mRNA, complete cds /cds	0.063	0.102	4 inconsistent probesets
AA522891	ni41b12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-979	0.063	0.102	4 consistent probesets
AI991040	wu36b05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-99	0.063	0.102	4 consistent probesets
U26266	Human deoxyhypusine synthase mRNA, complete cds /cds=	0.063	0.114	4 consistent probesets
W29030	55c4 Homo sapiens cDNA /gb=W29030 /gi=1308987 /	0.063	0.116	4 inconsistent probesets
AL080164	Homo sapiens mRNA; cDNA DKFZp564C1940 (from clone	0.063	0.116	4 consistent probesets
Y07604	H.sapiens mRNA for nucleoside-diphosphate kinase /cds=	0.063	0.116	4 consistent probesets
U39412	Homo sapiens alpha SNAP mRNA, complete cds /cds=(67,	0.063	0.118	4 consistent probesets
AF038952	Homo sapiens cofactor A protein mRNA, complete cds /cd	0.063	0.127	4 consistent probesets
AF004563	Homo sapiens hUNC18b alternatively-spliced mRNA, comp	0.063	0.139	4 consistent probesets
Y13834	Homo sapiens mRNA for farnesylated-proteins converting	0.063	0.144	4 consistent probesets
X92396	H.sapiens mRNA for novel gene in Xq28 region /cds=(114	0.063	0.144	4 consistent probesets
AL049924	Homo sapiens mRNA; cDNA DKFZp547G1110 (from clone	0.063	0.154	4 consistent probesets
U42412	Human 5-AMP-activated protein kinase, gamma-1 subunit	0.063	0.154	4 consistent probesets
AL050162	Homo sapiens mRNA; cDNA DKFZp586B2022 (from clone	0.063	0.155	4 consistent probesets
AB006190	Homo sapiens mRNA for aquaporin adipose, complete cds	0.063	0.169	4 consistent probesets
AF065485	Homo sapiens sorting nexin 4 mRNA, complete cds /cds=(	0.063	0.181	4 consistent probesets
U22815	Human LAR-interacting protein 1a mRNA, complete cds /c	0.063	0.281	4 consistent probesets
Y09443	H.sapiens mRNA for alkyl-dihydroxyacetonephosphate syn	0.062	0.086	4 inconsistent probesets
U72508	Human B7 mRNA, complete cds /cds=(112,1050) /gb=U725	0.061	0.229	3 consistent probesets
X04409	Human mRNA for coupling protein G(s) alpha-subunit (al	0.060	0.030	8 inconsistent probesets
W52024	zd13a03.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-34	0.060	0.050	8 inconsistent probesets
M13932	Human ribosomal protein S17 mRNA, complete cds /cds=(2	0.058	0.043	8 consistent probesets
D55654	Human mRNA for cytosolic malate dehydrogenase, comple	0.058	0.044	4 inconsistent probesets
M22806	Human prolyl 4-hydroxylase beta-subunit and disulfide	0.058	0.044	4 consistent probesets
M30938	HUMKUP Human Ku (p70/p80) subunit mRNA, compl	0.058	0.045	12 consistent probesets
U94855	Homo sapiens translation initiation factor 3 47 kDa su	0.058	0.047	4 consistent probesets
Z26876	H.sapiens gene for ribosomal protein L38 /cds=(110,322	0.058	0.049	4 inconsistent probesets
D00596	HUMTS1 Homo sapiens gene for thymidylate synt	0.058	0.050	4 consistent probesets
AI525834	PT1.3_06_D01.r Homo sapiens cDNA, 5 end /clone_end=5	0.058	0.053	4 consistent probesets
J03191	Human profilin mRNA, complete cds /cds=(127,549) /gb=J	0.058	0.055	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
X07732	Human hepatoma mRNA for serine protease hepsin /cds=U	0.058	0.057	4 consistent probesets
M94314	Homo sapiens ribosomal protein L30 mRNA, complete cds	0.058	0.057	4 inconsistent probesets
M28713	Homo sapiens NADH-cytochrome b5 reductase (b5R) gene	0.058	0.058	4 consistent probesets
U14968	Human ribosomal protein L27a mRNA, complete cds /cds=(	0.058	0.058	4 consistent probesets
M18667	Human pepsinogen C gene /cds=(73,1230) /gb=M18667 /gi	0.058	0.061	4 consistent probesets
X64330	H.sapiens mRNA for ATP-citrate lyase /cds=(84,3401) /g	0.058	0.062	4 consistent probesets
X06617	Human mRNA for ribosomal protein S11 /cds=(15,491) /gb	0.058	0.062	4 consistent probesets
D13630	Human mRNA for KIAA0005 gene, complete cds /cds=(80,1	0.058	0.064	4 consistent probesets
Z37986	H.sapiens mRNA for phenylalkylamine binding protein /c	0.058	0.064	4 consistent probesets
U17999	HSU17999 Homo sapiens cDNA /clone=B49B32B27 /gb=U1	0.058	0.064	4 consistent probesets
X74104	H.sapiens mRNA for TRAP beta subunit /cds=(50,601) /gb	0.058	0.064	4 consistent probesets
M64716	Human ribosomal protein S25 mRNA, complete cds /cds=(7	0.058	0.067	4 consistent probesets
X74801	H.sapiens Cctg mRNA for chaperonin /cds=(0,1634) /gb=X	0.058	0.068	4 consistent probesets
Y13492	Homo sapiens mRNA for smoothelin-B /cds=(219,2972) /gb	0.058	0.068	4 inconsistent probesets
M88458	Human ELP-1 mRNA sequence /cds=UNKNOWN /gb=M88	0.058	0.070	4 consistent probesets
D17793	Human mRNA for KIAA0119 gene, complete cds /cds=(51,1	0.058	0.070	4 consistent probesets
U33821	HSU33821 Homo sapiens tax1-binding protein TX	0.058	0.071	8 consistent probesets
Z21507	H.sapiens EF-1delta gene encoding human elongation fac	0.058	0.072	4 consistent probesets
AL041443	DKFZp434D0717_s1 Homo sapiens cDNA, 3 end /clone=D	0.058	0.073	4 consistent probesets
D14658	Human mRNA for KIAA0102 gene, complete cds /cds=(307	0.058	0.075	4 consistent probesets
L16842	HUMMITCORA Human ubiquinol cytochrome-c reduc	0.058	0.075	4 consistent probesets
L07633	Homo sapiens (clone 1950.2) interferon-gamma IEF SSP 5	0.058	0.076	4 consistent probesets
U22961	Human mRNA clone with similarity to L-glycerol-3-phosp	0.058	0.077	8 inconsistent probesets
U52100	Human XMP mRNA, complete cds /cds=(63,566) /gb=U521	0.058	0.077	4 consistent probesets
X82456	H.sapiens MLN50 mRNA /cds=(75,860) /gb=X82456 /gi=24	0.058	0.078	4 consistent probesets
AF007142	Homo sapiens clone 23938 mRNA sequence /cds=UNKNO	0.058	0.080	4 consistent probesets
AA631698	np79a08.s1 Homo sapiens cDNA /clone=IMAGE-1132502 /	0.058	0.080	4 consistent probesets
AI828168	wk32h09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.058	0.081	4 consistent probesets
D23661	Human mRNA for ribosomal protein L37, complete cds /cd	0.058	0.082	4 consistent probesets
U72355	Human Hsp27 ERE-TATA-binding protein (HET) mRNA, co	0.058	0.082	4 inconsistent probesets
AI708889	as86g01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.058	0.082	4 consistent probesets
Z25749	H.sapiens gene for ribosomal protein S7 /cds=(81,665)	0.058	0.083	4 consistent probesets
X60489	Human mRNA for elongation factor-1-beta /cds=(235,912)	0.058	0.083	4 consistent probesets
J04444	HUMCYC1A Human cytochrome c-1 gene, complete	0.058	0.085	4 consistent probesets
D28137	Human mRNA for BST-2, complete cds /cds=(9,551) /gb=D	0.058	0.085	4 consistent probesets
AF015553	Homo sapiens TFII-I protein (TFII-I) mRNA, complete cd	0.058	0.085	4 consistent probesets
AF042384	Homo sapiens BC-2 protein mRNA, complete cds /cds=(129	0.058	0.085	4 consistent probesets
AF010400	untitled /cds=(50,1063) /gb=AF010400 /gi=2612878 /	0.058	0.086	4 consistent probesets
AI345944	qp47e09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-19	0.058	0.086	4 inconsistent probesets
AI541336	pec1.2-7.A07.r Homo sapiens cDNA, 5 end /clone_end=5	0.058	0.086	4 consistent probesets
D21235	Human mRNA for HHR23A protein, complete cds /cds=(36,	0.058	0.088	4 consistent probesets
AF070649	Homo sapiens clone 24452 mRNA sequence /cds=UNKNO	0.058	0.090	4 consistent probesets
L42572	Homo sapiens p87/89 gene, complete cds /cds=(92,2368)	0.058	0.091	4 consistent probesets
M33308	Human vinculin mRNA, complete cds /cds=(50,3250) /gb=M	0.058	0.091	4 consistent probesets
X92098	H.sapiens mRNA for transmembrane protein rnp24 /cds=(2	0.058	0.091	4 consistent probesets
D83077	Homo sapiens mRNA for TPRD, complete cds /cds=(1396,7	0.058	0.092	4 consistent probesets
M77836	Human pyrroline 5-carboxylate reductase mRNA, complete	0.058	0.092	4 consistent probesets
H93123	yv05g07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.058	0.092	4 consistent probesets
U28386	Human nuclear localization sequence receptor hSRP1alph	0.058	0.093	4 consistent probesets
X76040	H.sapiens mRNA for Lon protease-like protein /cds=(61,	0.058	0.095	4 consistent probesets
AL050101	Homo sapiens mRNA; cDNA DKFZp586E1519 (from clone	0.058	0.095	4 consistent probesets
Y00486	Human APRT gene for adenine phosphoribosyltransferase	0.058	0.098	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
W27641	37d11 Homo sapiens cDNA /gb=W27641 /gi=1307715 /	0.058	0.100	4 consistent probesets
AL021937	dJ149A16.5 (PUTATIVE novel protein similar to mouse RG	0.058	0.101	4 consistent probesets
AB011114	Homo sapiens mRNA for KIAA0542 protein, complete cds /	0.058	0.101	6 inconsistent probesets
AI541308	pec1.2-4.F11.r Homo sapiens cDNA, 5 end /clone_end=5	0.058	0.102	4 consistent probesets
L29433	Human factor X (blood coagulation factor) gene /cds=(2	0.058	0.104	4 consistent probesets
AL080101	Homo sapiens mRNA; cDNA DKFZp564L0472 (from clone	0.058	0.107	4 consistent probesets
AF070640	Homo sapiens clone 24781 mRNA sequence /cds=UNKNO	0.058	0.107	4 consistent probesets
U41816	Human C-1 mRNA, complete cds /cds=(11,403) /gb=U4181	0.058	0.111	4 consistent probesets
L19605	Homo sapiens 56K autoantigen annexin XI gene mRNA, co	0.058	0.111	4 consistent probesets
U53003	Human GT335 mRNA, complete cds /cds=(84,890) /gb=U5	0.058	0.112	4 consistent probesets
AF027974	Homo sapiens clone LM1955 H105e3 gene, partial cds /cd	0.058	0.113	4 consistent probesets
U65090	Human carboxypeptidase D mRNA, complete cds /cds=(35	0.058	0.115	4 consistent probesets
U91512	Human adhesion molecule ninjurin mRNA, complete cds /c	0.058	0.118	4 consistent probesets
AL096779	Novel human gene mapping to chomosome 2213.3 similar	0.058	0.118	4 consistent probesets
X16135	Human mRNA for novel heterogeneous nuclear RNP protei	0.058	0.121	4 consistent probesets
L07765	Human carboxylesterase mRNA, complete cds /cds=(67,17	0.058	0.124	4 consistent probesets
AB024705	Homo sapiens mRNA for fls485, complete cds /cds=(246,1	0.058	0.124	4 consistent probesets
AB002336	Human mRNA for KIAA0338 gene, partial cds /cds=(0,2806	0.058	0.128	4 consistent probesets
M95623	Homo sapiens hydroxymethylbilane synthase gene, comple	0.058	0.132	4 consistent probesets
AB006572	Homo sapiens RMP mRNA for RPB5 meidating protein, co	0.058	0.132	4 consistent probesets
S72370	pyruvate carboxylase [human, kidney, mRNA, 4017 nt] /c	0.058	0.134	4 inconsistent probesets
AL046394	DKFZp434M217_r1 Homo sapiens cDNA, 5 end /clone=DK	0.058	0.140	4 inconsistent probesets
AF000416	Homo sapiens EXT-like protein 2 (EXTL2) mRNA, complete	0.058	0.141	4 consistent probesets
AF088219	Homo sapiens CC chemokine gene cluster, complete seque	0.058	0.152	4 consistent probesets
W28575	51f12 Homo sapiens cDNA /gb=W28575 /gi=1308730 /	0.058	0.154	4 consistent probesets
U51587	Homo sapiens Golgi complex autoantigen golgin-97 mRNA	0.058	0.167	4 consistent probesets
AF065854	Homo sapiens OR7E12P pseudogene, complete sequence	0.058	0.170	4 consistent probesets
AL080111	Homo sapiens mRNA; cDNA DKFZp586G2222 (from clone	0.058	0.172	4 consistent probesets
AJ224442	Homo sapiens mRNA for putative methyltransferase /cds=	0.058	0.174	4 consistent probesets
M20560	Human lipocortin-III mRNA, complete cds /cds=(46,1017)	0.058	0.176	4 consistent probesets
AF029729	Homo sapiens neuralized mRNA, complete cds /cds=(61,17	0.058	0.192	2 consistent probesets
AF039081	Homo sapiens Cre binding protein-like 2 mRNA, complete	0.058	0.196	4 consistent probesets
U45973	Human phosphatidylinositol (4,5)bisphosphate 5-phospha	0.058	0.201	4 consistent probesets
X52142	Human mRNA for CTP synthetase (EC 6.3.4.2) /cds=(75,18	0.058	0.207	4 consistent probesets
Z35402	HSECAD3 H.sapiens gene encoding E-cadherin, e	0.058	0.223	4 consistent probesets
U81992	Homo sapiens C2H2 zinc finger protein PLAGL1 (PLAGL1)	0.058	0.247	4 consistent probesets
X15875	Human mRNA for cAMP response element (CRE-BP1) binc	0.058	0.249	4 consistent probesets
AB006626	Homo sapiens mRNA for KIAA0288 gene, complete cds /co	0.058	0.273	4 consistent probesets
AA418537	zv93a09.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-76	0.058	0.287	2 consistent probesets
U83410	Human CUL-2 (cul-2) mRNA, complete cds /cds=(146,2383	0.058	0.424	2 consistent probesets
L41498	Homo sapiens longation factor 1-alpha 1 (PTI-1) mRNA,	0.056	0.033	8 inconsistent probesets
D88674	D88674 Homo sapiens mRNA for antizyme inhibit	0.056	0.087	8 inconsistent probesets
AA099265	zk84f07.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-489	0.056	0.162	8 consistent probesets
X96924	H.sapiens gene encoding mitochondrial citrate transpor	0.054	0.035	8 inconsistent probesets
AI557852	P6test.G05.r Homo sapiens cDNA, 5 end /clone_end=5 /	0.054	0.039	4 inconsistent probesets
J04755	Human ferritin H processed pseudogene, complete cds /c	0.054	0.041	4 consistent probesets
U14972	Human ribosomal protein S10 mRNA, complete cds /cds=(1	0.054	0.045	4 consistent probesets
L06499	Homo sapiens ribosomal protein L37a (RPL37A) mRNA, co	0.054	0.045	4 consistent probesets
X57958	H.sapiens mRNA for ribosomal protein L7 /cds=(22,783)	0.054	0.047	4 consistent probesets
M84711	HUMFTE1A Human v-fos transformation effector	0.054	0.048	4 inconsistent probesets
X65923	H.sapiens fau mRNA /cds=(56,457) /gb=X65923 /gi=31302	0.054	0.048	4 consistent probesets
M13934	Human ribosomal protein S14 gene, complete cds /cds=(0	0.054	0.050	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF054183	Homo sapiens GTP binding protein mRNA, complete cds /c	0.054	0.050	4 inconsistent probesets
U41635	Human OS-9 precurosor mRNA, complete cds /cds=(85,208	0.054	0.050	4 inconsistent probesets
X04106	Human mRNA for calcium dependent protease (small subu	0.054	0.052	4 consistent probesets
AF017115	Homo sapiens cytochrome c oxidase subunit IV precursor	0.054	0.053	4 consistent probesets
X63432	H.sapiens ACTB mRNA for mutant beta-actin (beta-actin)	0.054	0.055	4 consistent probesets
D63475	Human mRNA for KIAA0109 gene, complete cds /cds=(86,1	0.054	0.055	4 consistent probesets
AA083129	zn31a06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-54	0.054	0.057	4 inconsistent probesets
L05095	Homo sapiens ribosomal protein L30 mRNA, complete cds	0.054	0.057	4 consistent probesets
M33764	HUMSODB Human ornithine decarboxylase gene, c	0.054	0.058	4 consistent probesets
U73824	Human p97 mRNA, complete cds /cds=(306,3029) /gb=U73	0.054	0.060	4 consistent probesets
K00558	human alpha-tubulin mRNA, complete cds /cds=(67,1422)	0.054	0.060	4 inconsistent probesets
X04714	Human mRNA for apolipoprotein B-100 (apoB-100) /cds=(1	0.054	0.061	4 inconsistent probesets
X07979	Human mRNA for integrin beta 1 subunit /cds=(103,2499)	0.054	0.061	4 consistent probesets
AI494623	qz17b06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-20	0.054	0.061	4 consistent probesets
X79234	H.sapiens mRNA for ribosomal protein L11 /cds=(0,536)	0.054	0.062	4 consistent probesets
AA426364	zv61b06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-75	0.054	0.062	8 consistent probesets
X02317	Human mRNA for Cu/Zn superoxide dismutase (SOD) /cds=	0.054	0.063	4 inconsistent probesets
J02621	HUMHMG14 Human non-histone chromosomal protei	0.054	0.064	4 consistent probesets
X55284	Human H2 mRNA coding for an asialoglycoprotein recepto	0.054	0.064	4 consistent probesets
X03168	Human mRNA for S-protein /cds=(61,1497) /gb=X03168 /gi	0.054	0.067	4 consistent probesets
X04412	Human mRNA for plasma gelsolin /cds=(14,2362) /gb=X044	0.054	0.067	4 consistent probesets
M76180	Human aromatic amino acid decarboxylase (ddc) mRNA, c	0.054	0.068	4 inconsistent probesets
M74491	Human ADP-ribosylation factor 3 mRNA, complete cds /cd	0.054	0.068	4 consistent probesets
AA044823	zk72a10.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-48	0.054	0.070	4 consistent probesets
M16279	Human MIC2 mRNA, complete cds /cds=(177,734) /gb=M1	0.054	0.070	4 consistent probesets
U20982	Human insulin-like growth factor binding protein-4 (IG	0.054	0.070	4 consistent probesets
AL008726	dJ337O18.2 (Lysosomal Protective Protein precursor (EC	0.054	0.070	4 consistent probesets
AJ224875	Homo sapiens mRNA for putative glucosyltransferase, pa	0.054	0.072	4 inconsistent probesets
U37022	HSU37022 Human cyclin-dependent kinase 4 (CDK	0.054	0.072	4 consistent probesets
W26628	34a4 Homo sapiens cDNA /gb=W26628 /gi=1307471 /	0.054	0.075	4 consistent probesets
W28186	43c2 Homo sapiens cDNA /gb=W28186 /gi=1308134 /	0.054	0.075	4 consistent probesets
AF047436	Homo sapiens F1Fo-ATPase synthase f subunit mRNA, cor	0.054	0.077	4 consistent probesets
L77567	Homo sapiens mitochondrial citrate transport protein (	0.054	0.078	4 consistent probesets
AI743134	wg87f07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.054	0.078	4 consistent probesets
D16431	Human mRNA for hepatoma-derived growth factor, complet	0.054	0.080	4 consistent probesets
X81372	H.sapiens mRNA for biphenyl hydrolase-related protein	0.054	0.080	4 consistent probesets
U96078	Homo sapiens hyaluronoglucosaminidase 1 (HYAL1) mRNA	0.054	0.080	4 consistent probesets
D49400	Homo sapiens mRNA for vacuolar ATPase, complete cds /c	0.054	0.080	4 consistent probesets
M88108	Human p62 mRNA, complete cds /cds=(106,1437) /gb=M88	0.054	0.082	4 consistent probesets
Z48042	H.sapiens mRNA encoding GPI-anchored protein p137 /cds	0.054	0.082	4 inconsistent probesets
Z97630	Human DNA sequence from clone 466N1 on chromosome 2	0.054	0.083	8 inconsistent probesets
U90549	Human non-histone chromosomal protein (NHC) mRNA, co	0.054	0.084	4 inconsistent probesets
M90683	Human lymphocyte antigen (HLA-G1) mRNA, complete cds	0.054	0.085	4 consistent probesets
L03411	Human RD protein (RD) mRNA, complete cds /cds=(86,122	0.054	0.087	4 consistent probesets
L06328	Human voltage-dependent anion channel isoform 2 (VDAC)	0.054	0.087	4 consistent probesets
Y12711	H.sapiens mRNA for putative progesterone binding prote	0.054	0.088	4 consistent probesets
M86737	Human high mobility group box (SSRP1) mRNA, complete	0.054	0.095	4 consistent probesets
AF090988	Homo sapiens U5 snRNP-specific 40 kDa protein mRNA, c	0.054	0.095	4 inconsistent probesets
AF042379	Homo sapiens spindle pole body protein spc97 homolog G	0.054	0.095	4 inconsistent probesets
X67055	H.sapiens mRNA for inter-alpha-trypsin inhibitor heavy	0.054	0.096	4 consistent probesets
L36719	HUMMKK3A Homo sapiens MAP kinase kinase 3 (MK	0.054	0.096	4 consistent probesets
AI765280	wi73a08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.054	0.098	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
M80469	Human MHC class I HLA-J gene, exons 1-8 and complete c	0.054	0.098	4 consistent probesets
Y18046	Homo sapiens mRNA for FOP (FGFR1 oncogene partner) /	0.054	0.098	4 consistent probesets
AL049703	Human gene from PAC 179D3, chromosome X, isoform of	0.054	0.098	4 consistent probesets
M10058	Human asialoglycoprotein receptor H1 mRNA, complete cd	0.054	0.102	4 consistent probesets
L40027	HUMGLYSYN Homo sapiens glycogen synthase kina	0.054	0.102	4 inconsistent probesets
L38961	Human putative transmembrane protein precursor (B5) mR	0.054	0.102	4 consistent probesets
AL050156	Homo sapiens mRNA; cDNA DKFZp586N1020 (from clone	0.054	0.105	4 consistent probesets
Z50853	H.sapiens mRNA for CLPP /cds=(19,852) /gb=Z50853 /gi=9	0.054	0.107	4 consistent probesets
AF019767	Homo sapiens zinc finger protein (ZPR1) mRNA, complete	0.054	0.108	4 consistent probesets
S68616	Na+/H+ exchanger NHE-1 isoform [human, heart, mRNA, 4	0.054	0.111	4 consistent probesets
X76302	H.sapiens RY-1 mRNA for putative nucleic acid binding	0.054	0.115	4 inconsistent probesets
D21262	Human mRNA for KIAA0035 gene, partial cds /cds=(0,2125	0.054	0.117	4 consistent probesets
L26260	Human MHC class III HLA-RP gene, complete cds /cds=(12	0.054	0.120	4 consistent probesets
AF106941	Homo sapiens beta-arrestin 2 mRNA, complete cds /cds=(	0.054	0.121	4 consistent probesets
AB002370	Human mRNA for KIAA0372 gene, complete cds /cds=(297	0.054	0.122	4 inconsistent probesets
AF070626	Homo sapiens clone 24483 unknown mRNA, parital cds /cd	0.054	0.125	4 consistent probesets
AI651368	wb05d07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.054	0.127	4 consistent probesets
M34668	HUMPTPAAA Human protein tyrosine phosphatase	0.054	0.128	4 consistent probesets
AF045584	Homo sapiens PB39 mRNA, complete cds /cds=(76,1755) /	0.054	0.129	4 consistent probesets
AB030506	Homo sapiens mRNA for B9, complete cds /cds=(158,772)	0.054	0.132	4 consistent probesets
Y00264	Human mRNA for amyloid A4 precursor of Alzheimers dise	0.054	0.132	4 consistent probesets
X83425	H.sapiens LU gene for Lutheran blood group glycoprotei	0.054	0.133	4 consistent probesets
U69190	U69190 Homo sapiens cDNA /clone=27655 /gb=U69190 /g	0.054	0.133	4 consistent probesets
U90552	Human butyrophilin (BTF5) mRNA, complete cds /cds=(359	0.054	0.137	4 consistent probesets
M31452	Human proline-rich protein (PRP) mRNA, complete cds /c	0.054	0.138	4 consistent probesets
AL080091	Homo sapiens mRNA; cDNA DKFZp564L0862 (from clone	0.054	0.148	4 consistent probesets
AF012023	Homo sapiens integrin cytoplasmic domain associated pr	0.054	0.158	4 consistent probesets
S87759	S87759 protein phosphatase 2C alpha [human, t	0.054	0.167	8 inconsistent probesets
AL120687	DKFZp762F2110_r1 Homo sapiens cDNA, 5 end /clone=D	0.054	0.173	4 consistent probesets
D86326	Homo sapiens mRNA for p115, complete cds /cds=(175,306	0.054	0.174	4 consistent probesets
AB006780	Homo sapiens mRNA for galectin-3, complete cds /cds=(5	0.054	0.176	4 consistent probesets
AI766078	wh67g08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.054	0.180	4 consistent probesets
X89887	Homo sapiens mRNA for WD repeat protein (HIRA) /cds=(2	0.054	0.185	4 consistent probesets
AF057297	Homo sapiens ornithine decarboxylase antizyme 2 (OAZ2)	0.054	0.209	4 consistent probesets
X90780	Homo sapiens TNNI3 gene /cds=(143,775) /gb=X90780 /gi=	0.054	0.303	4 consistent probesets
U50196	HSU50196 Human adenosine kinase mRNA, complet	0.052	0.152	7 consistent probesets
D00763	HUMPSC9 Human mRNA for proteasome subunit HC9	0.052	0.056	8 inconsistent probesets
U24266	Human pyrroline-5-carboxylate dehydrogenase (P5CDh) mI	0.052	0.057	8 consistent probesets
AF043906	Homo sapiens T245 protein (T245) mRNA, complete cds /c	0.052	0.100	8 consistent probesets
J04617	HUMEF1A Human elongation factor EF-1-alpha ge	0.050	0.031	4 inconsistent probesets
X66945	HSNSAMTK H.sapiens N-sam mRNA for fibroblast	0.050	0.058	4 consistent probesets
M22382	Human mitochondrial matrix protein P1 (nuclear encoded	0.050	0.058	4 inconsistent probesets
AF029890	Homo sapiens hepatitis B virus X interacting protein (	0.050	0.061	4 inconsistent probesets
D38048	D38048 Human mRNA for proteasome subunit z, c	0.050	0.065	8 consistent probesets
X74008	H.sapiens mRNA for protein phosphatase 1 gamma /cds=(1	0.050	0.067	4 consistent probesets
X53793	H.sapiens ADE2H1 mRNA showing homologies to SAICAR	0.050	0.067	4 consistent probesets
X70218	HSPPX Homo sapiens mRNA for protein phosphata	0.050	0.068	4 inconsistent probesets
X94754	H.sapiens mRNA for yeast methionyl-tRNA synthetase hom	0.050	0.068	4 consistent probesets
J04794	Human aldehyde reductase mRNA, complete cds /cds=(60,	0.050	0.068	4 consistent probesets
AI688098	wc92f08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.050	0.070	6 inconsistent probesets
U31930	Human deoxyuridine nucleotidohydrolase mRNA, complete	0.050	0.070	4 consistent probesets
U29185	Homo sapiens prion protein (PrP) gene, complete cds /c	0.050	0.070	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
M10905	Human cellular fibronectin mRNA /cds=(0,2383) /gb=M109	0.050	0.071	4 inconsistent probesets
X51521	Human mRNA for ezrin /cds=(117,1877) /gb=X51521 /gi=31	0.050	0.071	4 consistent probesets
AF104398	Homo sapiens cornichon mRNA, complete cds /cds=(56,49(	0.050	0.076	4 consistent probesets
R87876	yo45h01.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-180	0.050	0.077	4 inconsistent probesets
L19183	Human MAC30 mRNA, 3 end /cds=(0,569) /gb=L19183 /gi=	0.050	0.079	4 consistent probesets
M82809	Human annexin IV (ANX4) mRNA, complete cds /cds=(73,1	0.050	0.079	4 consistent probesets
M32304	HUMMET Human metalloproteinase inhibitor mRNA	0.050	0.080	4 consistent probesets
M93036	HUMGA7A08 Human (clone 21726) carcinoma-assoc	0.050	0.081	4 consistent probesets
AB019392	Homo sapiens mRNA of muscle specific gene M9, complete	0.050	0.082	4 consistent probesets
Z23064	H.sapiens mRNA gene for hnRNP G protein /cds=(11,1186)	0.050	0.082	4 consistent probesets
L76200	HUMGUK1R Human guanylate kinase (GUK1) mRNA,	0.050	0.084	4 consistent probesets
AL031230	dJ73M23.2 (NAD+-dependent succinic semialdehyde dehyd	0.050	0.085	4 consistent probesets
M24069	Human DNA-binding protein A (dbpA) gene, 3 end /cds=(	0.050	0.085	4 inconsistent probesets
D49396	Human mRNA for Apo1_Human (MER5(Aop1-Mouse)-like	0.050	0.085	4 inconsistent probesets
AL022315	dJ1177I5.3 (Lectin, Galactose-binding, soluble, 2 (Gal	0.050	0.085	3 inconsistent probesets
Z56281	HSIRF3MR H.sapiens mRNA for interferon regula	0.050	0.087	3 consistent probesets
AF035316	Homo sapiens clone 23678 mRNA, partial cds /cds=(0,103	0.050	0.087	4 consistent probesets
AF006083	Homo sapiens actin-related protein Arp3 (ARP3) mRNA, c	0.050	0.087	4 consistent probesets
N29665	yw73e06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.050	0.089	4 consistent probesets
AL050369	Homo sapiens mRNA; cDNA DKFZp566J153 (from clone D	0.050	0.090	3 inconsistent probesets
M96824	Human nucleobindin precursor mRNA, complete cds /cds=(	0.050	0.091	2 inconsistent probesets
U15780	Human p82 (ST5) mRNA, alternatively spliced, complete	0.050	0.091	4 inconsistent probesets
X75252	H.sapiens phosphatidylethanolamine binding protein mRN	0.050	0.093	4 consistent probesets
AF085692	Homo sapiens multidrug resistance-associated protein 3	0.050	0.095	4 consistent probesets
AI201310	qf71b11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-175	0.050	0.096	4 consistent probesets
AI417075	tg78e09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-211	0.050	0.101	4 consistent probesets
AB011147	Homo sapiens mRNA for KIAA0575 protein, complete cds /	0.050	0.101	4 consistent probesets
AA522537	ni38e08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-979	0.050	0.101	4 consistent probesets
U75686	Homo sapiens polyadenylate binding protein mRNA, compl	0.050	0.102	4 consistent probesets
AL050021	Homo sapiens mRNA; cDNA DKFZp564D016 (from clone D	0.050	0.102	4 consistent probesets
Y07593	H.sapiens mRNA for 46 kDa coxsackievirus and adenoviru	0.050	0.102	4 consistent probesets
L33799	Human procollagen C-proteinase enhancer protein (PCOLC	0.050	0.109	4 consistent probesets
L05425	Homo sapiens autoantigen mRNA, complete cds /cds=(79,2	0.050	0.110	4 consistent probesets
AF038660	Homo sapiens chromosome 1p33-p34 beta-1,4-galactosyltr	0.050	0.112	4 consistent probesets
Z48482	HSMMPM2 H.sapiens mRNA for membrane-type matr	0.050	0.113	3 consistent probesets
M85289	Human heparan sulfate proteoglycan (HSPG2) mRNA, com	0.050	0.113	3 consistent probesets
AL096750	Homo sapiens mRNA; cDNA DKFZp434H244 (from clone D	0.050	0.113	2 consistent probesets
U66684	HSU66684 Homo sapiens cDNA /gb=U66684 /gi=1906569 /	0.050	0.118	3 consistent probesets
AF038406	Homo sapiens NADH dehydrogenase-ubiquinone Fe-S prot	0.050	0.118	4 consistent probesets
U87947	Human hematopoietic neural membrane protein (HNMP-1)	0.050	0.119	3 consistent probesets
AL049929	Homo sapiens mRNA; cDNA DKFZp547O0510 (from clone	0.050	0.125	4 consistent probesets
M93107	Homo sapiens heart (R)-3-hydroxybutyrate dehydrogenase	0.050	0.126	3 consistent probesets
AF070539	Homo sapiens clone 24433 myelodysplasia/myeloid leukem	0.050	0.130	4 consistent probesets
U86529	HSU86529 Human glutathione transferase Zeta 1	0.050	0.137	4 consistent probesets
M69177	Human monoamine oxidase B (MAOB) mRNA, complete co	0.050	0.138	4 consistent probesets
X69532	H.sapiens gene for inter-alpha-trypsin inhibitor heavy	0.050	0.144	3 consistent probesets
Y09160	H.sapiens Sub1.5 mRNA /cds=(435,3044) /gb=Y09160 /gi=	0.050	0.147	3 consistent probesets
X69910	H.sapiens p63 mRNA for transmembrane protein /cds=(84,	0.050	0.149	4 consistent probesets
AB014555	Homo sapiens mRNA for KIAA0655 protein, partial cds /c	0.050	0.151	1 consistent probesets
X60364	Human ALAS mRNA for 5-aminolevulinate synthase precur	0.050	0.151	1 consistent probesets
AF049891	Homo sapiens tyrosylprotein sulfotransferase-2 mRNA, c	0.050	0.153	3 consistent probesets
X63465	H.sapiens hGDS mRNA for smg GDS /cds=(0,1676) /gb=X6	0.050	0.156	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
X98261	H.sapiens mRNA for M-phase phosphoprotein, mpp5 /cds=	0.050	0.161	2 consistent probesets
U92315	Homo sapiens hydroxysteroid sulfotransferase SULT2B1b	0.050	0.161	2 consistent probesets
L00190	Human antithrombin III (ATIII) gene /cds=(46,1440) /gb	0.050	0.162	4 consistent probesets
U20657	HSU20657 Human ubiquitin protease (Unph) prot	0.050	0.164	6 consistent probesets
D89618	Homo sapiens mRNA for karyopherin alhph 3, complete cd	0.050	0.164	4 consistent probesets
AB029031	Homo sapiens mRNA for KIAA1108 protein, partial cds /c	0.050	0.164	3 consistent probesets
AB020698	Homo sapiens mRNA for KIAA0891 protein, partial cds /c	0.050	0.165	3 consistent probesets
AF052106	Homo sapiens clone 23781 mRNA sequence /cds=UNKNO	0.050	0.165	3 consistent probesets
K03195	Human (HepG2) glucose transporter gene mRNA, complete	0.050	0.166	4 consistent probesets
Y17108	Homo sapiens mRNA for rhomboid-related protein, comple	0.050	0.169	2 consistent probesets
X04297	Human mRNA for Na,K-ATPase alpha-subunit /cds=(318,33	0.050	0.171	4 consistent probesets
U02680	HSU02680 Human protein tyrosine kinase mRNA,	0.050	0.172	4 consistent probesets
AJ005892	Homo sapiens mRNA for JM23 protein, complete coding se	0.050	0.178	4 consistent probesets
X61587	H.sapiens rhoG mRNA for GTPase /cds=(129,704) /gb=X61	0.050	0.178	2 consistent probesets
D50920	Human mRNA for KIAA0130 gene, complete cds /cds=(73,3	0.050	0.178	2 consistent probesets
AB018270	Homo sapiens mRNA for KIAA0727 protein, partial cds /c	0.050	0.179	3 consistent probesets
AL080062	Homo sapiens mRNA; cDNA DKFZp564I122 (from clone D	0.050	0.182	2 consistent probesets
AA161065	zl50h04.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-505	0.050	0.183	3 consistent probesets
L42451	Homo sapiens pyruvate dehydrogenase kinase isoenzyme 2	0.050	0.192	3 inconsistent probesets
X74794	HSP1CDC21 H.sapiens P1-Cdc21 mRNA	0.050	0.201	3 consistent probesets
X90857	H.sapiens mRNA for -14 gene, containing globin regulat	0.050	0.202	1 consistent probesets
X14675	HSBCR3C Human bcr-abl mRNA 5 fragment (clone	0.050	0.202	1 consistent probesets
AB028996	Homo sapiens mRNA for KIAA1073 protein, complete cds /	0.050	0.202	1 consistent probesets
U78678	Human thioredoxin mRNA, nuclear gene encoding mitochor	0.050	0.202	1 consistent probesets
W28620	49c2 Homo sapiens cDNA /gb=W28620 /gi=1308568 /	0.050	0.202	1 consistent probesets
U49395	Human ionotropic ATP receptor P2X5a mRNA, complete cd	0.050	0.202	1 consistent probesets
AJ238097	Homo sapiens mRNA for Lsm5 protein /cds=(0,275) /gb=AJ	0.050	0.208	3 consistent probesets
D32002	Human mRNA for nuclear cap binding protein, complete c	0.050	0.209	3 consistent probesets
X67098	H.sapiens rTS alpha mRNA containing four open reading	0.050	0.214	2 consistent probesets
W25934	15b3 Homo sapiens cDNA /gb=W25934 /gi=1306057 /	0.050	0.223	3 consistent probesets
U49957	Human LIM protein (LPP) mRNA, partial cds /cds=(246,20	0.050	0.224	3 inconsistent probesets
U66616	HSU66616 Human SWI/SNF complex 170 KDa subuni	0.050	0.226	3 consistent probesets
D79995	Human mRNA for KIAA0173 gene, complete cds /cds=(207	0.050	0.231	4 consistent probesets
U42391	Human myosin-IXb mRNA, complete cds /cds=(0,6068) /gb	0.050	0.252	1 consistent probesets
AI345337	tb81g11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-206	0.050	0.252	1 consistent probesets
U31216	Human metabotropic glutamate receptor 1 beta (mGluR1be	0.050	0.252	1 consistent probesets
M13928	Human delta-aminolevulinate dehydratase mRNA, complete	0.050	0.252	1 consistent probesets
D88667	Homo sapiens mRNA for cerebroside sulfotransferase, co	0.050	0.252	2 consistent probesets
X71346	Homo sapiens HNF1-B mRNA /cds=UNKNOWN /gb=X7134	0.050	0.252	1 consistent probesets
W27858	39e3 Homo sapiens cDNA /gb=W27858 /gi=1307869 /	0.050	0.271	4 consistent probesets
U11037	Human Sel-1 like mRNA, complete cds /cds=(11,298) /gb=	0.050	0.282	3 consistent probesets
U79262	Human deoxyhypusine synthase mRNA, complete cds /cds=	0.050	0.302	1 consistent probesets
AA872560	oa16f04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-130	0.050	0.302	1 consistent probesets
AL050306	Human DNA sequence from clone 475B7 on chromosome X	0.050	0.302	1 consistent probesets
AL049382	Homo sapiens mRNA; cDNA DKFZp586K1318 (from clone	0.050	0.302	1 consistent probesets
Y15801	Homo sapiens mRNA for PRKY protein /cds=UNKNOWN /g	0.050	0.302	1 consistent probesets
AB002331	Human mRNA for KIAA0333 gene, partial cds /cds=(0,2977	0.050	0.302	1 consistent probesets
H23429	H23429 ym52d12.s1 Soares infant brain 1NIB Ho	0.050	0.302	1 consistent probesets
AB023967	Homo sapiens mRNA for Rod1, complete cds /cds=(46,161	0.050	0.348	3 consistent probesets
AB018329	Homo sapiens mRNA for KIAA0786 protein, partial cds /c	0.050	0.353	1 consistent probesets
AB019410	Homo sapiens mRNA, expressed in fibroblasts of periodo	0.050	0.353	1 consistent probesets
AF048732	Homo sapiens cyclin T2b mRNA, complete cds /cds=(0,219	0.050	0.353	1 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
M24594	HUMII56KD Human interferon-inducible 56 Kd pr	0.050	0.359	3 consistent probesets
W28275	44g12 Homo sapiens cDNA /gb=W28275 /gi=1308241 /	0.050	0.371	3 consistent probesets
W28510	48f3 Homo sapiens cDNA /gb=W28510 /gi=1308521 /	0.050	0.403	1 consistent probesets
AB023226	Homo sapiens mRNA for KIAA1009 protein, complete cds /	0.050	0.403	1 consistent probesets
AL080196	Homo sapiens mRNA; cDNA DKFZp434C212 (from clone D	0.050	0.406	2 consistent probesets
W25984	17e5 Homo sapiens cDNA /gb=W25984 /gi=1306251 /	0.050	0.427	3 consistent probesets
L42324	HUMFRCG Homo sapiens (clone GPCR W) G protein	0.050	0.454	1 consistent probesets
U80456	Human transcription factor SIM2 long form mRNA, comple	0.050	0.454	1 consistent probesets
AJ005016	Homo sapiens mRNA for putative ABC transporter, partia	0.050	0.454	1 consistent probesets
AF070523	Homo sapiens JWA protein mRNA, complete cds /cds=(115	0.050	0.475	2 consistent probesets
AA156237	zl50c09.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-505	0.050	0.478	2 consistent probesets
D86407	Homo sapiens DNA for apoER2 /cds=(236,3127) /gb=D864	0.050	0.500	2 consistent probesets
AB023157	Homo sapiens mRNA for KIAA0940 protein, complete cds /	0.050	0.502	2 consistent probesets
AL080220	Homo sapiens mRNA; cDNA DKFZp586P0123 (from clone	0.050	0.504	1 consistent probesets
AL049365	Homo sapiens mRNA; cDNA DKFZp586A0618 (from clone	0.050	0.554	1 consistent probesets
AB028994	Homo sapiens mRNA for KIAA1071 protein, partial cds /c	0.050	0.554	1 consistent probesets
AF027156	Homo sapiens alpha 1,2-mannosidase IB mRNA, complete	0.050	0.554	1 consistent probesets
L40371	Homo sapiens thyroid receptor interactor (TRIP4) mRNA,	0.050	0.580	2 consistent probesets
AF041260	Homo sapiens AIBC1 (AIBC1) mRNA, complete cds /cds=(	0.050	0.605	1 consistent probesets
U60062	Human FEZ1-T mRNA, alternatively spliced form, complet	0.050	0.634	1 consistent probesets
U00930	Human clone C4E 1.63 (CAC)n/(GTG)n repeat-containing r	0.050	0.756	1 consistent probesets
Z49194	H.sapiens mRNA for oct-binding factor /cds=(523,1293)	0.050	0.907	1 consistent probesets
L23320	Human replication factor C large subunit mRNA, complet	0.050	0.958	1 consistent probesets
AL049351	Homo sapiens mRNA; cDNA DKFZp566C093 (from clone D	0.050	0.958	1 consistent probesets
N32859	yw88d02.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.050	1.361	1 consistent probesets
M17886	Human acidic ribosomal phosphoprotein P1 mRNA, comple	0.050	0.042	8 consistent probesets
U62531	Human AE2 anion exchanger (SLC4A2) mRNA, complete c	0.050	0.067	4 inconsistent probesets
D50310	HUMCYI Human mRNA for cyclin I, complete cds"	0.050	0.087	4 inconsistent probesets
AL050118	Homo sapiens mRNA; cDNA DKFZp586C201 (from clone D	0.050	0.118	3 consistent probesets
X53416	Human mRNA for actin-binding protein (filamin) (ABP-28	0.050	0.126	2 consistent probesets
L28997	Homo sapiens ARL1 mRNA, complete cds /cds=(144,689) /	0.050	0.130	4 consistent probesets
AF093420	Homo sapiens Hsp70 binding protein HspBP1 mRNA, comp	0.050	0.147	2 consistent probesets
AF043117	Homo sapiens ubiquitin-fusion degradation protein 2 (U	0.050	0.174	4 consistent probesets
U40572	Human beta2-syntrophin (SNT B2) mRNA, complete cds /c	0.050	0.227	4 inconsistent probesets
D13243	Homo sapiens gene for pyruvate kinase L /cds=(39,1838)	0.050	0.232	2 consistent probesets
U39447	Human placenta copper monamine oxidase mRNA, comple	0.050	0.249	4 consistent probesets
M33318	HUMCPIIA3A Human cytochrome P450IIA3 (CYP2A3)	0.050	0.273	2 consistent probesets
AL050035	Homo sapiens mRNA; cDNA DKFZp566H0124 (from clone	0.050	0.378	2 consistent probesets
AF071202	Homo sapiens ABC transporter MOAT-B (MOAT-B) mRNA,	0.050	0.413	2 consistent probesets
L11695	Human activin receptor-like kinase (ALK-5) mRNA, compl	0.050	0.559	2 consistent probesets
U54996	Human protein ZW10 homolog (HZW10) mRNA, complete	0.050	0.566	2 consistent probesets
AB007976	Homo sapiens mRNA, chromosome 1 specific transcript KI	0.050	0.590	2 consistent probesets
X15334	Human gene for creatine kinase B (EC 2.7.3.2) /cds=(80	0.048	0.056	8 consistent probesets
U49283	Human NAD+-specific isocitrate dehydrogenase beta subu	0.048	0.057	8 inconsistent probesets
M22430	HUMRASFAB Human RASF-A PLA2 mRNA, complete cd	0.048	0.090	8 consistent probesets
M97935	Homo sapiens transcription factor ISGF-3 mRNA, complet	0.047	0.078	17 inconsistent probesets
M17885	Human acidic ribosomal phosphoprotein P0 mRNA, comple	0.046	0.039	4 consistent probesets
W26633	34b1 Homo sapiens cDNA /gb=W26633 /gi=1307476 /	0.046	0.041	4 inconsistent probesets
D13315	Human mRNA for lactoyl glutathione lyase /cds=(87,641)	0.046	0.041	4 inconsistent probesets
X04098	Human mRNA for cytoskeletal gamma-actin /cds=(73,1200	0.046	0.050	4 consistent probesets
X64364	H.sapiens mRNA for M6 antigen /cds=(57,866) /gb=X64364	0.046	0.051	4 inconsistent probesets
AI653621	tz21b11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-228	0.046	0.052	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
X53281	H.sapiens BTF3b mRNA /cds=(239,727) /gb=X53281 /gi=29	0.046	0.053	4 inconsistent probesets
AF006043	Homo sapiens 3-phosphoglycerate dehydrogenase mRNA,	0.046	0.054	4 consistent probesets
M12523	Human serum albumin (ALB) gene, complete cds /cds=(39,	0.046	0.055	4 consistent probesets
X56009	Human GSA mRNA for alpha subunit of GsGTP binding pro	0.046	0.055	4 inconsistent probesets
X53463	Human mRNA for glutathione peroxidase-like protein /cd	0.046	0.058	4 consistent probesets
S60099	APPH=amyloid precursor protein homolog [human, placent	0.046	0.058	4 inconsistent probesets
S57501	protein phosphatase type 1 catalytic subunit [human, m	0.046	0.058	4 consistent probesets
X13546	Human HMG-17 gene for non-histone chromosomal protein	0.046	0.060	4 inconsistent probesets
V00572	Human mRNA encoding phosphoglycerate kinase /cds=(79	0.046	0.061	4 consistent probesets
X16832	Human mRNA for cathepsin H (EC 3.4.22.16) /cds=(34,104	0.046	0.061	4 consistent probesets
L38696	Homo sapiens autoantigen p542 mRNA, complete cds /cds=	0.046	0.061	4 consistent probesets
M62839	Human apolipoprotein H mRNA, complete cds /cds=UNKN(	0.046	0.064	4 consistent probesets
L03532	Human M4 protein mRNA, complete cds /cds=(11,2200) /gb	0.046	0.064	4 consistent probesets
D16562	Human mRNA for ATP synthase gamma-subunit (L-type), c	0.046	0.064	4 inconsistent probesets
T89651	yd99a05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-11(	0.046	0.064	4 consistent probesets
AF099149	Homo sapiens TRIAD1 type I mRNA, complete cds /cds=(1	0.046	0.064	4 inconsistent probesets
U25182	Human antioxidant enzyme AOE37-2 mRNA, complete cds	0.046	0.064	4 consistent probesets
D43950	Human mRNA for KIAA0098 gene, partial cds /cds=(0,1642	0.046	0.064	4 inconsistent probesets
U89322	Homo sapiens nucleophosmin phosphoprotein/B23 (NPM) g	0.046	0.065	4 consistent probesets
D86973	Human mRNA for KIAA0219 gene, partial cds /cds=(0,7239	0.046	0.068	4 consistent probesets
D16294	Human mRNA for mitochondrial 3-oxoacyl-CoA thiolase, c	0.046	0.068	4 consistent probesets
X16064	Human mRNA for translationally controlled tumor protei	0.046	0.070	4 consistent probesets
AA631972	fmfc39 Homo sapiens cDNA /clone=CR7-5 /gb=AA631972 /	0.046	0.070	4 consistent probesets
L01124	Human ribosomal protein S13 (RPS13) mRNA, complete co	0.046	0.070	4 consistent probesets
N90862	zb11b06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-30	0.046	0.070	4 consistent probesets
Y13374	Homo sapiens mRNA for putatively prenylated protein /c	0.046	0.072	4 inconsistent probesets
AB007618	Homo sapiens mRNA for COX7RP, complete cds /cds=(40,	0.046	0.072	4 consistent probesets
L38941	Homo sapiens ribosomal protein L34 (RPL34) mRNA, comp	0.046	0.073	4 consistent probesets
AL038662	DKFZp566I0346_r1 Homo sapiens cDNA, 5 end /clone=DK	0.046	0.076	4 consistent probesets
AL096737	Homo sapiens mRNA; cDNA DKFZp434F152 (from clone D	0.046	0.077	4 consistent probesets
AB014587	Homo sapiens mRNA for KIAA0687 protein, partial cds /c	0.046	0.078	4 consistent probesets
X04526	Human liver mRNA for beta-subunit signal transducing p	0.046	0.079	4 consistent probesets
AF054174	Homo sapiens histone macroH2A1.2 mRNA, complete cds /	0.046	0.079	4 consistent probesets
X13293	HSBMYB Human mRNA for B-myb gene	0.046	0.081	4 consistent probesets
AF006088	Homo sapiens Arp2/3 protein complex subunit p16-Arc (A	0.046	0.081	4 consistent probesets
U47077	HSU47077 Homo sapiens DNA-dependent protein k	0.046	0.081	8 inconsistent probesets
AF067575	untitled /cds=(21,2564) /gb=AF067575 /gi=3789867 /	0.046	0.082	4 inconsistent probesets
D38076	Human mRNA for RanBP1 (Ran-binding protein 1), comple	0.046	0.086	4 consistent probesets
M81601	HUMTEFSII Human transcription elongation fact	0.046	0.086	4 consistent probesets
AI525393	PT1.1_07_A11.r Homo sapiens cDNA, 5 end /clone_end=5	0.046	0.086	4 consistent probesets
AA142942	zl43c04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-504	0.046	0.088	4 consistent probesets
U49070	Human peptidyl-prolyl isomerase and essential mitotic	0.046	0.088	4 consistent probesets
U44772	Human palmitoyl protein thioesterase mRNA, complete cd	0.046	0.088	4 inconsistent probesets
AI816034	au44e05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.046	0.089	4 inconsistent probesets
AB014511	Homo sapiens mRNA for KIAA0611 protein, partial cds /c	0.046	0.090	4 consistent probesets
D14657	Human mRNA for KIAA0101 gene, complete cds /cds=(61,3	0.046	0.091	4 consistent probesets
U54559	Homo sapiens translation initiation factor eIF3 p40 su	0.046	0.091	4 consistent probesets
AF004709	Homo sapiens stress-activated protein kinase 4 mRNA, c	0.046	0.092	4 consistent probesets
D00632	HUMGSHPXA Homo sapiens mRNA for glutathione p	0.046	0.093	4 consistent probesets
AJ238096	Homo sapiens mRNA for Lsm4 protein /cds=(23,442) /gb=A	0.046	0.094	4 inconsistent probesets
AA203303	zx55b01.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-446	0.046	0.095	4 consistent probesets
AL037557	DKFZp564H2472_r1 Homo sapiens cDNA, 5 end /clone=D	0.046	0.095	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AA205857	zq50e04.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-645	0.046	0.095	4 consistent probesets
U59057	Human beta-A4 crystallin (CRYBA4) mRNA, complete cds /	0.046	0.095	4 consistent probesets
U74612	Human hepatocyte nuclear factor-3/fork head homolog 11	0.046	0.096	4 consistent probesets
AB006198	Homo sapiens mRNA for SART-1, complete cds /cds=(38,2	0.046	0.098	4 inconsistent probesets
Y11395	H.sapiens mRNA for p40 /cds=(104,1303) /gb=Y11395 /gi=	0.046	0.098	4 consistent probesets
W51774	zc48b04.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-325	0.046	0.098	4 consistent probesets
AA310786	EST181572 Homo sapiens cDNA, 5 end /clone=ATCC-156	0.046	0.098	4 consistent probesets
Z22555	H.sapiens encoding CLA-1 mRNA /cds=(69,1598) /gb=Z225	0.046	0.098	4 consistent probesets
U50733	Human dynamitin mRNA, complete cds /cds=(78,1298) /gb=	0.046	0.098	4 consistent probesets
M22760	Homo sapiens nuclear-encoded mitochondrial cytochrome	0.046	0.100	4 consistent probesets
U78302	Human 2,4-dienoyl-CoA reductase gene /cds=(73,1080) /g	0.046	0.100	4 consistent probesets
M37400	Human cytosolic aspartate aminotransferase mRNA, compl	0.046	0.104	4 consistent probesets
AF071309	Homo sapiens OPA-containing protein mRNA, complete cd	0.046	0.105	4 consistent probesets
D63998	Human mRNA for golgi alpha-mannosidaseII, complete cds	0.046	0.109	4 consistent probesets
Y10387	H.sapiens mRNA for PAPS synthetase /cds=(36,1910) /gb=	0.046	0.111	4 consistent probesets
X75593	H.sapiens mRNA for rab 13 /cds=(139,750) /gb=X75593 /g	0.046	0.115	4 consistent probesets
S81916	phosphoglycerate kinase {alternatively spliced} [human	0.046	0.116	4 inconsistent probesets
X06745	HSPOLAR Human mRNA for DNA polymerase alpha-s	0.046	0.116	4 consistent probesets
AJ012409	Homo sapiens mRNA for hypothetical protein, clone YR-2	0.046	0.116	4 consistent probesets
U33053	HSU33053 Human lipid-activated protein kinase	0.046	0.117	4 consistent probesets
W27152	23h11 Homo sapiens cDNA /gb=W27152 /gi=1306731 /	0.046	0.118	4 consistent probesets
X92475	H.sapiens mRNA for ITBA1 protein /cds=(284,1069) /gb=X	0.046	0.118	4 consistent probesets
U23143	Human mitochondrial serine hydroxymethyltransferase ge	0.046	0.121	4 consistent probesets
Y11651	H.sapiens mRNA for phosphate cyclase /cds=(170,1270) /	0.046	0.122	4 consistent probesets
U64444	HSU64444 Homo sapiens ubiquitin fusion-degrad	0.046	0.122	4 consistent probesets
AI700633	we38g03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.046	0.125	4 consistent probesets
J04080	Human complement component C1r mRNA, complete cds /	0.046	0.127	4 consistent probesets
AL049951	Homo sapiens mRNA; cDNA DKFZp564O0122 (from clone	0.046	0.127	4 consistent probesets
X59871	Human TCF-1 mRNA for T cell factor 1 (splice form C) /	0.046	0.133	4 inconsistent probesets
U95032	Human growth-arrest-specific protein 2 mRNA, complete	0.046	0.133	4 consistent probesets
AB020695	Homo sapiens mRNA for KIAA0888 protein, partial cds /c	0.046	0.137	4 consistent probesets
U93181	Homo sapiens nuclear dual-specificity phosphatase (SBF	0.046	0.145	4 consistent probesets
D88152	Homo sapiens mRNA for acetyl-coenzyme A transporter, c	0.046	0.145	4 consistent probesets
Z28339	H.sapiens mRNA for delta 4-3-oxosteroid 5 beta-reducta	0.046	0.160	4 consistent probesets
AF091086	Homo sapiens clone 640 unknown mRNA, complete sequer	0.046	0.163	4 consistent probesets
M14565	Human cholesterol side-chain cleavage enzyme P450scc m	0.046	0.174	4 consistent probesets
AF067853	Homo sapiens adenylosuccinate lyase (ADSL) mRNA, alter	0.046	0.193	4 consistent probesets
L40388	HUMTRIP15M Homo sapiens thyroid receptor inte	0.046	0.213	4 consistent probesets
AF064804	Homo sapiens transcription factor SUPT3H (SUPT3H) mRN	0.046	0.248	4 consistent probesets
AF001891	Homo sapiens clone lambda MEN1 region unknown protein	0.045	0.071	7 inconsistent probesets
W28518	48a1 Homo sapiens cDNA /gb=W28518 /gi=1308466 /	0.044	0.213	3 consistent probesets
X58965	HSNM23H2G H.sapiens RNA for nm23-H2 gene	0.044	0.034	8 consistent probesets
J04031	HUMMDMCSF Human methylenetetrahydrofolate deh	0.044	0.063	8 consistent probesets
M63488	HUMRPA70KD Human replication protein A 70kDa	0.044	0.082	8 consistent probesets
U50410	Human heparan sulphate proteoglycan (OCI5) mRNA, com	0.042	0.035	4 inconsistent probesets
X55715	Human Hums3 mRNA for 40S ribosomal protein s3 /cds=(2	0.042	0.035	4 consistent probesets
AI540925	PEC1.2_15_A02.r Homo sapiens cDNA, 5 end /clone_end=	0.042	0.037	4 inconsistent probesets
X56932	H.sapiens mRNA for 23 kD highly basic protein /cds=(17	0.042	0.039	4 consistent probesets
AL022326	dJ333H23.2.2 (Synaptogyrin 1A (SYNGR1A)) /cds=(43,744	0.042	0.043	4 consistent probesets
AL031670	similar to Zinc finger, C3HC4 type (RING finger); matc	0.042	0.045	4 consistent probesets
Z49148	H.sapiens mRNA for ribosomal protein L29 /cds=(29,508)	0.042	0.047	4 inconsistent probesets
D78361	HUMODAZ Human mRNA for ornithine decarboxylas	0.042	0.048	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF022229	Homo sapiens translation initiation factor 6 (eIF6) mR	0.042	0.050	4 inconsistent probesets
X15183	Human mRNA for 90-kDa heat-shock protein /cds=(60,2258	0.042	0.053	4 consistent probesets
U03271	Human F-actin capping protein beta subunit mRNA, compl	0.042	0.055	4 inconsistent probesets
AL031295	Human DNA sequence from clone 886K2 on chromosome 1	0.042	0.055	4 consistent probesets
X03342	Human mRNA for ribosomal protein L32 /cds=(34,441) /gb	0.042	0.056	4 consistent probesets
U14971	Human ribosomal protein S9 mRNA, complete cds /cds=(35	0.042	0.057	4 consistent probesets
AI541542	libtest16.A02.r Homo sapiens cDNA, 5 end /clone_end=5	0.042	0.058	4 consistent probesets
AI912041	wd84b06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.042	0.060	4 inconsistent probesets
D23662	HUMULP Homo sapiens mRNA for ubiquitin-like p	0.042	0.061	4 consistent probesets
X58536	Human mRNA for HLA class I locus C heavy chain /cds=(1	0.042	0.061	4 consistent probesets
U73377	Human p66shc (SHC) mRNA, complete cds /cds=(194,1945	0.042	0.062	4 consistent probesets
D29012	HUMPSY Human mRNA for proteasome subunit Y, c	0.042	0.064	4 consistent probesets
M94250	HUMMKXX Human retinoic acid inducible factor	0.042	0.064	4 consistent probesets
AI741833	wg29e04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.042	0.064	4 consistent probesets
L13278	Homo sapiens zeta-crystallin/quinone reductase mRNA, c	0.042	0.067	4 inconsistent probesets
X73066	HSNM23H1A H.sapiens NM23-H1 mRNA	0.042	0.068	4 consistent probesets
N95406	zb80g12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-30	0.042	0.068	4 consistent probesets
X95404	H.sapiens mRNA for non-muscle type cofilin /cds=(51,55	0.042	0.069	4 consistent probesets
D14043	Human mRNA for MGC-24, complete cds /cds=(79,648) /gb	0.042	0.069	4 inconsistent probesets
J03909	HUMIIP Human gamma-interferon-inducible prote	0.042	0.070	8 consistent probesets
M64110	Human caldesmon mRNA, complete cds /cds=(111,1727) /g	0.042	0.070	4 inconsistent probesets
R98910	yr31d04.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-206	0.042	0.070	4 consistent probesets
X96752	H.sapiens mRNA for L-3-hydroxyacyl-CoA dehydrogenase /	0.042	0.072	4 inconsistent probesets
AF027204	Homo sapiens putative tetraspan transmembrane protein	0.042	0.072	4 consistent probesets
X75861	H.sapiens TEGT gene /cds=(40,753) /gb=X75861 /gi=45625	0.042	0.072	4 inconsistent probesets
S65738	actin depolymerizing factor [human, fetal brain, mRNA,	0.042	0.074	4 inconsistent probesets
AF050640	Homo sapiens NADH-ubiquinone oxidoreductase NDUFS2 s	0.042	0.075	4 consistent probesets
AI541256	pec1.2-3.F11.r Homo sapiens cDNA, 5 end /clone_end=5	0.042	0.076	4 consistent probesets
AL096879	Novel human mRNA similar to C. elegans gene WP-CE186	0.042	0.078	4 consistent probesets
M77349	HUMTGFBIG Human transforming growth factor-be	0.042	0.078	4 consistent probesets
AI740522	wg16b07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.042	0.078	4 consistent probesets
M60858	Human nucleolin gene, complete cds /cds=(111,2234) /gb	0.042	0.078	4 consistent probesets
U86602	Human nucleolar protein p40 mRNA, complete cds /cds=(1	0.042	0.080	4 inconsistent probesets
J05272	Human IMP dehydrogenase type 1 mRNA complete cds /cd	0.042	0.081	4 consistent probesets
L20941	Human ferritin heavy chain mRNA, complete cds /cds=(20	0.042	0.082	4 consistent probesets
S72008	hCDC10=CDC10 homolog [human, fetal lung, mRNA, 2314	0.042	0.082	4 consistent probesets
AB024301	Homo sapiens mRNA for RuvB-like DNA helicase TIP49b, c	0.042	0.082	4 consistent probesets
AA189161	zq45g01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-63	0.042	0.084	4 consistent probesets
AF070535	Homo sapiens clone 24432 mRNA sequence /cds=UNKNO	0.042	0.086	4 inconsistent probesets
U33760	HSU33760 Human cyclin A/CDK2-associated p19 (	0.042	0.086	4 consistent probesets
N73769	za61g08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-29	0.042	0.088	4 consistent probesets
D87444	Human mRNA for KIAA0255 gene, complete cds /cds=(340	0.042	0.089	4 consistent probesets
AF027302	Homo sapiens TNF-alpha stimulated ABC protein (ABC50)	0.042	0.091	4 consistent probesets
AF038957	Homo sapiens translation initiation factor 4e mRNA, co	0.042	0.091	4 consistent probesets
J00314	HUMTBBM40 Human beta-tubulin gene, clone m40"	0.042	0.091	4 consistent probesets
D16469	Human mRNA for ORF, Xq terminal portion /cds=(1353,219	0.042	0.093	4 inconsistent probesets
AA883868	am26e11.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-14	0.042	0.097	4 consistent probesets
W22541	69B4 Homo sapiens cDNA /clone=(not-directional) /gb=W2	0.042	0.098	4 consistent probesets
AB029003	Homo sapiens mRNA for KIAA1080 protein, partial cds /c	0.042	0.098	4 consistent probesets
Z99129	Human DNA sequence from clone 425C14 on chromosome	0.042	0.100	4 inconsistent probesets
AF091083	Homo sapiens clone 628 unknown mRNA, complete sequer	0.042	0.105	4 consistent probesets
M23161	Human transposon-like element mRNA /cds=UNKNOWN /g	0.042	0.107	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
X65488	H.sapiens U21.1 mRNA /cds=(41,2461) /gb=X65488 /gi=32	0.042	0.107	4 consistent probesets
AL050015	Homo sapiens mRNA; cDNA DKFZp564O243 (from clone D	0.042	0.107	4 consistent probesets
M65131	Human methylmalonyl-CoA mutase (MCM) mRNA, complet	0.042	0.108	4 consistent probesets
D63478	Human mRNA for KIAA0144 gene, complete cds /cds=(106	0.042	0.108	4 consistent probesets
D50692	HUMAMY1 Homo sapiens mRNA for c-myc binding p	0.042	0.114	4 inconsistent probesets
AF014398	Homo sapiens myo-inositol monophosphatase 2 mRNA, co	0.042	0.115	4 consistent probesets
X15674	Human pTR5 mRNA for repetitive sequence /cds=UNKNOV	0.042	0.116	4 consistent probesets
W52999	zc02f10.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-321	0.042	0.117	4 consistent probesets
M38449	HUMTGFBA Human transforming growth factor-bet	0.042	0.120	4 consistent probesets
AI400326	tg89c03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-211	0.042	0.121	4 consistent probesets
L33881	HUMPKCI Human protein kinase C iota isoform,	0.042	0.124	8 consistent probesets
U18932	Human heparan sulfate-N-deacetylase/N-sulfotransferase	0.042	0.128	4 consistent probesets
W28257	44c1 Homo sapiens cDNA /gb=W28257 /gi=1308205 /	0.042	0.128	4 inconsistent probesets
U37122	Human adducin gamma subunit mRNA, complete cds /cds=	0.042	0.129	4 consistent probesets
AL050275	Homo sapiens mRNA; cDNA DKFZp566D213 (from clone D	0.042	0.130	4 consistent probesets
AJ224326	Homo sapiens mRNA for putative ribulose-5-phosphate-ep	0.042	0.135	4 consistent probesets
AL109672	Homo sapiens mRNA full length insert cDNA clone EUROIN	0.042	0.136	4 consistent probesets
X85019	H.sapiens mRNA for UDP-GalNAc-polypeptide N-acetylgala	0.042	0.136	4 consistent probesets
AF039704	Homo sapiens lysosomal pepstatin insensitive protease	0.042	0.141	4 consistent probesets
U63825	Human hepatitis delta antigen interacting protein A (d	0.042	0.158	4 consistent probesets
U79272	Human clone 23720 mRNA sequence /cds=UNKNOWN /gb	0.042	0.166	4 consistent probesets
H94881	yu57f07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-230	0.042	0.168	4 consistent probesets
D63485	Human mRNA for KIAA0151 gene, complete cds /cds=(326	0.042	0.173	4 consistent probesets
AB002312	Human mRNA for KIAA0314 gene, partial cds /cds=(0,3723	0.042	0.189	4 consistent probesets
S69790	S69790 Brush-1=tumor suppressor {3 region} [	0.042	0.202	4 inconsistent probesets
D12625	HUMNF1ISO Human mRNA for NF1 protein isoform	0.042	0.205	4 consistent probesets
U69198	U69198 Homo sapiens cDNA /clone=c-32h10 /gb=U69198 /	0.042	0.219	4 consistent probesets
N30625	yw72f06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-257	0.042	0.245	4 consistent probesets
X57985	H.sapiens genes for histones H2B.1 and H2A /cds=(42,42	0.042	0.248	4 consistent probesets
U81561	Human protein tyrosine phosphatase receptor pi (PTPRP)	0.042	0.265	4 consistent probesets
U88620	Human 8-hydroxyguanine glycosylase (hMMH) mRNA, com	0.042	0.275	4 consistent probesets
M24194	Human MHC protein homologous to chicken B complex pro	0.040	0.029	8 inconsistent probesets
M34064	HUMNCADH Human N-cadherin mRNA, complete cds"	0.040	0.068	8 inconsistent probesets
M10943	Human metallothionein-If gene (hMT-If) /cds=(0,185) /g	0.039	0.101	3 consistent probesets
AF050145	Homo sapiens iduronate-2-sulfatase (IDS) mRNA, complet	0.039	0.194	3 consistent probesets
D28915	Human gene for hepatitis C-associated microtubular agg	0.039	0.369	3 consistent probesets
AF102803	untitled /cds=(2,2722) /gb=AF102803 /gi=4092760 /	0.038	0.108	7 consistent probesets
U46116	HSPTPRG28 Human receptor tyrosine phosphatase	0.038	0.146	7 consistent probesets
U34995	Human normal keratinocyte substraction library mRNA, c	0.038	0.032	4 inconsistent probesets
X17206	Human mRNA for LLRep3 /cds=(240,905) /gb=X17206 /gi=	0.038	0.034	4 inconsistent probesets
M88249	Human inter-alpha-trypsin inhibitor light chain (ITI)	0.038	0.047	4 inconsistent probesets
X67309	H.sapiens gene for ribosomal protein S6 /cds=(42,791)	0.038	0.047	4 consistent probesets
AF054187	Homo sapiens alpha NAC mRNA, complete cds /cds=(309,9	0.038	0.050	8 inconsistent probesets
AI989422	ws25a09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.038	0.052	4 inconsistent probesets
X71129	H.sapiens mRNA for electron transfer flavoprotein beta	0.038	0.053	4 inconsistent probesets
U58048	Human metallopeptidase PRSM1 mRNA, complete cds /cds	0.038	0.053	4 inconsistent probesets
L06132	Human voltage-dependent anion channel isoform 1 (VDAC)	0.038	0.055	4 inconsistent probesets
AL050268	Homo sapiens mRNA; cDNA DKFZp564B163 (from clone D	0.038	0.057	8 inconsistent probesets
X15573	Human liver-type 1-phosphofructokinase (PFKL) mRNA, co	0.038	0.058	4 consistent probesets
Y10805	H.sapiens mRNA for arginine methyltransferase, splice	0.038	0.060	4 inconsistent probesets
X98248	H.sapiens mRNA for sortilin /cds=(21,2522) /gb=X98248	0.038	0.061	4 inconsistent probesets
AI347088	qp60d09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-19	0.038	0.062	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
Y13286	Homo sapiens mRNA for GDP dissociation inhibitor beta	0.038	0.062	4 consistent probesets
M19961	Human cytochrome c oxidase subunit Vb (coxVb) mRNA, c	0.038	0.064	4 consistent probesets
U56637	Human capping protein alpha subunit isoform 1 mRNA, co	0.038	0.066	4 consistent probesets
L19185	Human natural killer cell enhancing factor (NKEFB) mRN	0.038	0.067	4 consistent probesets
AI540958	PEC1.2_15_H01.r Homo sapiens cDNA, 5 end /clone_end=	0.038	0.068	4 inconsistent probesets
U61397	HSU61397 Human ubiquitin-homology domain prot	0.038	0.068	4 inconsistent probesets
AF006082	Homo sapiens actin-related protein Arp2 (ARP2) mRNA, c	0.038	0.070	4 inconsistent probesets
D87442	Human mRNA for KIAA0253 gene, partial cds /cds=(0,2127	0.038	0.070	4 consistent probesets
D84273	Homo sapiens mRNA for Asparaginyl tRNA Synthetase, cor	0.038	0.070	4 consistent probesets
AL080212	Homo sapiens mRNA; cDNA DKFZp586H0723 (from clone	0.038	0.072	4 consistent probesets
AA926959	om68h08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-15	0.038	0.072	4 consistent probesets
D78151	HUM26SPSP Human mRNA for 26S proteasome subun	0.038	0.072	4 consistent probesets
AF044209	Homo sapiens nuclear receptor co-repressor N-CoR mRNA	0.038	0.073	4 inconsistent probesets
AB019517	Homo sapiens PKIG mRNA for protein kinase inhibitor ga	0.038	0.078	4 consistent probesets
U31383	Human G protein gamma-10 subunit mRNA, complete cds /	0.038	0.081	4 inconsistent probesets
AB021288	Homo sapiens mRNA for beta 2-microglobulin, complete c	0.038	0.082	4 consistent probesets
U22055	Human 100 kDa coactivator mRNA, complete cds /cds=(26	0.038	0.083	4 inconsistent probesets
AI799802	wc43d09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.038	0.086	4 consistent probesets
AF104421	Homo sapiens isolate normal patient 1 uroporphyrinogen	0.038	0.087	4 consistent probesets
U35451	Homo sapiens heterochromatin protein p25 mRNA, comple	0.038	0.088	4 consistent probesets
D14705	HUMALPHAC Human mRNA for alpha-catenin, compl	0.038	0.088	4 consistent probesets
AF006484	Homo sapiens putative oral tumor suppressor protein (d	0.038	0.090	4 consistent probesets
U46571	Human tetratricopeptide repeat protein (tpr2) mRNA, co	0.038	0.090	4 inconsistent probesets
AL035398	Human DNA sequence from clone 796I17 on chromosome	0.038	0.091	4 consistent probesets
X91257	H.sapiens mRNA for seryl-tRNA synthetase /cds=(75,1619	0.038	0.092	4 consistent probesets
M57567	Human ADP-ribosylation factor (hARF5) mRNA, complete c	0.038	0.093	4 consistent probesets
U79270	Human clone 23707 mRNA, partial cds /cds=(0,460) /gb=U	0.038	0.093	4 consistent probesets
M58600	Human heparin cofactor II (HCF2) gene, exons 1 through	0.038	0.094	4 consistent probesets
AL031668	Human DNA sequence from clone 64K7 on chromosome 20	0.038	0.094	4 consistent probesets
W28979	54e8 Homo sapiens cDNA /gb=W28979 /gi=1308927 /	0.038	0.094	4 consistent probesets
AL031659	dJ343K2.2.1 (ribophorin II (isoform 1)) /cds=(284,2179	0.038	0.098	4 consistent probesets
AB009398	Homo sapiens mRNA for 26S proteasome subunit p40.5, co	0.038	0.098	4 consistent probesets
AL035079	dJ53C18.1 (Catalase) /cds=(74,1657) /gb=AL035079 /gi=4	0.038	0.098	4 consistent probesets
U69609	Human transcriptional repressor (GCF2) mRNA, complete	0.038	0.102	4 inconsistent probesets
W28330	45d4 Homo sapiens cDNA /gb=W28330 /gi=1308278 /	0.038	0.103	4 inconsistent probesets
D13634	Human mRNA for KIAA0009 gene, complete cds /cds=(17,9	0.038	0.103	4 consistent probesets
U81006	Human p76 mRNA, complete cds /cds=(133,2124) /gb=U81	0.038	0.106	4 consistent probesets
AA203354	zx58b07.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-446	0.038	0.107	4 consistent probesets
AF075599	Homo sapiens ubiquitin conjugating enzyme 12 (UBC12) m	0.038	0.107	4 consistent probesets
U07736	Human quinone oxidoreductase2 (NQO2) gene /cds=(273,9	0.038	0.108	4 consistent probesets
S74221	S74221 IK=IK factor [human, leukemic cells K5	0.038	0.108	4 consistent probesets
S69272	cytoplasmic antiproteinase=38 kda intracellular serine	0.038	0.110	4 consistent probesets
D29963	Homo sapiens mRNA for CD151, complete cds /cds=(84,84	0.038	0.112	4 consistent probesets
X82206	H.sapiens mRNA for alpha-centractin /cds=(66,1196) /gb	0.038	0.117	4 consistent probesets
U64028	Human NADH-ubiquinone oxidoreductase subunit B13 mRN	0.038	0.118	4 consistent probesets
X60484	H.sapiens H4/e gene for H4 histone /cds=(0,311) /gb=X6	0.038	0.119	4 consistent probesets
L02320	Human radixin mRNA, complete cds /cds=(30,1781) /gb=L0	0.038	0.123	4 inconsistent probesets
AL080155	Homo sapiens mRNA; cDNA DKFZp434J154 (from clone D	0.038	0.124	4 consistent probesets
AF018631	untitled /cds=(35,1666) /gb=AF018631 /gi=2674074 /	0.038	0.133	4 consistent probesets
AB020648	Homo sapiens mRNA for KIAA0841 protein, partial cds /c	0.038	0.134	4 consistent probesets
U28413	Human Cockayne syndrome complementation group A CSA	0.038	0.135	4 consistent probesets
AF068195	Homo sapiens putative glialblastoma cell differentiati	0.038	0.135	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF015308	Homo sapiens nucleolar protein (MSP58) mRNA, complete	0.038	0.136	4 consistent probesets
U05569	Human alphaA-crystallin (CRYA1) mRNA, complete cds /cd	0.038	0.156	4 consistent probesets
AA746355	oa56f02.r1 Homo sapiens cDNA /clone=IMAGE-1308987 /g	0.038	0.157	4 consistent probesets
AJ224538	Homo sapiens mRNA for AMP-activated protein kinase bet	0.038	0.185	4 inconsistent probesets
AL079277	Homo sapiens mRNA full length insert cDNA clone EUROIM	0.038	0.202	4 consistent probesets
AA143321	zo37d01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-58	0.038	0.207	4 consistent probesets
X55079	Human lysosomal alpha-glucosidase gene exon 1 /cds=(44	0.037	0.085	4 inconsistent probesets
AB003184	Homo sapiens mRNA for ISLR, complete cds /cds=(98,1384	0.037	0.098	4 inconsistent probesets
M58526	Human alpha-5 collagen type IV (COL4A5) mRNA, 3 end /	0.037	0.132	4 consistent probesets
M21188	Human insulin-degrading enzyme (IDE) mRNA, complete c	0.037	0.189	4 inconsistent probesets
AF051323	Homo sapiens Src-associated adaptor protein (SAPS) mRN	0.037	0.201	4 consistent probesets
AB020682	Homo sapiens mRNA for KIAA0875 protein, partial cds /c	0.036	0.193	6 consistent probesets
U19796	HSU19796 Human melanoma antigen p15 mRNA, com	0.035	0.051	8 inconsistent probesets
J02902	HUMP2A Human protein phosphatase 2A regulator	0.035	0.051	8 consistent probesets
X78136	H.sapiens hnRNP-E2 mRNA /cds=(22,1119) /gb=X78136 /g	0.035	0.052	8 inconsistent probesets
X75755	H.sapiens PR264 gene /cds=(109,774) /gb=X75755 /gi=455	0.035	0.060	8 inconsistent probesets
L07493	HUM14RPA Homo sapiens replication protein A 1	0.035	0.080	8 consistent probesets
X02162	Human mRNA for apolipoprotein AI (apo AI)= /cds=(86,88	0.033	0.045	4 consistent probesets
M58459	Human ribosomal protein (RPS4Y) isoform mRNA, complet	0.033	0.045	4 consistent probesets
AF052124	Homo sapiens clone 23810 osteopontin mRNA, complete c	0.033	0.053	4 inconsistent probesets
AI701049	wc78b08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.033	0.055	4 consistent probesets
M62762	Human vacuolar H+ ATPase proton channel subunit mRNA	0.033	0.055	4 consistent probesets
K02215	HUMANG Human angiotensinogen mRNA, complete C	0.033	0.056	4 consistent probesets
M31520	Human ribosomal protein S24 mRNA /cds=(142,543) /gb=M	0.033	0.057	4 consistent probesets
X74929	H.sapiens KRT8 mRNA for keratin 8 /cds=(59,1510) /gb=X	0.033	0.058	4 inconsistent probesets
M25915	Human complement cytolysis inhibitor (CLI) mRNA, compl	0.033	0.058	4 inconsistent probesets
D89667	D89667 Homo sapiens mRNA for c-myc binding pr	0.033	0.059	4 inconsistent probesets
AA522698	ni39d03.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-979	0.033	0.060	4 inconsistent probesets
T79616	yd71e06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-11	0.033	0.060	4 consistent probesets
AI525652	PT1.3_04_C04.r Homo sapiens cDNA, 5 end /clone_end=5	0.033	0.060	4 inconsistent probesets
U14969	Human ribosomal protein L28 mRNA, complete cds /cds=(2	0.033	0.060	4 consistent probesets
AI935551	wo97g09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.033	0.061	4 consistent probesets
U14970	Human ribosomal protein S5 mRNA, complete cds /cds=(37	0.033	0.061	4 consistent probesets
AJ133534	Homo sapiens mRNA for prenylated Rab acceptor 1 /cds=(	0.033	0.061	4 consistent probesets
Y09022	H.sapiens mRNA for Not56-like protein /cds=(31,1347) /	0.033	0.062	4 consistent probesets
AF037643	Homo sapiens 60S ribosomal protein L12 (RPL12) pseudog	0.033	0.062	4 consistent probesets
AI557240	PT2.1_15_C11.r Homo sapiens cDNA, 3 end /clone_end=3	0.033	0.062	4 consistent probesets
X60036	H.sapiens mRNA for mitochondrial phosphate carrier pro	0.033	0.064	4 consistent probesets
U25849	Human red cell-type low molecular weight acid phosphat	0.033	0.064	4 consistent probesets
M15182	Human beta-glucuronidase mRNA, complete cds /cds=(26,1	0.033	0.064	4 consistent probesets
M80244	Human E16 mRNA, complete cds /cds=(310,1035) /gb=M80	0.033	0.065	4 consistent probesets
Y00815	Human mRNA for LCA-homolog. LAR protein (leukocyte an	0.033	0.068	4 consistent probesets
AF054175	Homo sapiens mitochondrial proteolipid 68MP homolog mR	0.033	0.068	4 consistent probesets
X07173	Human mRNA for second protein of inter-alpha-trypsin i	0.033	0.069	4 consistent probesets
J04977	HUMKUANT Human Ku autoimmune antigen gene, co	0.033	0.070	4 consistent probesets
AA477898	zu34f08.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-739	0.033	0.070	4 consistent probesets
L28101	Homo sapiens kallistatin (PI4) gene, exons 1-4, comple	0.033	0.072	4 consistent probesets
M64098	Human high density lipoprotein binding protein (HBP) m	0.033	0.072	4 inconsistent probesets
U02493	Human 54 kDa protein mRNA, complete cds /cds=(85,1500	0.033	0.072	4 consistent probesets
U90913	Human clone 23665 mRNA sequence /cds=UNKNOWN /gb	0.033	0.073	4 consistent probesets
L08044	Human intestinal trefoil factor mRNA, complete cds /cd	0.033	0.074	4 inconsistent probesets
Y00345	Human mRNA for polyA binding protein /cds=(502,2403) /	0.033	0.074	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF020038	Homo sapiens NADP-dependent isocitrate dehydrogenase (	0.033	0.078	4 consistent probesets
X78283	H.sapiens mRNA for aryl sulfotransferase (ST1A3) /cds=	0.033	0.078	4 consistent probesets
AA526812	ni92a08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-984	0.033	0.078	4 consistent probesets
U24578	Human RP1 and complement C4B precursor (C4B) genes,	0.033	0.078	4 inconsistent probesets
M24398	Human parathymosin mRNA, complete cds /cds=(300,608)	0.033	0.079	3 consistent probesets
M26326	Human keratin 18 mRNA, complete cds /cds=(51,1343) /gb	0.033	0.079	4 consistent probesets
Z49107	H.sapiens mRNA for galectin /cds=(87,1058) /gb=Z49107	0.033	0.080	4 consistent probesets
AB028974	Homo sapiens mRNA for KIAA1051 protein, partial cds /c	0.033	0.080	4 inconsistent probesets
AL050283	Homo sapiens mRNA; cDNA DKFZp586K0919 (from clone	0.033	0.082	4 consistent probesets
AF013759	Homo sapiens calumein (Calu) mRNA, complete cds /cds=(	0.033	0.084	4 consistent probesets
AW007731	wt68d11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	0.033	0.085	4 consistent probesets
AF044414	Homo sapiens alpha mannosidase 6A8B (6a8b) mRNA, cor	0.033	0.085	4 consistent probesets
AI827793	wf33b11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.033	0.085	4 consistent probesets
AB007937	Homo sapiens mRNA for KIAA0468 protein, complete cds /	0.033	0.086	4 inconsistent probesets
AB020718	Homo sapiens mRNA for KIAA0911 protein, complete cds /	0.033	0.086	4 consistent probesets
AL046940	DKFZp586I0517_r1 Homo sapiens cDNA, 5 end /clone=DK	0.033	0.087	4 consistent probesets
AB016869	Homo sapiens mRNA for p70 ribosomal S6 kinase beta, co	0.033	0.089	4 consistent probesets
M36803	Human hemopexin gene /cds=(28,1416) /gb=M36803 /gi=4	0.033	0.090	3 consistent probesets
AB009285	Homo sapiens BCNT mRNA, complete cds /cds=(109,1008	0.033	0.091	4 consistent probesets
N50520	yy89b05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-28(	0.033	0.091	4 consistent probesets
AF035555	Homo sapiens short chain L-3-hydroxyacyl-CoA dehydroge	0.033	0.091	4 consistent probesets
AF035287	Homo sapiens clone 23742 mRNA, partial cds /cds=(0,466	0.033	0.093	4 consistent probesets
AL080199	Homo sapiens mRNA; cDNA DKFZp434E082 (from clone D	0.033	0.094	4 consistent probesets
Z24727	H.sapiens tropomyosin isoform mRNA, complete CDS /cds=	0.033	0.095	4 consistent probesets
W27949	39h3 Homo sapiens cDNA /gb=W27949 /gi=1307897 /	0.033	0.100	4 consistent probesets
AB014460	Homo sapiens TSC2, NTHL1/NTH1 and SLC9A3R2/E3KAR	0.033	0.101	3 consistent probesets
AF020736	Homo sapiens ATPase homolog mRNA, complete cds /cds=	0.033	0.103	4 consistent probesets
X99270	H.sapiens Xq28, 2000bp sequence contg. ORF /cds=(544,1	0.033	0.104	4 consistent probesets
L40386	HUMDP2M Human DP-2 mRNA, complete cds"	0.033	0.104	4 consistent probesets
AF052510	Homo sapiens CTP-phosphocholine cytidylyltransferase b	0.033	0.105	5 consistent probesets
U40152	Human origin recognition complex 1 (HsORC1) mRNA, con	0.033	0.110	3 consistent probesets
U42031	Human 54 kDa progesterone receptor-associated immunop	0.033	0.110	4 consistent probesets
U51698	HSU51698 Homo sapiens cDNA /gb=U51698 /gi=1255268 /	0.033	0.111	4 consistent probesets
M15036	Human vitamin K-dependent plasma protein S mRNA, com	0.033	0.111	4 consistent probesets
Z11793	H.sapiens mRNA for selenoprotein P /cds=(36,1181) /gb=	0.033	0.114	4 consistent probesets
AF067730	Homo sapiens TLS-associated protein TASR-2 mRNA, com	0.033	0.115	4 consistent probesets
N36926	yy38e04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-27	0.033	0.116	4 consistent probesets
J05249	HUMREPA Human replication protein A 32-kDa su	0.033	0.118	4 consistent probesets
Z14093	H.sapiens mRNA for branched chain decarboxylase alpha	0.033	0.120	4 consistent probesets
X95384	Homo sapiens mRNA for translational inhibitor protein	0.033	0.122	4 inconsistent probesets
U60899	Human lysosomal alpha-mannosidase (manB) gene /cds=(3	0.033	0.122	4 consistent probesets
U34994	HSU34994 Homo sapiens DNA dependent protein k	0.033	0.123	4 consistent probesets
W27517	31h6 Homo sapiens cDNA /gb=W27517 /gi=1307321 /	0.033	0.123	3 consistent probesets
S79219	S79219 metastasis-associated gene [human, hig	0.033	0.128	4 consistent probesets
AA034074	zi06c05.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-429	0.033	0.131	4 consistent probesets
AB006202	Homo sapiens mRNA for cytochrome b small subunit of co	0.033	0.132	4 consistent probesets
U56418	Human lysophosphatidic acid acyltransferase-beta mRNA,	0.033	0.140	4 consistent probesets
AB018280	Homo sapiens mRNA for KIAA0737 protein, complete cds /	0.033	0.141	4 consistent probesets
AF006621	Homo sapiens embryonic lung protein (HUEL) mRNA, comp	0.033	0.141	4 consistent probesets
AL035081	H.sapiens mRNA similar to Xenopus laevis mRNA for KDEL	0.033	0.141	4 consistent probesets
AB019529	Homo sapiens mRNA for OGG1 protein type 2c, partial cd	0.033	0.153	3 consistent probesets
X79882	H.sapiens lrp mRNA /cds=(105,2795) /gb=X79882 /gi=8958	0.033	0.159	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AB018353	Homo sapiens mRNA for KIAA0810 protein, partial cds /c	0.033	0.161	4 consistent probesets
L04490	Homo sapiens (clone CC6) NADH-ubiquinone oxidoreducta	0.033	0.162	4 consistent probesets
D38537	Human mRNA for protoporphyrinogen oxidase, complete co	0.033	0.163	3 consistent probesets
L11702	HUMPHOSPDP Human phospholipase D mRNA, comple	0.033	0.168	1 consistent probesets
U16031	HSU16031 Human transcription factor IL-4 Stat	0.033	0.171	4 consistent probesets
AF073771	Homo sapiens RNA polymerase II termination factor mRNA	0.033	0.176	3 consistent probesets
AF104942	Homo sapiens ABC transporter MOAT-C (MOAT-C) mRNA	0.033	0.189	3 consistent probesets
AL009266	H. sapiens cDNA similar to C. elegans RNA binding prot	0.033	0.191	3 consistent probesets
J05158	Human carboxypeptidase N mRNA, 3 end /cds=(0,1610) /g	0.033	0.200	3 consistent probesets
AL049758	dJ437M21.3 (novel Src homology domain 3 containing pro	0.033	0.208	3 consistent probesets
AB028985	Homo sapiens mRNA for KIAA1062 protein, partial cds /c	0.033	0.223	3 consistent probesets
M26747	HUMCERBA Human c-erbA mRNA, complete cds"	0.033	0.235	1 consistent probesets
M62810	Human mitochondrial transcription factor 1 mRNA, compl	0.033	0.243	4 consistent probesets
U38964	HSU38964 Human PMS2 related (hPMSR2) gene, co	0.033	0.245	3 consistent probesets
D86965	Human mRNA for KIAA0210 gene, complete cds /cds=(166	0.033	0.256	3 consistent probesets
X75304	H.sapiens giantin mRNA /cds=(126,9905) /gb=X75304 /gi=	0.033	0.256	3 consistent probesets
M32313	HUM5AR Human steroid 5-alpha-reductase mRNA,	0.033	0.263	3 consistent probesets
AI796281	wh49a09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.033	0.269	1 consistent probesets
AL080110	Homo sapiens mRNA; cDNA DKFZp586G1922 (from clone	0.033	0.329	3 consistent probesets
M33197	Human glyceraldehyde-3-phosphate dehydrogenase (GAPD	0.032	0.030	12 inconsistent probesets
M33336	HUMCAMPPK Human cAMP-dependent protein kinase	0.032	0.037	12 consistent probesets
U12255	Human IgG Fc receptor hFcRn mRNA, complete cds /cds=(	0.031	0.053	8 inconsistent probesets
X68742	HSINTASA H.sapiens mRNA for integrin, alpha s	0.031	0.123	8 consistent probesets
AI767675	wh38d11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.031	0.088	7 inconsistent probesets
AI985964	wr79d08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.031	0.084	6 inconsistent probesets
J03592	Human ADP/ATP translocase mRNA, 3 end, clone pHAT8	0.029	0.042	8 inconsistent probesets
M55914	HUMCMYCQ Human c-myc binding protein (MBP-1)	0.029	0.047	4 consistent probesets
X64707	H.sapiens BBC1 mRNA /cds=(51,686) /gb=X64707 /gi=293	0.029	0.048	4 inconsistent probesets
S75463	P43=mitochondrial elongation factor homolog [human, li	0.029	0.050	4 consistent probesets
AL050282	Homo sapiens mRNA; cDNA DKFZp586H2219 (from clone	0.029	0.051	8 inconsistent probesets
M11353	HUMHISH3C Human H3.3 histone class C mRNA, co	0.029	0.055	4 inconsistent probesets
AB028972	Homo sapiens mRNA for KIAA1049 protein, partial cds /c	0.029	0.057	8 consistent probesets
M22632	Human mitochondrial aspartate aminotransferase mRNA, c	0.029	0.060	4 inconsistent probesets
D63486	Human mRNA for KIAA0152 gene, complete cds /cds=(128	0.029	0.061	4 consistent probesets
X15414	Human mRNA for aldose reductase (EC 1.1.1.2) /cds=(45,	0.029	0.062	4 consistent probesets
X57152	HSCKIIBE Human gene for casein kinase II subu	0.029	0.064	4 consistent probesets
U18009	Human chromosome 17q21 mRNA clone LF113 /cds=(0,93	0.029	0.064	4 inconsistent probesets
D78177	Homo sapiens mRNA for quinolinate phosphoribosyl trans	0.029	0.072	4 consistent probesets
X64037	H.sapiens mRNA for RNA polymerase II associated protei	0.029	0.075	4 consistent probesets
D37931	Human mRNA for RNase 4, complete cds /cds=(27,470) /gb	0.029	0.075	4 consistent probesets
AB017644	Homo sapiens mRNA for ubiquitin-conjugating enzyme E2,	0.029	0.075	4 consistent probesets
U18300	HSU18300 Human damage-specific DNA binding pr	0.029	0.075	4 consistent probesets
X91788	H.sapiens mRNA for Icln protein /cds=(88,801) /gb=X917	0.029	0.075	4 consistent probesets
D14662	Human mRNA for KIAA0106 gene, complete cds /cds=(43,7	0.029	0.076	4 consistent probesets
AF046889	Homo sapiens lysyl hydroxylase isoform 3 (PLOD3) mRNA,	0.029	0.080	4 consistent probesets
M63573	Human secreted cyclophilin-like protein (SCYLP) mRNA,	0.029	0.081	4 consistent probesets
U41514	Human UDP-GalNAc-polypeptide N-acetylgalactosaminyltra	0.029	0.082	4 consistent probesets
M13970	HUMGFI21 Human insulin-like growth factor (IG	0.029	0.084	4 inconsistent probesets
AF010187	Homo sapiens FGF-1 intracellular binding protein (FIBP	0.029	0.085	4 consistent probesets
X79353	H.sapiens XAP-4 mRNA for GDP-dissociation inhibitor /c	0.029	0.085	4 consistent probesets
X55110	Human mRNA for neurite outgrowth-promoting protein /cd	0.029	0.086	4 consistent probesets
AA669799	ag36c04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-11	0.029	0.086	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
W28931	56f3 Homo sapiens cDNA /gb=W28931 /gi=1309086 /	0.029	0.087	4 consistent probesets
X84908	H.sapiens mRNA for phosphorylase-kinase, beta subunit	0.029	0.091	4 consistent probesets
L11245	Human (clone A19) C4b-binding protein beta-chain mRNA,	0.029	0.093	4 consistent probesets
Y00971	Human mRNA for phosphoriobosyl pyrophosphate syntheta	0.029	0.094	4 consistent probesets
AJ237946	Homo sapiens mRNA for DEAD Box Protein 5 /cds=(0,1439	0.029	0.096	4 consistent probesets
W26659	34d2 Homo sapiens cDNA /gb=W26659 /gi=1307502 /	0.029	0.097	4 consistent probesets
X59892	H.sapiens mRNA for IFN-inducible gamma2 protein /cds=(	0.029	0.098	4 consistent probesets
AA004795	zh96a06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-42	0.029	0.098	4 consistent probesets
U61263	Human acetolactate synthase homolog mRNA, complete co	0.029	0.098	4 consistent probesets
M93405	Human methylmalonate semialdehyde dehydrogenase gene	0.029	0.101	4 consistent probesets
M95929	Human homeobox protein (PHOX1) mRNA, 3 end /cds=(0,	0.029	0.105	4 consistent probesets
Y10183	H.sapiens mRNA for MEMD protein /cds=(0,1748) /gb=Y10	0.029	0.112	4 consistent probesets
AJ005893	Homo sapiens mRNA for JM26 protein, complete CDS (clo	0.029	0.113	4 consistent probesets
AL120500	DKFZp761M078_s1 Homo sapiens cDNA, 3 end /clone=DK	0.029	0.113	4 consistent probesets
AI961040	wq58f01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.029	0.114	4 consistent probesets
X59766	H.sapiens mRNA for Zn-alpha2-glycoprotein /cds=(10,918	0.029	0.115	4 consistent probesets
U54645	Human adenylate kinase 2B (adk2b) gene, complete cds /	0.029	0.120	4 consistent probesets
J04131	Human gamma-glutamyl transpeptidase (GGT) protein mR	0.029	0.128	4 consistent probesets
M60725	HUMP70S6KB Human p70 ribosomal S6 kinase alph	0.029	0.159	4 inconsistent probesets
AL049989	Homo sapiens mRNA; cDNA DKFZp564F052 (from clone D	0.029	0.167	4 consistent probesets
X73114	H.sapiens mRNA for slow MyBP-C /cds=(81,3452) /gb=X73	0.029	0.175	4 consistent probesets
X59543	Human mRNA for M1 subunit of ribonucleotide reductase	0.029	0.183	4 consistent probesets
D63875	Human mRNA for KIAA0155 gene, complete cds /cds=(86,3	0.029	0.188	4 consistent probesets
U09578	HSU09578 Homo sapiens MAPKAP kinase (3pK) mRN	0.029	0.190	4 consistent probesets
U64105	HSU64105 Human guanine nucleotide exchange fa	0.029	0.199	4 consistent probesets
Y08134	H.sapiens mRNA for ASM-like phosphodiesterase 3b /cds=	0.029	0.216	4 consistent probesets
U00943	Human clone A9A2BRB2 (CAC)n/(GTG)n repeat-containing	0.029	0.227	4 consistent probesets
AB010812	Homo sapiens Nrf3 mRNA for NF-E2-related factor 3, com	0.029	0.238	4 consistent probesets
U50553	HSU50553 Homo sapiens helicase like protein 2	0.029	0.240	4 consistent probesets
AA917672	on46d09.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-15	0.029	0.255	4 consistent probesets
U37359	Homo sapiens MRE11 homologue hMre11 mRNA, complet	0.029	0.309	4 consistent probesets
J03810	Human liver glucose transporter-like protein (GLUT2),	0.028	0.226	3 consistent probesets
AF010315	Homo sapiens Pig11 (PIG11) mRNA, complete cds /cds=(1	0.028	0.259	3 consistent probesets
AA977163	oq25a04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-15	0.027	0.050	8 inconsistent probesets
X03473	HSHIS10G Human gene for histone H1(0)	0.027	0.064	8 inconsistent probesets
AF006751	Homo sapiens ES/130 mRNA, complete cds /cds=(70,3003	0.027	0.065	8 consistent probesets
U41745	HSU41745 Human PDGF associated protein mRNA,	0.027	0.073	8 inconsistent probesets
L00635	HUMFPTB Human farnesyl-protein transferase be	0.026	0.093	7 consistent probesets
L13210	Human Mac-2 binding protein mRNA, complete cds /cds=(1	0.025	0.053	4 consistent probesets
U28686	Human putative RNA binding protein RNPL mRNA, comple	0.025	0.057	4 consistent probesets
L06498	Homo sapiens ribosomal protein S20 (RPS20) mRNA, com	0.025	0.058	4 consistent probesets
X60188	HSERK1 Human ERK1 mRNA for protein serine/thr	0.025	0.060	4 inconsistent probesets
Z49878	H.sapiens mRNA for guanidinoacetate N-methyltransferas	0.025	0.060	4 consistent probesets
L13939	Homo sapiens beta adaptin (BAM22) mRNA, complete cds	0.025	0.061	4 consistent probesets
X02544	Human mRNA for alpha1-acid glycoprotein (orosomucoid)	0.025	0.064	4 consistent probesets
U37408	Homo sapiens phosphoprotein CtBP mRNA, complete cds /	0.025	0.066	8 inconsistent probesets
M10050	Human liver fatty acid binding protein (FABP) mRNA, co	0.025	0.070	4 consistent probesets
D13627	Human mRNA for KIAA0002 gene, complete cds /cds=(28,1	0.025	0.074	4 consistent probesets
AL035304	H.sapiens gene from PAC 295C6, similar to rat PO44 /cd	0.025	0.074	4 inconsistent probesets
D50918	Human mRNA for KIAA0128 gene, partial cds /cds=(0,1276	0.025	0.075	4 consistent probesets
U91932	Homo sapiens AP-3 complex sigma3A subunit mRNA, com	0.025	0.080	4 consistent probesets
AI017574	ou23f10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-162	0.025	0.082	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF035309	Homo sapiens clone 23598 mRNA, complete cds /cds=(880	0.025	0.084	4 consistent probesets
AF068754	Homo sapiens heat shock factor binding protein 1 HSBP1	0.025	0.085	4 consistent probesets
AL050159	Homo sapiens mRNA; cDNA DKFZp586A0522 (from clone	0.025	0.086	4 consistent probesets
X05276	Human mRNA for fibroblast tropomyosin TM30 (pl) /cds=(	0.025	0.089	4 consistent probesets
D14686	Human gene for glycine cleavage system T-protein /cds=	0.025	0.091	4 consistent probesets
AA524058	ng33b12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-93	0.025	0.097	4 consistent probesets
AB026894	Homo sapiens mRNA for NESCA, complete cds /cds=(125,	0.025	0.098	4 consistent probesets
AF023462	Homo sapiens peroxisomal phytanoyl-CoA alpha-hydroxyla	0.025	0.102	4 consistent probesets
Y08685	H.sapiens mRNA for serine palmitoyltransferase, subuni	0.025	0.104	4 consistent probesets
AF009615	Homo sapiens ADAM10 (ADAM10) mRNA, complete cds /c	0.025	0.108	4 consistent probesets
AF061034	Homo sapiens FIP2 alternatively translated mRNA, compl	0.025	0.110	8 inconsistent probesets
X13710	H.sapiens unspliced mRNA for glutathione peroxidase /c	0.025	0.112	4 consistent probesets
AB020712	Homo sapiens mRNA for KIAA0905 protein, complete cds /	0.025	0.117	4 consistent probesets
Y14391	Homo sapiens mRNA for putative GTP-binding protein /cd	0.025	0.120	4 consistent probesets
AJ010277	Homo sapiens mRNA for TBX19 protein /cds=(51,1397) /gb	0.025	0.128	4 consistent probesets
AB018268	Homo sapiens mRNA for KIAA0725 protein, partial cds /c	0.025	0.133	4 consistent probesets
U43784	HSU43784 Human mitogen activated protein kina	0.025	0.135	4 consistent probesets
AI871359	wl81b11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.025	0.147	2 consistent probesets
AJ224335	Homo sapien mRNA for putative secretory protein, hBET3	0.025	0.156	4 consistent probesets
X93209	H.sapiens mRNA for NRD1 convertase /cds=UNKNOWN /g	0.025	0.159	4 consistent probesets
X95715	HSARAPROT H.sapiens mRNA for anthracycline re	0.025	0.178	2 consistent probesets
AF042800	Homo sapiens suppressor of white apricot homolog 2 (SW	0.025	0.178	2 consistent probesets
M34480	Human platelet glycoprotein IIb (GPIIb) mRNA, complete	0.025	0.178	2 consistent probesets
AF006740	Homo sapiens Coch-5B2 mRNA, complete cds /cds=(56,17	0.025	0.212	4 consistent probesets
L40392	Homo sapiens (clone S164) mRNA, 3 end of cds /cds=(0,	0.025	0.214	2 consistent probesets
L15409	HUMHIPLIND Homo sapiens (clone g7) von Hippel	0.025	0.214	2 consistent probesets
X56832	H.sapiens ENO3 gene for muscle specific enolase /cds=(	0.025	0.215	2 consistent probesets
AB006533	Homo sapiens RecQ5 mRNA for DNA helicase, complete c	0.025	0.245	2 consistent probesets
AB004922	AB004922S1 Homo sapiens gene for Smad 3, exon	0.025	0.249	2 consistent probesets
AI459274	tk11f11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-215	0.025	0.294	2 consistent probesets
AJ005801	Homo sapiens mRNA for protein phosphatase 2C (beta) /c	0.025	0.296	4 consistent probesets
AA996066	os33d01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.025	0.308	2 consistent probesets
W28558	48f7 Homo sapiens cDNA /gb=W28558 /gi=1308524 /	0.025	0.308	2 consistent probesets
AL096717	Homo sapiens mRNA; cDNA DKFZp564P0662 (from clone	0.025	0.441	2 consistent probesets
N98607	zb82d01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-31	0.025	0.478	2 consistent probesets
AB023215	Homo sapiens mRNA for KIAA0998 protein, partial cds /c	0.025	0.481	2 consistent probesets
AF055016	Homo sapiens clone 24774 unknown mRNA, partial cds /cd	0.025	0.524	2 consistent probesets
AF000987	Homo sapiens eIF-1A, Y isoform (EIF1AY) mRNA, complet	0.025	0.530	2 consistent probesets
J03077	Human co-beta glucosidase (proactivator) mRNA, complet	0.025	0.062	4 inconsistent probesets
AA768912	nz82h06.s1 Homo sapiens cDNA /clone=IMAGE-1302011 /	0.025	0.066	8 inconsistent probesets
X57352	Human 1-8U gene from interferon-inducible gene family	0.025	0.093	4 consistent probesets
U67322	Human HBV associated factor (XAP4) mRNA, complete cds	0.025	0.095	4 consistent probesets
H10201	ym02c07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-46	0.025	0.104	4 consistent probesets
Z68907	H.sapiens mRNA for NAD (H)-specific isocitrate dehydro	0.025	0.111	4 consistent probesets
W26651	34c5 Homo sapiens cDNA /gb=W26651 /gi=1307494 /	0.025	0.118	4 consistent probesets
AF100153	Homo sapiens connector enhancer of KSR-like protein CN	0.025	0.136	4 consistent probesets
AF052149	Homo sapiens clone 24733 mRNA sequence /cds=UNKNO	0.025	0.148	4 consistent probesets
AI362017	qy39a10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-20	0.025	0.152	4 consistent probesets
L05628	HUMMRPX Human multidrug resistance-associated	0.025	0.282	2 consistent probesets
AB007958	Homo sapiens mRNA, chromosome 1 specific transcript KI	0.025	0.287	2 consistent probesets
X53795	Human R2 mRNA for an inducible membrane protein /cds=	0.025	0.336	2 consistent probesets
AL050267	Homo sapiens mRNA; cDNA DKFZp564A032 (from clone D	0.025	0.469	2 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AB029343	Homo sapiens HCR (a-helix coiled-coil rod homologue) g	0.024	0.181	7 consistent probesets
AA527880	nh86h10.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-96	0.023	0.078	8 consistent probesets
AF023268	Homo sapiens clk2 kinase (CLK2), propin1, cote1, gluco	0.023	0.078	8 inconsistent probesets
U70063	HSU70063 Human acid ceramidase mRNA, complete	0.023	0.115	8 inconsistent probesets
AB018551	Homo sapiens ATPBL mRNA for coiled-coil protein, compl	0.022	0.136	3 consistent probesets
AJ224162	Homo sapiens mRNA for putative lipoic acid synthetase,	0.022	0.200	3 consistent probesets
AL050082	Homo sapiens mRNA; cDNA DKFZp566J2446 (from clone	0.022	0.359	3 consistent probesets
X89416	HSRNAPPP5 H.sapiens mRNA for protein phosphat	0.021	0.095	7 consistent probesets
Z28407	H.sapiens mRNA for ribosomal protein L8 /cds=(43,816)	0.021	0.035	4 consistent probesets
X01683	Human mRNA for alpha 1-antitrypsin /cds=(38,1294) /gb=	0.021	0.035	4 inconsistent probesets
J04173	Homo sapiens phosphoglycerate mutase (PGAM-B) mRNA	0.021	0.048	4 consistent probesets
M30448	Human casein kinase II beta subunit mRNA, complete cds	0.021	0.050	4 inconsistent probesets
X52851	Human cyclophilin gene for cyclophilin (EC 5.2.1.8) /c	0.021	0.050	4 consistent probesets
U05861	Human hepatic dihydrodiol dehydrogenase gene /cds=(26,	0.021	0.050	4 consistent probesets
X63527	H.sapiens mRNA for ribosomal protein L19 /cds=(28,618)	0.021	0.053	4 consistent probesets
J04765	HUMOSTRO Human osteopontin mRNA, complete cds	0.021	0.055	4 inconsistent probesets
X67247	H.sapiens rpS8 gene for ribosomal protein S8 /cds=(23,	0.021	0.055	4 consistent probesets
M19828	HUMAPB21 Human apolipoprotein B-100 (apoB) ge	0.021	0.057	4 inconsistent probesets
X02152	Human mRNA for lactate dehydrogenase-A (LDH-A, EC 1.1	0.021	0.057	4 consistent probesets
D26598	HUMPSH1 Human mRNA for proteasome subunit HsC	0.021	0.061	4 consistent probesets
AI857469	wl57f04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-242	0.021	0.062	4 inconsistent probesets
M60784	Human U1 snRNP-specific protein A gene /cds=(137,985)	0.021	0.064	4 inconsistent probesets
AA877215	ob15e02.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-13	0.021	0.067	4 inconsistent probesets
L40397	Homo sapiens (clone S31i125) mRNA, 3 end of cds /cds=	0.021	0.072	4 inconsistent probesets
D16480	Homo sapiens mRNA for mitochondrial enoyl-CoA hydratas	0.021	0.072	4 consistent probesets
AF047181	Homo sapiens NADH-ubiquinone oxidoreductase subunit C	0.021	0.074	4 inconsistent probesets
X69550	H.sapiens mRNA for rho GDP-dissociation Inhibitor 1 /c	0.021	0.076	4 consistent probesets
X02761	Human mRNA for fibronectin (FN precursor) /cds=(0,6987	0.021	0.077	4 inconsistent probesets
W27939	39g3 Homo sapiens cDNA /gb=W27939 /gi=1307887 /	0.021	0.077	4 consistent probesets
M64992	Human prosomal protein P30-33K (pros-30) mRNA, comple	0.021	0.077	4 consistent probesets
L07548	Human aminoacylase-1 (ACY1) mRNA, complete cds /cds=	0.021	0.078	4 consistent probesets
J04058	Human electron transfer flavoprotein alpha-subunit mRN	0.021	0.080	4 inconsistent probesets
M16447	HUMDHPRA Human dihydropteridine reductase (hD	0.021	0.080	4 consistent probesets
X13973	Human mRNA for ribonuclease/angiogenin inhibitor (RAI)	0.021	0.081	4 consistent probesets
U67122	HSU67122 Human ubiquitin-related protein SUMO	0.021	0.082	4 consistent probesets
X70940	H.sapiens mRNA for elongation factor 1 alpha-2 /cds=(8	0.021	0.083	4 consistent probesets
M28209	HUMRAB1A Homo sapiens GTP-binding protein (RA	0.021	0.084	4 inconsistent probesets
M94630	Homo sapiens hnRNP-C like protein mRNA, complete cds /	0.021	0.085	4 consistent probesets
U70735	Homo sapiens 34 kDa Mov34 homolog mRNA, complete co	0.021	0.086	4 consistent probesets
AA420624	nc61c12.r1 Homo sapiens cDNA /clone=IMAGE-745750 /gb	0.021	0.087	8 inconsistent probesets
AF112219	Homo sapiens esterase D mRNA, complete cds /cds=(183,1	0.021	0.087	4 consistent probesets
X76488	H.sapiens mRNA for lysosomal acid lipase /cds=(145,134	0.021	0.088	4 inconsistent probesets
X92896	H.sapiens mRNA for ITBA2 protein /cds=(10,327) /gb=X92	0.021	0.092	4 consistent probesets
Y16752	Homo sapiens mRNA for secretagogin, complete CDS /cds=	0.021	0.093	4 consistent probesets
D14446	Human HFREP-1 mRNA for unknown protein, complete cds	0.021	0.093	4 inconsistent probesets
U77594	Human tazarotene-induced gene 2 (TIG2) mRNA, complete	0.021	0.095	4 consistent probesets
AI052224	oz21a02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.021	0.096	4 inconsistent probesets
X59303	Human G7a mRNA for valyl-tRNA synthetase /cds=(219,40	0.021	0.098	4 consistent probesets
M93056	Human mononcyte/neutrophil elastase inhibitor mRNA seq	0.021	0.098	4 inconsistent probesets
U44111	Human histamine N-methyltransferase (HNMT) gene /cds=(	0.021	0.101	4 inconsistent probesets
X17644	Human GST1-Hs mRNA for GTP-binding protein /cds=(648	0.021	0.104	4 consistent probesets
AF034546	Homo sapiens sorting nexin 3 (SNX3) mRNA, complete cds	0.021	0.106	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U58766	Human FX protein mRNA, complete cds /cds=(74,1039) /gb	0.021	0.107	4 consistent probesets
Z24459	H.sapiens MTCP1 gene, exons 2A to 7 (and joined mRNA)	0.021	0.108	8 consistent probesets
X76538	H.sapiens Mpv17 mRNA /cds=(29,559) /gb=X76538 /gi=434	0.021	0.110	4 inconsistent probesets
D82061	Homo sapiens mRNA for a member of the short-chain alco	0.021	0.111	4 consistent probesets
X74837	H.sapiens HUMM9 mRNA /cds=(689,2566) /gb=X74837 /gi=	0.021	0.111	4 inconsistent probesets
M12036	HUMHER2B Human tyrosine kinase-type receptor	0.021	0.112	4 inconsistent probesets
L33930	HUMCD24B Homo sapiens CD24 signal transducer	0.021	0.113	4 consistent probesets
AF004222	Homo sapiens RTN2-A (RTN2) mRNA, complete cds /cds=	0.021	0.116	4 consistent probesets
D42138	D42138 Homo sapiens mRNA for PIG-B, complete	0.021	0.116	4 consistent probesets
X76648	H.sapiens mRNA for glutaredoxin /cds=(63,383) /gb=X766	0.021	0.117	4 consistent probesets
D14697	Human mRNA for KIAA0003 gene, complete cds /cds=(114	0.021	0.121	4 consistent probesets
AL050225	Homo sapiens mRNA; cDNA DKFZp586M1523 (from clone	0.021	0.122	4 consistent probesets
Z95118	Human DNA sequence from clone 354J5 on chromosome 6	0.021	0.122	4 consistent probesets
AF049140	Homo sapiens MMS2 (MMS2) mRNA, complete cds /cds=(	0.021	0.123	4 inconsistent probesets
AI017935	ou43h10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	0.021	0.124	4 consistent probesets
U36336	Human lysosome-associated membrane protein-2b (LAMP2	0.021	0.127	4 consistent probesets
AL049786	Novel human gene mapping to chomosome 13 /cds=(370,7	0.021	0.130	4 consistent probesets
R38263	yc92c11.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-234	0.021	0.133	4 consistent probesets
D87469	Human mRNA for KIAA0279 gene, partial cds /cds=(0,7226	0.021	0.145	4 consistent probesets
U90441	Human prolyl 4-hydroxylase alpha (II) subunit mRNA, co	0.021	0.147	4 consistent probesets
X12458	Human P3 gene /cds=(494,1927) /gb=X12458 /gi=35187 /	0.021	0.147	4 consistent probesets
D31815	HUMSMP30 Human mRNA for SMP-30 (senescence ma	0.021	0.149	4 consistent probesets
AB029027	Homo sapiens mRNA for KIAA1104 protein, complete cds /	0.021	0.150	4 consistent probesets
AF035262	Homo sapiens BAF57 (BAF57) gene, complete cds /cds=(0	0.021	0.158	4 consistent probesets
U50708	Human branched chain alpha-ketoacid dehydrogenase E1 b	0.021	0.174	8 consistent probesets
L22569	Homo sapiens cathepsin B mRNA, 3 UTR with a stem-loop	0.021	0.235	4 consistent probesets
W26762	12d8 Homo sapiens cDNA /gb=W26762 /gi=1305846 /	0.021	0.238	4 consistent probesets
X15422	Human mRNA for mannose-binding protein C /cds=(65,811	0.021	0.332	4 consistent probesets
M19267	Human tropomyosin mRNA, complete cds /cds=(286,1140)	0.019	0.058	8 inconsistent probesets
L09159	Homo sapiens RHOA proto-oncogene multi-drug-resistance	0.017	0.041	4 inconsistent probesets
X67951	H.sapiens mRNA for proliferation-associated gene (pag)	0.017	0.049	4 consistent probesets
Z37166	H.sapiens BAT1 mRNA for nuclear RNA helicase (DEAD fa	0.017	0.055	4 consistent probesets
X01388	Human mRNA for pre-apolipoprotein CIII /cds=(71,370) /	0.017	0.055	4 inconsistent probesets
D13643	Human mRNA for KIAA0018 gene, complete cds /cds=(38,1	0.017	0.060	4 inconsistent probesets
AF044671	Homo sapiens MM46 mRNA, complete cds /cds=(78,431) /g	0.017	0.061	4 consistent probesets
L17131	Human high mobility group protein (HMG-I(Y)) gene exon	0.017	0.061	4 consistent probesets
M20786	Human alpha-2-plasmin inhibitor gene /cds=(4,1479) /gb	0.017	0.062	4 inconsistent probesets
M26252	Human TCB gene encoding cytosolic thyroid hormone-bind	0.017	0.063	4 inconsistent probesets
AL096714	Homo sapiens mRNA; cDNA DKFZp564E242 (from clone D	0.017	0.064	4 consistent probesets
U83246	Homo sapiens copine I mRNA, complete cds /cds=(156,176	0.017	0.067	4 consistent probesets
X79389	H.sapiens GSTT1 mRNA /cds=(0,722) /gb=X79389 /gi=510	0.017	0.067	4 consistent probesets
N98670	yy66d08.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-278	0.017	0.072	4 consistent probesets
U21090	HSU21090 Human DNA polymerase delta small sub	0.017	0.080	4 consistent probesets
X78706	H.sapiens mRNA for carnitine acetyltransferase /cds=(0	0.017	0.080	4 consistent probesets
AB017563	Homo sapiens IGSF4 gene /cds=(0,1328) /gb=AB017563 /g	0.017	0.080	4 consistent probesets
U84573	Homo sapiens lysyl hydroxylase isoform 2 (PLOD2) mRNA,	0.017	0.080	4 consistent probesets
X69433	H.sapiens mRNA for mitochondrial isocitrate dehydrogen	0.017	0.085	4 consistent probesets
AL022318	bK150C2.3 (PUTATIVE novel protein similar to APOBEC1 (	0.017	0.088	8 consistent probesets
AL050290	Homo sapiens mRNA; cDNA DKFZp586G1923 (from clone	0.017	0.088	4 consistent probesets
U55206	Homo sapiens human gamma-glutamyl hydrolase (hGH) m	0.017	0.091	4 consistent probesets
L25899	Human ribosomal protein L10 mRNA, complete cds /cds=(1	0.017	0.091	4 consistent probesets
AF052159	Homo sapiens clone 24416 mRNA sequence /cds=UNKNO	0.017	0.097	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AI570572	tm78c02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-21	0.017	0.097	3 consistent probesets
L49380	Homo sapiens clone B4 transcription factor ZFM1 mRNA,	0.017	0.097	3 inconsistent probesets
W25936	15b5 Homo sapiens cDNA /gb=W25936 /gi=1306059 /	0.017	0.098	3 inconsistent probesets
X63679	H.sapiens mRNA for TRAMP protein /cds=(121,1245) /gb=X	0.017	0.098	4 consistent probesets
AJ011779	Homo sapiens mRNA for SEC63 protein /cds=(98,2380) /gb	0.017	0.098	4 consistent probesets
S70154	cytosolic acetoacetyl-coenzyme A thiolase [human, live	0.017	0.100	4 consistent probesets
U11313	Human sterol carrier protein-X/sterol carrier protein-	0.017	0.102	4 consistent probesets
AJ001258	Homo sapiens mRNA for NIPSNAP1 protein /cds=(254,110	0.017	0.105	4 consistent probesets
AI547262	PN001_AH_H03.r Homo sapiens cDNA, 5 end /clone_end=	0.017	0.107	4 consistent probesets
D00749	HUMCD7G3 Human T cell surface antigen CD7 gen	0.017	0.111	4 consistent probesets
X85237	H.sapiens mRNA for splicing factor SF3a120 /cds=(97,24	0.017	0.118	4 consistent probesets
X82207	H.sapiens mRNA for beta-centractin (PC3) /cds=(56,1186	0.017	0.119	3 consistent probesets
AF014118	Homo sapiens membrane-associated kinase (Myt1) mRNA,	0.017	0.123	3 consistent probesets
J05257	Homo sapiens (clones MDP4, MDP7) microsomal dipeptida	0.017	0.123	3 consistent probesets
AA883502	am25h07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-14	0.017	0.125	4 inconsistent probesets
U28811	Human cysteine-rich fibroblast growth factor receptor	0.017	0.125	4 inconsistent probesets
D87989	Human mRNA for UDP-galactose transporter related isozy	0.017	0.140	3 consistent probesets
X81789	H.sapiens mRNA for splicing factor SF3a60 /cds=(565,20	0.017	0.141	4 consistent probesets
AI888563	wn33a05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.017	0.147	3 consistent probesets
M65066	HUMRIB Human cAMP-dependent protein kinase re	0.017	0.148	3 consistent probesets
U60644	Human HU-K4 mRNA, complete cds /cds=(487,1800) /gb=U	0.017	0.152	4 consistent probesets
U83659	HSU83659 Human multidrug resistance-associate	0.017	0.153	3 consistent probesets
AL050008	Homo sapiens mRNA; cDNA DKFZp564A063 (from clone D	0.017	0.156	4 consistent probesets
W28479	47d8 Homo sapiens cDNA /gb=W28479 /gi=1308427 /	0.017	0.158	3 consistent probesets
J04621	Human heparan sulfate proteoglycan (HSPG) core protein	0.017	0.158	4 consistent probesets
AF036715	Homo sapiens syntaxin 8 mRNA, complete cds /cds=(0,710	0.017	0.162	4 consistent probesets
AF035281	Homo sapiens clone 23903 mRNA sequence /cds=UNKNO	0.017	0.164	4 consistent probesets
AJ004832	Homo sapiens mRNA for neuropathy target esterase /cds=	0.017	0.165	3 consistent probesets
AF032119	Homo sapiens hCASK (CASK) mRNA, complete cds /cds=(	0.017	0.173	4 consistent probesets
L19267	Homo sapiens 59 protein mRNA, 3 end /cds=(0,1661) /gb	0.017	0.175	3 consistent probesets
U61262	Human neogenin mRNA, complete cds /cds=(136,4521) /gb	0.017	0.176	3 consistent probesets
D78611	Human MEST mRNA, complete cds /cds=(223,1230) /gb=D	0.017	0.184	4 inconsistent probesets
AB023192	Homo sapiens mRNA for KIAA0975 protein, partial cds /c	0.017	0.186	3 consistent probesets
AJ001016	Homo sapiens mRNA encoding RAMP3 /cds=(29,475) /gb=	0.017	0.190	3 consistent probesets
L17128	Homo sapiens (clone H4/H16) gamma-glutamic carboxylase	0.017	0.196	4 consistent probesets
AF072836	Homo sapiens Sox-like transcriptional factor mRNA, com	0.017	0.198	3 consistent probesets
AL031714	Human DNA sequence from clone 356B7 on chromosome 1	0.017	0.205	3 consistent probesets
AF017656	Homo sapiens G protein beta 5 subunit mRNA, complete c	0.017	0.211	4 consistent probesets
AA772359	ai43d10.s1 Homo sapiens cDNA, 3 end /clone=1359763 /c	0.017	0.228	3 consistent probesets
L07261	Human alpha adducin mRNA, partial cds including altern	0.017	0.231	3 consistent probesets
U02388	HSU02388 Human cytochrome P450 4F2 (CYP4F2) m	0.017	0.238	4 consistent probesets
AF001434	Human Hpast (HPAST) mRNA, complete cds /cds=(255,185	0.017	0.244	3 consistent probesets
AL049250	Homo sapiens mRNA; cDNA DKFZp564D113 (from clone D	0.017	0.245	3 inconsistent probesets
AL050148	Homo sapiens mRNA; cDNA DKFZp586G1520 (from clone	0.017	0.250	3 consistent probesets
Y12059	H.sapiens HUNKI mRNA /cds=(222,2390) /gb=Y12059 /gi=3	0.017	0.253	3 consistent probesets
AJ006470	Homo sapiens mRNA for cartilage-associated protein (CA	0.017	0.281	3 consistent probesets
D87464	Human mRNA for KIAA0274 gene, complete cds /cds=(124	0.017	0.284	3 consistent probesets
D90084	Human pyruvate dehydrogenase (EC 1.2.4.1) alpha subuni	0.017	0.286	4 consistent probesets
AB020637	Homo sapiens mRNA for KIAA0830 protein, partial cds /c	0.017	0.313	3 consistent probesets
M13194	HUMERCC1 Human excision repair protein (ERCC1	0.015	0.061	12 consistent probesets
M21186	Human neutrophil cytochrome b light chain p22 phagocyt	0.013	0.047	4 consistent probesets
X69392	H.sapiens mRNA for ribosomal protein L26 /cds=(6,443)	0.013	0.053	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U97105	Homo sapiens N2A3 mRNA, complete cds /cds=(1336,3054	0.013	0.057	4 consistent probesets
J02943	Human corticosteroid binding globulin mRNA, complete c	0.013	0.061	4 consistent probesets
L24521	Human transformation-related protein mRNA, 3 end /cds	0.013	0.064	4 consistent probesets
AB014560	Homo sapiens mRNA for KIAA0660 protein, complete cds /	0.013	0.064	4 consistent probesets
U25789	Human ribosomal protein L21 mRNA, complete cds /cds=(3	0.013	0.067	4 consistent probesets
X97064	H.sapiens mRNA for Sec23A isoform, 2748bp /cds=(159,24	0.013	0.067	4 consistent probesets
M37583	Human histone (H2A.Z) mRNA, complete cds /cds=(106,49	0.013	0.068	4 consistent probesets
U49897	Homo sapiens phenylalanine hydroxylase (PAH) mRNA, co	0.013	0.070	4 consistent probesets
AA845575	ak04e09.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-14	0.013	0.074	4 consistent probesets
Z97353	Human DNA sequence from clone 90L6 on chromosome 22	0.013	0.074	4 consistent probesets
X92972	H.sapiens mRNA for protein phosphatase 6 /cds=(21,938)	0.013	0.076	4 consistent probesets
AF035295	Homo sapiens clone 23623 mRNA, partial cds /cds=(0,399	0.013	0.081	4 consistent probesets
AF054184	Homo sapiens Sec61 gamma mRNA, complete cds /cds=(9	0.013	0.082	4 consistent probesets
X78669	H.sapiens ERC-55 mRNA /cds=(66,1019) /gb=X78669 /gi=4	0.013	0.084	8 inconsistent probesets
X69819	HSICAM3RN H.sapiens ICAM-3 mRNA	0.013	0.085	4 consistent probesets
AF059202	Homo sapiens ACAT related gene product 1 mRNA, comple	0.013	0.085	4 inconsistent probesets
X72964	H.sapiens mRNA for caltractin /cds=(47,565) /gb=X72964	0.013	0.087	4 consistent probesets
AL050202	Homo sapiens mRNA; cDNA DKFZp586E2023 (from clone	0.013	0.087	4 consistent probesets
AF054987	Homo sapiens clone 23831 aldolase C mRNA, complete cd	0.013	0.087	4 consistent probesets
AF079221	Homo sapiens BCL2/adenovirus E1B 19kDa-interacting pro	0.013	0.093	4 inconsistent probesets
Z47087	H.sapiens mRNA for RNA polymerase II elongation factor	0.013	0.094	4 consistent probesets
AI127424	qb75b02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-17	0.013	0.095	4 consistent probesets
D84307	Homo sapiens mRNA for phosphoethanolamine cytidylyltra	0.013	0.102	4 consistent probesets
D16827	HUMSSTR5 Human gene for fifth somatostatin re	0.013	0.105	4 consistent probesets
X97074	H.sapiens mRNS for clathrin-associated protein /cds=(1	0.013	0.107	4 consistent probesets
AI885170	wl90e10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	0.013	0.108	4 consistent probesets
AB006746	Homo sapiens hMmTRA1b mRNA, complete cds /cds=(256	0.013	0.108	4 consistent probesets
U77413	Human O-linked GlcNAc transferase mRNA, complete cds /	0.013	0.113	4 consistent probesets
U41766	Human metalloprotease/disintegrin/cysteine-rich protei	0.013	0.116	4 consistent probesets
D83200	Homo sapiens mRNA expressed in placenta /cds=UNKNOV	0.013	0.125	4 consistent probesets
D32257	Human GTF3A mRNA for Xenopus transcription factor IIIA	0.013	0.127	4 consistent probesets
AJ002190	Homo sapiens cDNA for dihydroxyacetone phosphate acylt	0.013	0.149	4 consistent probesets
AA905543	oj86h04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-150	0.013	0.179	4 consistent probesets
W28994	54h7 Homo sapiens cDNA /gb=W28994 /gi=1308960 /	0.013	0.180	4 consistent probesets
U59299	Homo sapiens putative monocarboxylate transporter MCT	0.013	0.183	4 consistent probesets
AF041474	Homo sapiens BAF53a (BAF53a) mRNA, complete cds /cds	0.012	0.126	4 consistent probesets
X07315	Human gene for PP15 (placental protein 15) /cds=(99,48	0.012	0.131	4 consistent probesets
AF052732	Homo sapiens 10-formyltetrahydrofolate dehydrogenase m	0.012	0.138	4 inconsistent probesets
L00972	Human cystathionine-beta-synthase (CBS) mRNA /cds=UN	0.012	0.180	4 consistent probesets
AF022991	Homo sapiens Rigui (RIGUI) mRNA, complete cds /cds=(18	0.011	0.120	3 consistent probesets
AF052288	untitled /cds=(0,2988) /gb=AF052288 /gi=3510692 /	0.011	0.233	3 consistent probesets
X66503	Human adenylosuccinate synthetase mRNA /cds=(0,1367) /	0.011	0.310	3 consistent probesets
X97065	H.sapiens mRNA for Sec23B isoform, 2450bp /cds=(105,24	0.011	0.341	3 consistent probesets
AI762547	wh92e05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.011	0.343	3 consistent probesets
AL080119	Homo sapiens mRNA; cDNA DKFZp564M2423 (from clone	0.010	0.056	8 inconsistent probesets
AI375033	ta66e10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-204	0.010	0.095	8 consistent probesets
M74091	HUMCLNA Human cyclin mRNA	0.010	0.156	8 inconsistent probesets
AA733050	zg79b05.s1 Homo sapiens cDNA, 3 end /clone=399537 /cl	0.010	0.126	7 consistent probesets
M22919	Human nonmuscle/smooth muscle alkali myosin light chai	0.008	0.042	4 inconsistent probesets
X70326	H.sapiens MacMarcks mRNA /cds=(13,600) /gb=X70326 /g	0.008	0.055	4 consistent probesets
X56597	Human humFib mRNA for fibrillarin /cds=(59,1024) /gb=X	0.008	0.059	4 consistent probesets
AI436567	ti03b09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-212	0.008	0.060	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D49489	Human mRNA for protein disulfide isomerase-related pro	0.008	0.061	4 consistent probesets
D87735	Homo sapiens mRNA for ribosomal protein L14, complete	0.008	0.061	4 consistent probesets
Z50022	H.sapiens mRNA for surface glycoprotein /cds=(93,635)	0.008	0.064	4 consistent probesets
U06863	Human follistatin-related protein precursor mRNA, comp	0.008	0.068	4 inconsistent probesets
AF012434	untitled /cds=(38,394) /gb=AF012434 /gi=2352914 /	0.008	0.070	4 consistent probesets
M16660	Human 90-kDa heat-shock protein gene, cDNA, complete c	0.008	0.070	4 consistent probesets
M37435	HUMCSDF1 Human macrophage-specific colony-sti	0.008	0.074	4 inconsistent probesets
Z49835	H.sapiens mRNA for protein disulfide isomerase /cds=(1	0.008	0.075	4 inconsistent probesets
U19718	Human microfibril-associated glycoprotein (MFAP2) mRNA	0.008	0.076	4 consistent probesets
X52882	Human t-complex polypeptide 1 gene /cds=(21,1691) /gb=	0.008	0.078	4 consistent probesets
U47924	Human chromosome 12p13 sequence /cds=(535,2301) /gb=	0.008	0.080	4 consistent probesets
J00068	Human adult skeletal muscle alpha-actin mRNA /cds=(103	0.008	0.080	4 inconsistent probesets
X72727	H.sapiens tunp mRNA for transformation upregulated nuc	0.008	0.080	4 consistent probesets
M69023	Human globin gene /cds=UNKNOWN /gb=M69023 /gi=183	0.008	0.081	4 consistent probesets
AF001601	Homo sapiens paraoxonase (PON2) mRNA, complete cds /	0.008	0.086	4 consistent probesets
L13977	Human prolylcarboxypeptidase mRNA, complete cds /cds=	0.008	0.088	4 consistent probesets
L40391	Homo sapiens (clone s153) mRNA fragment /cds=UNKNO	0.008	0.091	4 consistent probesets
M31303	HUMOP18A Human oncoprotein 18 (Op18) gene, co	0.008	0.091	4 inconsistent probesets
M68840	Human monoamine oxidase A (MAOA) mRNA, complete co	0.008	0.092	4 consistent probesets
AJ005940	Homo sapiens mRNA for GTP-binding protein /cds=(65,116	0.008	0.096	4 consistent probesets
AB024327	Homo sapiens pt-wd mRNA for WD-40 repeat protein, com	0.008	0.096	4 consistent probesets
L37127	HUMRPIA Homo sapiens RNA polymerase II mRNA,	0.008	0.100	4 consistent probesets
U70987	HSU70987 Human GAP binding protein p62dok (DO	0.008	0.101	2 consistent probesets
U37100	Homo sapiens aldose reductase-like peptide mRNA, compl	0.008	0.104	4 inconsistent probesets
M37721	Human peptidylglycine alpha-amidating monooxygenase m	0.008	0.104	4 consistent probesets
M30474	Human kidney gamma-glutamyl transpeptidase type II mRN	0.008	0.105	4 consistent probesets
X01630	Human mRNA for argininosuccinate synthetase /cds=(75,1	0.008	0.107	4 consistent probesets
Z48501	H.sapiens mRNA for polyadenylate binding protein II /c	0.008	0.111	4 inconsistent probesets
L07594	HUMTGFB3C Human transforming growth factor-be	0.008	0.118	4 inconsistent probesets
U05040	Human FUSE binding protein mRNA, complete cds /cds=(2	0.008	0.119	4 consistent probesets
U02556	Human RP3 mRNA, complete cds /cds=(68,418) /gb=U025	0.008	0.121	4 consistent probesets
U31525	Human glycogenin mRNA, complete cds /cds=(127,1128) /g	0.008	0.122	4 consistent probesets
W28190	43c6 Homo sapiens cDNA /gb=W28190 /gi=1308138 /	0.008	0.124	4 consistent probesets
Z30093	H.sapiens mRNA for basic transcription factor 2, 34 kD	0.008	0.130	4 inconsistent probesets
AA913812	ol39a08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-152	0.008	0.131	4 consistent probesets
U72882	HSU72882 Human interferon-induced leucine zip	0.008	0.132	4 consistent probesets
AF089816	Homo sapiens RGS-GAIP interacting protein GIPC mRNA,	0.008	0.133	4 consistent probesets
AB028449	Homo sapiens mRNA for Helicase-MOI, complete cds /cds=	0.008	0.135	4 consistent probesets
AF012281	Homo sapiens PDZ domain containing-protein (PDZK1) mR	0.008	0.145	4 consistent probesets
AF106861	Homo sapiens attractin-2 (ATRN) mRNA, complete cds /cd	0.008	0.148	4 consistent probesets
D10040	Homo sapiens mRNA for long-chain acyl-CoA synthetase,	0.008	0.148	4 consistent probesets
N25122	yx19d10.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-262	0.008	0.150	4 consistent probesets
AF093771	Homo sapiens mitoxantrone resistance protein 1 mRNA, p	0.008	0.159	4 consistent probesets
AF005392	Homo sapiens alpha tubulin (TUBA2) gene, partial cds /	0.008	0.172	4 inconsistent probesets
AB005289	Homo sapiens mRNA for ABC transporter 7 protein, compl	0.008	0.175	4 consistent probesets
U80744	Homo sapiens CTG4a mRNA, complete cds /cds=(387,818	0.008	0.178	4 consistent probesets
AF069765	Homo sapiens signal recognition particle 72 (SRP72) mR	0.008	0.207	4 consistent probesets
U59185	Human putative monocarboxylate transporter (MCT) mRNA	0.008	0.207	4 consistent probesets
AF070606	Homo sapiens clone 24411 mRNA sequence /cds=UNKNO	0.008	0.211	4 consistent probesets
X00351	Human mRNA for beta-actin (_5, _M, _	0.007	0.043	16 inconsistent probesets
M65254	HUMP2B Protein phosphatase 2A 65 kDa regulato	0.006	0.083	11 inconsistent probesets
D38047	HUMPSP31 Human mRNA for 26S proteasome subuni	0.005	0.104	7 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF023612	Homo sapiens Dim1p homolog mRNA, complete cds /cds=(	0.005	0.145	7 consistent probesets
X55954	Human mRNA for HL23 ribosomal protein homologue /cds=	0.004	0.050	8 inconsistent probesets
J03826	Human adrenodoxin reductase mRNA, complete cds /cds=(	0.004	0.051	4 inconsistent probesets
X53777	Human L23 mRNA for putative ribosomal protein /cds=(13	0.004	0.058	4 consistent probesets
U37230	Human ribosomal protein L23a mRNA, complete cds /cds=(	0.004	0.062	4 consistent probesets
M86400	HUMPHPLA2 Human phospholipase A2 mRNA, comple	0.004	0.064	4 consistent probesets
AF010313	Homo sapiens Pig8 (PIG8) mRNA, complete cds /cds=(72,1	0.004	0.064	4 inconsistent probesets
M34539	HUMFKBP Human FK506-binding protein (FKBP) mR	0.004	0.064	4 inconsistent probesets
M86667	HUMNAP H.sapiens NAP (nucleosome assembly pro	0.004	0.067	8 inconsistent probesets
AF038958	Homo sapiens synaptic glycoprotein SC2 spliced variant	0.004	0.068	4 consistent probesets
M36341	Human ADP-ribosylation factor 4 (ARF4) mRNA, complete	0.004	0.069	4 inconsistent probesets
L29277	HUMAPRF Homo sapiens DNA-binding protein (APR	0.004	0.070	8 inconsistent probesets
M29065	Human hnRNP A2 protein mRNA /cds=(155,1180) /gb=M29	0.004	0.071	4 inconsistent probesets
AL120815	DKFZp762F172_r1 Homo sapiens cDNA, 5 end /clone=DK	0.004	0.071	4 consistent probesets
U50523	HSU50523 Human BRCA2 region, mRNA sequence CG	0.004	0.074	4 consistent probesets
U10860	Human guanosine 5-monophosphate synthase mRNA, com	0.004	0.078	4 consistent probesets
U49188	Human placenta (Diff33) mRNA, complete cds /cds=(106,1	0.004	0.081	4 consistent probesets
AF081281	Homo sapiens lysophospholipase (LPL1) mRNA, complete (	0.004	0.082	4 consistent probesets
AD001528	Homo sapiens spermidine aminopropyltransferase mRNA, c	0.004	0.085	4 consistent probesets
M81600	Human NAD(P)H-quinone oxireductase gene /cds=(111,935	0.004	0.085	4 consistent probesets
L25081	HUMRHOCA Homo sapiens GTPase (rhoC) mRNA, com	0.004	0.085	4 inconsistent probesets
D17530	Homo sapiens mRNA for drebrin E, complete cds /cds=(97	0.004	0.091	4 consistent probesets
U61837	Homo sapiens putative cyclin G1 interacting protein mR	0.004	0.093	4 consistent probesets
M90357	Human basic transcription factor 3a (BTF3a) gene /cds=	0.004	0.093	4 consistent probesets
L29254	Human (clone P1-5) L-iditol-2 dehydrogenase gene /cds=	0.004	0.094	4 consistent probesets
X04828	Human mRNA for G(i) protein alpha-subunit (adenylate c	0.004	0.095	4 consistent probesets
M69040	Human SF2p33 mRNA, complete cds /cds=(124,870) /gb=M	0.004	0.097	4 consistent probesets
S72869	H4(D10S170)=putative cytoskeletal protein [human, thyr	0.004	0.097	4 inconsistent probesets
Y00318	Human mRNA for complement control protein factor I /cd	0.004	0.099	4 consistent probesets
AI360249	qy84f07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-201	0.004	0.101	4 consistent probesets
AF060902	Homo sapiens vesicle soluble NSF attachment protein re	0.004	0.110	4 consistent probesets
U64520	Human synaptobrevin-3 mRNA, complete cds /cds=(24,326	0.004	0.113	4 consistent probesets
U50079	HSU50079 Human histone deacetylase HD1 mRNA,	0.004	0.115	4 consistent probesets
AL080080	Homo sapiens mRNA; cDNA DKFZp564E1962 (from clone	0.004	0.116	4 consistent probesets
W28255	44b8 Homo sapiens cDNA /gb=W28255 /gi=1308203 /	0.004	0.118	4 consistent probesets
U83411	Homo sapiens carboxypeptidase Z precursor, mRNA, comp	0.004	0.124	4 consistent probesets
W22110	64F11 Homo sapiens cDNA /clone=(not-directional) /gb=W	0.004	0.129	4 inconsistent probesets
AI867349	wi24g10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-239	0.004	0.134	4 consistent probesets
AB028953	Homo sapiens mRNA for KIAA1030 protein, partial cds /c	0.004	0.136	4 consistent probesets
X60592	Human CDw40 mRNA for nerve growth factor receptor-rela	0.004	0.136	4 consistent probesets
AB006629	Homo sapiens mRNA for KIAA0291 gene, partial cds /cds=	0.004	0.157	4 consistent probesets
AW026656	wv15c06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-99	0.004	0.162	4 inconsistent probesets
M14648	HUMVTNR Human cell adhesion protein (vitronec	0.004	0.163	8 consistent probesets
AJ010071	Homo sapiens for TOM1-like protein /cds=(30,1460) /gb=	0.004	0.184	4 consistent probesets
AA808961	nw16h03.s1 Homo sapiens cDNA /clone=IMAGE-1240661 /	0.004	0.186	4 consistent probesets
AF061258	Homo sapiens LIM protein mRNA, complete cds /cds=(83,1	0.004	0.205	4 consistent probesets
AF022813	Homo sapiens tetraspan (NAG-2) mRNA, complete cds /cds	0.004	0.217	4 consistent probesets
U92980	Homo sapiens clone DT1P1A10 mRNA, CAG repeat region	0.004	0.316	4 consistent probesets
AI692348	wd85g12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	0.004	0.388	4 consistent probesets
AI526079	DU3.2-7.G09 Homo sapiens cDNA, 3 end /clone_end=3 /g	0.000	0.039	4 inconsistent probesets
X80822	H.sapiens mRNA for ORF /cds=(133,555) /gb=X80822 /gi=5	0.000	0.041	4 consistent probesets
AI033692	ow26f02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-164	0.000	0.044	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D78134	Homo sapiens mRNA for CIRP, complete cds /cds=(80,598	0.000	0.050	4 consistent probesets
D29643	Human mRNA for KIAA0115 gene, complete cds /cds=(106	0.000	0.055	4 consistent probesets
X81817	H.sapiens BAP31 mRNA /cds=(73,813) /gb=X81817 /gi=55(	0.000	0.061	4 consistent probesets
X17620	HSNM23 Human mRNA for Nm23 protein, involved	0.000	0.062	4 inconsistent probesets
X69391	H.sapiens mRNA for ribosomal protein L6 /cds=(26,892)	0.000	0.067	4 inconsistent probesets
AL050161	Homo sapiens mRNA; cDNA DKFZp586B0222 (from clone	0.000	0.068	4 consistent probesets
AF034544	Homo sapiens delta7-sterol reductase mRNA, complete cd	0.000	0.068	4 inconsistent probesets
AA206524	zq58b03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-645	0.000	0.077	4 consistent probesets
D31885	Human mRNA for KIAA0069 gene, partial cds /cds=(0,680)	0.000	0.078	4 consistent probesets
AA845349	ak01g01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-14(	0.000	0.082	4 consistent probesets
U24105	Homo sapiens coatomer protein (COPA) mRNA, complete (	0.000	0.082	4 consistent probesets
X71490	H.sapiens mRNA for vacuolar proton ATPase, subunit D /	0.000	0.091	4 consistent probesets
AF006087	Homo sapiens Arp2/3 protein complex subunit p20-Arc (A	0.000	0.093	8 inconsistent probesets
U76010	Human putative zinc transporter ZnT-3 (ZnT-3) mRNA, co	0.000	0.095	4 consistent probesets
W25986	17e7 Homo sapiens cDNA /gb=W25986 /gi=1306253 /	0.000	0.096	4 consistent probesets
D26068	Human mRNA for KIAA0038 gene, partial cds /cds=(0,694)	0.000	0.096	4 inconsistent probesets
AB018269	Homo sapiens mRNA for KIAA0726 protein, complete cds /	0.000	0.098	3 consistent probesets
L36983	Homo sapiens dynamin (DNM) mRNA, complete cds /cds=(	0.000	0.098	3 inconsistent probesets
AF000652	Homo sapiens syntenin (sycl) mRNA, complete cds /cds=(	0.000	0.098	4 consistent probesets
U10323	Human nuclear factor NF45 mRNA, complete cds /cds=(39,	0.000	0.098	4 consistent probesets
D87012	Human (lambda) DNA for immunoglobin light chain /cds=(	0.000	0.101	1 consistent probesets
X66364	HSSTHPKE H.sapiens mRNA PSSALRE for serine/th	0.000	0.101	1 consistent probesets
D42123	Homo sapiens mRNA for ESP1/CRP2, complete cds /cds=(	0.000	0.101	3 consistent probesets
L41143	Homo sapiens expressed pseudo TCTA mRNA at t(1;3) tran	0.000	0.101	1 consistent probesets
J05243	Human nonerythroid alpha-spectrin (SPTAN1) mRNA, comp	0.000	0.101	4 inconsistent probesets
L11285	HUMMEK2NF Homosapiens ERK activator kinase (M	0.000	0.103	4 consistent probesets
Z26317	H.sapiens mRNA for desmoglein 2 /cds=(11,3364) /gb=Z26	0.000	0.111	3 inconsistent probesets
AF007140	Homo sapiens clone 23711 unknown mRNA, partial cds /cd	0.000	0.112	4 consistent probesets
L41162	Homo sapiens collagen alpha 3 type IX (COL9A3) mRNA, c	0.000	0.113	4 consistent probesets
X83928	HSTAFII28 H.sapiens mRNA for transcription fa	0.000	0.114	8 inconsistent probesets
AF000573	Homo sapiens homogentisate 1,2-dioxygenase gene, comp	0.000	0.118	4 consistent probesets
AA010078	ze16d01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-35(	0.000	0.122	4 consistent probesets
X76732	H.sapiens mRNA for NEFA protein /cds=(219,1481) /gb=X7	0.000	0.129	4 consistent probesets
AI761818	wi62g02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23(	0.000	0.131	3 consistent probesets
U08998	Human TAR RNA binding protein 2 (TRBP2) mRNA, compl	0.000	0.135	3 consistent probesets
X54637	HSTYK2 Human tyk2 mRNA for non-receptor prote	0.000	0.139	3 consistent probesets
U17969	HSU17969 Human initiation factor eIF-5A gene,	0.000	0.143	2 consistent probesets
U75362	Homo sapiens isopeptidase T-3 (ISOT-3) mRNA, complete	0.000	0.143	2 consistent probesets
AF091092	Homo sapiens clone 683 unknown mRNA, complete sequen	0.000	0.143	4 consistent probesets
AL049654	Novel human mRNA similar to mouse gene PICK1 (TR-Q62	0.000	0.143	2 consistent probesets
M16941	Human MHC class II HLA-DR7-associated glycoprotein bet	0.000	0.147	4 inconsistent probesets
X96484	H.sapiens mRNA for DGCR6 protein /cds=(422,676) /gb=X9	0.000	0.151	1 consistent probesets
Z29481	H.sapiens mRNA for 3-hydroxyanthranilic acid dioxygena	0.000	0.151	1 consistent probesets
U94592	Human uncoupling protein homolog (UCPH) mRNA, comple	0.000	0.151	1 consistent probesets
L78207	Homo sapiens sulfonylurea receptor (SUR1) mRNA, comple	0.000	0.154	4 consistent probesets
AF069517	Homo sapiens RNA binding protein DEF-3 mRNA, complete	0.000	0.156	8 consistent probesets
X95762	H.sapiens mRNA for aminopeptidase P-like /cds=(0,1871)	0.000	0.161	2 consistent probesets
AB021981	Homo sapiens mRNA for UDP-N-acetylglucosamine transpo	0.000	0.177	4 consistent probesets
AA586695	nn42h06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-10	0.000	0.178	2 consistent probesets
AF070531	Homo sapiens clone 24764 mRNA sequence /cds=UNKNO	0.000	0.179	3 consistent probesets
X07228	Human mRNA for hepatic triglyceride lipase (HTGL) /cds	0.000	0.187	4 consistent probesets
AI189226	qd04h11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-17	0.000	0.191	2 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
N74607	za55a01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-29	0.000	0.194	3 consistent probesets
AF093119	Homo sapiens UPH1 (UPH1) mRNA, complete cds /cds=(0	0.000	0.194	3 consistent probesets
AB011124	Homo sapiens mRNA for KIAA0552 protein, complete cds /	0.000	0.199	4 consistent probesets
AL050376	Homo sapiens mRNA; cDNA DKFZp586J101 (from clone D	0.000	0.202	1 consistent probesets
AB007455	Homo sapiens mRNA for P53TG1-A, complete cds /cds=(1	0.000	0.202	1 consistent probesets
AF026977	Homo sapiens microsomal glutathione S-transferase 3 (M	0.000	0.202	1 consistent probesets
AB023151	Homo sapiens mRNA for KIAA0934 protein, partial cds /c	0.000	0.202	1 consistent probesets
U46193	Human renal cell carcinoma antigen RAGE-3 mRNA, comp	0.000	0.202	1 consistent probesets
D13640	Human mRNA for KIAA0015 gene, complete cds /cds=(106	0.000	0.202	1 consistent probesets
AF055027	Homo sapiens clone 24658 mRNA sequence /cds=UNKNO	0.000	0.203	3 inconsistent probesets
M14218	Human argininosuccinate lyase mRNA, complete cds /cds=	0.000	0.203	2 consistent probesets
U18671	HSU18671 Human Stat2 gene, complete cds"	0.000	0.208	3 consistent probesets
M54927	Human myelin proteolipid protein mRNA, complete cds /c	0.000	0.214	2 consistent probesets
AL031387	dJ596C15.1.1 (novel protein) (isoform 1) /cds=(0,488)	0.000	0.214	2 consistent probesets
U91543	Homo sapiens zinc-finger helicase (hZFH) mRNA, complet	0.000	0.214	2 consistent probesets
D49357	Human mRNA for S-adenosylmethionine synthetase, comp	0.000	0.215	2 consistent probesets
U63289	Human RNA-binding protein CUG-BP/hNab50 (NAB50) mR	0.000	0.215	2 inconsistent probesets
AF087020	Homo sapiens protein zero related protein (PZR) mRNA,	0.000	0.221	3 inconsistent probesets
AF031469	Homo sapiens MHC class I related protein 1 isoform D (	0.000	0.225	3 consistent probesets
AL049370	Homo sapiens mRNA; cDNA DKFZp586D0918 (from clone	0.000	0.226	3 consistent probesets
W26851	17b12 Homo sapiens cDNA /gb=W26851 /gi=1306214 /	0.000	0.231	3 consistent probesets
M81590	Homo sapiens serotonin 1D receptor (5-HT1D~) mRNA, co	0.000	0.235	1 consistent probesets
U46746	Human dystrobrevin-epsilon mRNA, complete cds /cds=(33	0.000	0.238	2 consistent probesets
AL049415	Homo sapiens mRNA; cDNA DKFZp586N2119 (from clone	0.000	0.252	1 consistent probesets
X53414	Human mRNA for peroxisomal L-alanine-glyoxylate aminot	0.000	0.252	1 consistent probesets
K01383	Human metallothionein-I-A gene, complete coding sequen	0.000	0.252	1 consistent probesets
AF087693	Homo sapiens veli 1 mRNA, complete cds /cds=(162,863)	0.000	0.259	2 consistent probesets
X15218	HSSKIR Human ski oncogene mRNA	0.000	0.292	3 consistent probesets
AF007876	Homo sapiens Na,K-ATPase beta 2 subunit gene, complete	0.000	0.293	1 consistent probesets
U56421	Human olfactory receptor (OLF3) gene, complete cds /cd	0.000	0.293	1 consistent probesets
AF000421	Homo sapiens TTF-I interacting peptide 12 mRNA, partia	0.000	0.297	3 consistent probesets
AF022913	Homo sapiens GPI transamidase mRNA, complete cds /cds	0.000	0.302	1 consistent probesets
AL050405	Novel human gene mapping to chomosome X /cds=(39,100	0.000	0.302	1 consistent probesets
AJ000414	Homo sapiens mRNA for Cdc42-interacting protein 4 (CIP	0.000	0.302	1 consistent probesets
X13589	HSAROMAT Human mRNA for aromatase (estrogen s	0.000	0.302	1 consistent probesets
L37360	HUMEFL2 Homo sapiens (clone hEHK1-L) EHK1 rec	0.000	0.302	1 consistent probesets
Y00970	Human mRNA for acrosin (EC 3.4.21.10) /cds=(16,1281) /	0.000	0.302	1 consistent probesets
L20965	HUMPDEA Human phosphodiesterase mRNA, complet	0.000	0.302	1 consistent probesets
X92110	H.sapiens mRNA for hcgVIII protein /cds=UNKNOWN /gb=)	0.000	0.302	1 consistent probesets
W72733	zd77h11.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-346	0.000	0.302	1 consistent probesets
U32324	HSU32324 Human interleukin-11 receptor alpha	0.000	0.302	1 consistent probesets
AL080202	Homo sapiens mRNA; cDNA DKFZp434F172 (from clone D	0.000	0.302	1 consistent probesets
D14664	Human mRNA for KIAA0022 gene, complete cds /cds=(184	0.000	0.304	2 consistent probesets
D63390	Homo sapiens mRNA for acetylhydrolase IB beta-subunit,	0.000	0.312	3 consistent probesets
AB007972	Homo sapiens mRNA, chromosome 1 specific transcript KI	0.000	0.330	3 consistent probesets
U10868	Human aldehyde dehydrogenase ALDH7 mRNA, complete	0.000	0.350	2 consistent probesets
X80910	H.sapiens PPP1CB mRNA /cds=(258,1241) /gb=X80910 /g	0.000	0.353	1 consistent probesets
L14754	Human DNA-binding protein (SMBP2) mRNA, complete cds	0.000	0.353	1 consistent probesets
AB000712	Homo sapiens hCPE-R mRNA for CPE-receptor, complete	0.000	0.353	1 consistent probesets
D50030	Homo sapiens gene for hepatocyte growth factor activat	0.000	0.353	1 consistent probesets
M29386	HUMPRLA Human prolactin mRNA, 3 end"	0.000	0.353	1 consistent probesets
X07948	Human mRNA for transition protein 1 (TP1) /cds=(58,225	0.000	0.353	1 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
D84239	Homo sapiens mRNA for IgG Fc binding protein, complete	0.000	0.356	2 consistent probesets
L38951	Homo sapiens importin beta subunit mRNA, complete cds	0.000	0.360	3 consistent probesets
AB023152	Homo sapiens mRNA for KIAA0935 protein, partial cds /c	0.000	0.395	3 consistent probesets
M94345	Homo sapiens macrophage capping protein mRNA, comple	0.000	0.403	1 consistent probesets
L28957	Homo sapiens CTP-phosphocholine cytidyltransferase mRN	0.000	0.403	1 consistent probesets
AB014534	Homo sapiens mRNA for KIAA0634 protein, partial cds /c	0.000	0.403	1 consistent probesets
D26158	Homo sapiens mRNA for PLE21 protein, complete cds /cds	0.000	0.454	1 consistent probesets
U48213	Human D-site binding protein gene, promoter region and	0.000	0.454	1 consistent probesets
AF093239	Homo sapiens MRS1 mRNA, complete cds /cds=(348,1616	0.000	0.504	1 consistent probesets
AI337192	qx88h10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-20	0.000	0.504	1 consistent probesets
J03910	HUMMT2A Human (clone 14VS) metallothionein-IG	0.000	0.504	1 consistent probesets
X13403	Human mRNA for octamer-binding protein Oct-1 /cds=(59,	0.000	0.504	1 consistent probesets
AL031666	Human DNA sequence from clone 569M23 on chromosome	0.000	0.510	2 consistent probesets
AF041853	Homo sapiens kinesin family member protein KIF3A mRNA	0.000	0.554	1 consistent probesets
M31659	Human GT mitochondrial solute carrier protein homologu	0.000	0.554	1 consistent probesets
Y15228	Homo sapiens mRNA for leukemia associated gene 2 /cds=	0.000	0.605	1 consistent probesets
AL080133	Homo sapiens mRNA; cDNA DKFZp434G173 (from clone D	0.000	0.605	1 consistent probesets
X66087	HSAMYB2 H.sapiens a-myb mRNA /NOTE=replacemen	0.000	0.642	2 consistent probesets
AF064801	Homo sapiens multiple membrane spanning receptor TRC8	0.000	0.756	1 consistent probesets
D87074	Human mRNA for KIAA0237 gene, complete cds /cds=(475	0.000	0.958	1 consistent probesets
M61832	Human S-adenosylhomocysteine hydrolase (AHCY) mRNA	-0.004	0.053	4 consistent probesets
M16961	Human alpha-2-HS-glycoprotein alpha and beta chain mRN	-0.004	0.056	4 inconsistent probesets
D45248	HUMPHPA28A Human mRNA for proteasome activato	-0.004	0.056	8 consistent probesets
U10324	Human nuclear factor NF90 mRNA, complete cds /cds=(264	-0.004	0.057	8 inconsistent probesets
S85655	prohibitin [human, mRNA, 1043 nt] /cds=(50,868) /gb=S8	-0.004	0.065	4 inconsistent probesets
L36055	Human 4E-binding protein 1 mRNA, complete cds /cds=(0,	-0.004	0.067	4 consistent probesets
AF059203	Homo sapiens acyl coenzyme A-cholesterol acyltransfera	-0.004	0.069	4 consistent probesets
X12671	Human gene for heterogeneous nuclear ribonucleoprotein	-0.004	0.070	4 consistent probesets
U37055	HSU37055 Human hepatocyte growth factor-like	-0.004	0.070	4 consistent probesets
U49785	Human D-dopachrome tautomerase mRNA, complete cds /	-0.004	0.073	4 inconsistent probesets
M64673	HUMHSF1 Human heat shock factor 1 (TCF5) mRNA	-0.004	0.074	8 inconsistent probesets
X77794	HSCYCG1 H.sapiens mRNA for cyclin G1	-0.004	0.084	4 inconsistent probesets
M21259	Human Alu repeats in the region 5 to the small nuclear	-0.004	0.085	4 consistent probesets
J02683	Human ADP/ATP carrier protein mRNA, complete cds /cds=	-0.004	0.090	4 consistent probesets
D13988	HUMRABGDI Human rab GDI mRNA, complete cds"	-0.004	0.097	4 inconsistent probesets
Z35093	H.sapiens mRNA for SURF-1 /cds=(14,916) /gb=Z35093 /gi	-0.004	0.098	4 inconsistent probesets
AF054825	Homo sapiens VAMP5 mRNA, complete cds /cds=(57,407)	-0.004	0.098	4 consistent probesets
D86062	Human mRNA for KNP-Ib, complete cds /cds=(18,731) /gb=	-0.004	0.101	4 consistent probesets
M33519	Human HLA-B-associated transcript 3 (BAT3) mRNA, comp	-0.004	0.102	4 inconsistent probesets
M28211	HUMRAB4A Homo sapiens GTP-binding protein (RA	-0.004	0.103	8 consistent probesets
W28869	53h2 Homo sapiens cDNA /gb=W28869 /gi=1308880 /	-0.004	0.105	4 consistent probesets
D16581	Human mRNA for 8-oxo-dGTPase, complete cds /cds=(26,4	-0.004	0.108	4 consistent probesets
X04366	Human mRNA for calcium activated neutral protease larg	-0.004	0.115	4 consistent probesets
X98654	H.sapiens mRNA for DRES9 protein /cds=(189,3923) /gb=X	-0.004	0.140	4 consistent probesets
AA978353	oq40b07.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-15	-0.004	0.143	4 consistent probesets
AI806379	wf27b10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	-0.004	0.145	4 consistent probesets
W28170	43a12 Homo sapiens cDNA /gb=W28170 /gi=1308118 /	-0.004	0.185	4 inconsistent probesets
U28424	Human protein kinase inhibitor p58 mRNA, complete cds	-0.006	0.198	3 consistent probesets
S66213	integrin alpha 6B [human, mRNA Partial, 528 nt] /cds=(	-0.006	0.104	8 inconsistent probesets
AF053356	Homo sapiens chromosome 7q22 sequence /cds=(253,1275	-0.007	0.081	7 inconsistent probesets
U01337	HSU01337 Human Ser/Thr protein kinase (A-RAF-	-0.008	0.048	8 inconsistent probesets
M85234	Human nuclease sensitive element binding protein-1 mRN	-0.008	0.066	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
L76517	HUMPS1MRNA Homo sapiens (clone cc44) senilin	-0.008	0.067	4 consistent probesets
M69039	Human pre-mRNA splicing factor SF2p32, complete sequer	-0.008	0.067	4 inconsistent probesets
Z82022	H.sapiens mRNA for GlcNac-1-P transferase /cds=(103,13	-0.008	0.067	4 consistent probesets
AB019408	Homo sapiens mRNA, expressed in fibroblasts of periodo	-0.008	0.075	4 consistent probesets
AL049324	Homo sapiens mRNA; cDNA DKFZp564D246 (from clone D	-0.008	0.077	4 consistent probesets
AL050383	Homo sapiens mRNA; cDNA DKFZp547E0510 (from clone	-0.008	0.080	4 inconsistent probesets
U44839	HSU44839 Human putative ubiquitin C-terminal	-0.008	0.082	4 consistent probesets
AF030249	Homo sapiens putative dienoyl-CoA isomerase (ECH1) gen	-0.008	0.083	4 consistent probesets
S80071	hPROT=brain-specific L-proline transporter [human, hip	-0.008	0.083	4 consistent probesets
AF095154	Homo sapiens C1q-related factor mRNA, complete cds /cd	-0.008	0.084	4 consistent probesets
U37221	Human cyclophilin-like protein mRNA, partial cds /cds=	-0.008	0.085	4 inconsistent probesets
U47025	Human fetal brain glycogen phosphorylase B mRNA, compl	-0.008	0.089	4 consistent probesets
U24166	Homo sapiens EB1 mRNA, complete cds /cds=(64,870) /gb	-0.008	0.091	4 inconsistent probesets
AI557272	PT2.1_15_G02.r Homo sapiens cDNA, 3 end /clone_end=3	-0.008	0.094	4 consistent probesets
X83218	H.sapiens mRNA for ATP synthase /cds=(36,677) /gb=X832	-0.008	0.094	4 consistent probesets
Y08915	H.sapiens mRNA for alpha 4 protein /cds=(8,1027) /gb=Y	-0.008	0.098	4 consistent probesets
D50914	Human mRNA for KIAA0124 gene, partial cds /cds=(0,2048	-0.008	0.102	4 consistent probesets
AB014584	Homo sapiens mRNA for KIAA0684 protein, partial cds /c	-0.008	0.104	4 consistent probesets
W28732	50h7 Homo sapiens cDNA /gb=W28732 /gi=1308680 /	-0.008	0.108	4 consistent probesets
L47738	Homo sapiens inducible protein mRNA, complete cds /cds	-0.008	0.111	4 inconsistent probesets
L07540	HUMPOLACCA Human replication factor C, 36-kDa	-0.008	0.114	8 inconsistent probesets
X77196	H.sapiens mRNA for lysosome-associated membrane prote	-0.008	0.122	4 consistent probesets
R93527	yq35f10.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-197	-0.008	0.125	4 consistent probesets
AB002450	Homo sapiens mRNA from chromosome 5q21-22, clone-A3	-0.008	0.128	4 inconsistent probesets
S74728	antiquitin=26g turgor protein homolog [human, kidney,	-0.008	0.132	4 consistent probesets
AB011086	Homo sapiens mRNA for KIAA0514 protein, complete cds /	-0.008	0.132	4 consistent probesets
L41668	Homo sapiens UDP-galactose-4-epimerase (GALE) mRNA,	-0.008	0.132	4 consistent probesets
D63780	Homo sapiens mRNA for YSK1, complete cds /cds=(114,13	-0.008	0.137	4 consistent probesets
U94319	Human autoantigen DFS70 mRNA, partial cds /cds=(0,1055	-0.008	0.137	4 consistent probesets
L49229	HUMRB1AADL Homo sapiens retinoblastoma suscep	-0.008	0.181	4 inconsistent probesets
U72263	Human multiple exostoses type II protein EXT2.I mRNA,	-0.008	0.188	4 consistent probesets
AB000095	Homo sapiens mRNA for hepatocyte growth factor activat	-0.008	0.189	4 consistent probesets
AF052167	Homo sapiens clone 24749 and 24750 mRNA sequences /c	-0.008	0.196	4 consistent probesets
AA524802	nh33h11.s1 Homo sapiens cDNA /clone=IMAGE-954213 /g	-0.008	0.242	2 consistent probesets
U09284	Human PINCH protein mRNA, complete cds /cds=(119,106	-0.008	0.244	4 consistent probesets
D10511	Homo sapiens MAT gene for mitochondrial acetoacetyl-Co	-0.008	0.295	2 consistent probesets
U79274	Human clone 23733 mRNA, complete cds /cds=(416,1237)	-0.008	0.317	4 consistent probesets
AF000430	AF000430 Homo sapiens dynamin-like protein mR	-0.010	0.154	7 consistent probesets
AB020685	Homo sapiens mRNA for KIAA0878 protein, complete cds /	-0.010	0.259	7 consistent probesets
U12022	Human calmodulin (CALM1) gene /cds=(199,648) /gb=U12(	-0.012	0.119	7 consistent probesets
AA675900	g02504r Homo sapiens cDNA, 5 end /clone=g02504 /clone	-0.012	0.091	8 inconsistent probesets
AI961669	wt65e11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	-0.012	0.098	4 inconsistent probesets
L20433	Human octamer binding transcription factor 1 (OTF1) mR	-0.012	0.100	4 inconsistent probesets
AB028952	Homo sapiens mRNA for KIAA1029 protein, complete cds /	-0.012	0.143	4 consistent probesets
U40490	Human nicotinamide nucleotide transhydrogenase mRNA, r	-0.012	0.149	4 inconsistent probesets
AF084523	Homo sapiens cellular repressor of E1A-stimulated gene	-0.012	0.201	4 consistent probesets
U01244	Human fibulin-1D mRNA, complete cds /cds=(10,2121) /gb	-0.012	0.203	4 inconsistent probesets
M34181	Human testis-specific cAMP-dependent protein kinase ca	-0.012	0.318	4 consistent probesets
D12765	HUME1AF Human mRNA for E1A-F	-0.013	0.050	4 inconsistent probesets
X59417	H.sapiens PROS-27 mRNA /cds=(62,802) /gb=X59417 /gi=	-0.013	0.060	4 consistent probesets
Y00281	Human mRNA for ribophorin I /cds=(137,1960) /gb=Y00281	-0.013	0.075	4 inconsistent probesets
AI346580	qp51f08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-192	-0.013	0.076	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF053070	Homo sapiens NADH-ubiquinone dehydrogenase 51 kDa su	-0.013	0.086	4 consistent probesets
M95627	Homo sapiens angio-associated migratory cell protein (	-0.013	0.095	4 consistent probesets
S82470	S82470 BB1=malignant cell expression-enhanced	-0.013	0.102	4 consistent probesets
AL120559	DKFZp761B219_r1 Homo sapiens cDNA, 5 end /clone=DK	-0.013	0.111	4 inconsistent probesets
Y10659	HSIL13RA H.sapiens IL-13Ra mRNA	-0.013	0.114	4 consistent probesets
M90516	Human glutamine-fructose-6-phosphate amidotransferase	-0.013	0.115	4 inconsistent probesets
M75106	Human prepro-plasma carboxypeptidase B mRNA, complet	-0.013	0.121	4 consistent probesets
U41387	Human Gu protein mRNA, partial cds /cds=(0,2405) /gb=U	-0.013	0.127	4 consistent probesets
W28610	49b12 Homo sapiens cDNA /gb=W28610 /gi=1308558 /	-0.013	0.162	4 consistent probesets
AF064093	Homo sapiens KE04p mRNA, complete cds /cds=(89,1129)	-0.013	0.167	4 consistent probesets
U14187	Human receptor tyrosine kinase ligand LERK-3 (EPLG3) m	-0.013	0.175	4 consistent probesets
L14595	Human alanine/serine/cysteine/threonine transporter (A	-0.013	0.204	4 consistent probesets
J05682	Human subunit C of V-ATPase (vat C) mRNA, 3 end /cds=	-0.013	0.204	4 consistent probesets
M92642	Homo sapiens alpha-1 type XVI collagen (COL16A1) mRNA	-0.013	0.262	4 consistent probesets
W28616	49b9 Homo sapiens cDNA /gb=W28616 /gi=1308564 /	-0.015	0.095	8 inconsistent probesets
L12535	Human RSU-1/RSP-1 mRNA, complete cds /cds=(827,166	-0.015	0.101	8 consistent probesets
AF004430	Homo sapiens hD54+ins2 isoform (hD54) mRNA, complete	-0.017	0.075	4 consistent probesets
J04973	Human cytochrome bc-1 complex core protein II mRNA, co	-0.017	0.076	4 consistent probesets
D13641	Human mRNA for KIAA0016 gene, complete cds /cds=(101	-0.017	0.085	4 consistent probesets
AA978033	oq55e04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-15	-0.017	0.088	4 consistent probesets
U91316	Human acyl-CoA thioester hydrolase mRNA, complete cds	-0.017	0.091	4 consistent probesets
D87453	Human mRNA for KIAA0264 gene, partial cds /cds=(0,1249	-0.017	0.091	4 consistent probesets
M35416	HUMRALBA Human GTP-binding protein (RALB) mRN	-0.017	0.092	8 inconsistent probesets
X13839	Human mRNA for vascular smooth muscle alpha-actin /cds	-0.017	0.093	4 consistent probesets
D63391	Human mRNA for platelet activating factor acetylhydrol	-0.017	0.093	4 consistent probesets
M31642	Human hypoxanthine phosphoribosyltransferase (HPRT) m	-0.017	0.093	4 consistent probesets
L76702	Homo sapiens protein phosphatase 2A B56-delta (PP2A) m	-0.017	0.098	4 consistent probesets
U78110	Human prepro-neurturin mRNA, complete cds /cds=(0,593)	-0.017	0.100	4 consistent probesets
M55150	Human fumarylacetoacetate hydrolase mRNA, complete cd	-0.017	0.107	4 consistent probesets
M63138	HUMCATD5 Human cathepsin D (catD) gene, exons	-0.017	0.108	4 consistent probesets
X77567	H.sapiens mRNA for InsP3 5-phosphatase /cds=(99,1337)	-0.017	0.119	3 consistent probesets
M84526	Human adipsin/complement factor D mRNA, complete cds ,	-0.017	0.119	3 consistent probesets
U32519	Human GAP SH3 binding protein mRNA, complete cds /cds	-0.017	0.119	4 consistent probesets
D90282	Human carbamyl phosphate synthetase I (EC 6.3.4.16) mR	-0.017	0.125	4 consistent probesets
M17733	Human thymosin beta-4 mRNA, complete cds /cds=(77,211	-0.017	0.131	4 inconsistent probesets
AB011177	Homo sapiens mRNA for KIAA0605 protein, complete cds /	-0.017	0.133	4 consistent probesets
AF038961	Homo sapiens SL15 protein mRNA, complete cds /cds=(16,	-0.017	0.141	3 consistent probesets
L05568	Human Na+/Cl- dependent serotonin transporter mRNA, co	-0.017	0.145	4 consistent probesets
D86960	Human mRNA for KIAA0205 gene, complete cds /cds=(227	-0.017	0.147	4 consistent probesets
AF084645	Homo sapiens orphan nuclear receptor (PAR1) mRNA, com	-0.017	0.147	4 consistent probesets
AF040990	Homo sapiens roundabout 1 (robo1) mRNA, complete cds /	-0.017	0.153	3 consistent probesets
X15949	HSIRF2 Human mRNA for interferon regulatory f	-0.017	0.159	3 consistent probesets
M63967	Human mitochondrial aldehyde dehydrogenase x gene, com	-0.017	0.165	3 consistent probesets
AF070617	Homo sapiens clone 24812 mRNA sequence /cds=UNKNO	-0.017	0.166	3 consistent probesets
Y09445	HSTBX5 H.sapiens mRNA for transcription facto	-0.017	0.172	3 consistent probesets
X65633	HSACTHR H.sapiens ACTH-R gene for adrenocorti	-0.017	0.181	4 consistent probesets
AB018302	Homo sapiens mRNA for KIAA0759 protein, partial cds /c	-0.017	0.186	3 consistent probesets
R93981	yt73d01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-276	-0.017	0.186	3 consistent probesets
AF004849	Homo sapiens PKY protein kinase mRNA, complete cds /co	-0.017	0.202	3 consistent probesets
AI590869	tw88g03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-22(	-0.017	0.204	3 consistent probesets
Z46389	Homo sapiens encoding vasodilator-stimulated phosphopr	-0.017	0.213	3 consistent probesets
N25429	yx76b02.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-26	-0.017	0.218	3 consistent probesets

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AJ131186	Homo sapiens mRNA for nuclear matrix protein NMP200 /c	-0.017	0.227	3 inconsistent probesets
AJ005577	Homo sapiens pfkfb2 gene, exons 1 to 15 /cds=(93,1607)	-0.017	0.228	3 consistent probesets
AF007157	Homo sapiens clone 23856 unknown mRNA, partial cds /cd	-0.017	0.247	3 consistent probesets
AB011136	Homo sapiens mRNA for KIAA0564 protein, partial cds /c	-0.017	0.253	4 consistent probesets
U24267	Human pyrroline-5-carboxylate dehydrogenase (P5CDh) m	-0.017	0.253	3 consistent probesets
X73079	Homo sapiens encoding Polymeric immunoglobulin recepto	-0.017	0.314	3 consistent probesets
U45984	Homo sapiens CCR6 chemokine receptor (CMKBR6) gene,	-0.017	0.330	3 consistent probesets
M62424	Human thrombin receptor mRNA, complete cds /cds=(224,1	-0.017	0.348	4 consistent probesets
D14710	Human mRNA for ATP synthase alpha subunit, complete cd	-0.021	0.041	4 inconsistent probesets
H98552	yv97h03.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	-0.021	0.057	4 inconsistent probesets
AA173896	zp03b02.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-59	-0.021	0.069	4 consistent probesets
AI541042	pec1.2-1.D12.r Homo sapiens cDNA, 5 end /clone_end=5	-0.021	0.078	4 consistent probesets
U28389	Human dematin 52 kDa subunit mRNA, complete cds /cds=	-0.021	0.085	4 consistent probesets
U05770	Human annexin V (ANX5) gene /cds=(164,1126) /gb=U057	-0.021	0.098	4 consistent probesets
AI525633	PT1.3_04_A08.r Homo sapiens cDNA, 5 end /clone_end=5	-0.021	0.102	4 consistent probesets
X82554	HSSPHAR H.sapiens SPHAR gene for cyclin-relat	-0.021	0.102	8 consistent probesets
AF001691	Homo sapiens 195 kDa cornified envelope precursor mRNA	-0.021	0.117	4 consistent probesets
D38549	Human mRNA for KIAA0068 gene, partial cds /cds=(0,3816	-0.021	0.117	4 consistent probesets
W28790	54g3 Homo sapiens cDNA /gb=W28790 /gi=1308945 /	-0.021	0.118	4 consistent probesets
U72515	Human C3f mRNA, complete cds /cds=(117,1262) /gb=U72	-0.021	0.119	4 consistent probesets
AA149428	zl26a05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-503	-0.021	0.121	4 consistent probesets
AI655458	tt13a03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-224	-0.021	0.122	4 consistent probesets
U83460	Human high-affinity copper uptake protein (hCTR1) mRNA	-0.021	0.127	4 consistent probesets
U83598	HSU83598 Human death domain receptor 3 solubl	-0.021	0.135	4 consistent probesets
M97936	Human transcription factor ISGF-3 mRNA sequence /cds=U	-0.021	0.137	4 consistent probesets
AI087268	oz77e01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	-0.021	0.141	4 consistent probesets
U82319	Homo sapiens clone YDD19 mRNA sequence /cds=(0,206)	-0.021	0.161	4 inconsistent probesets
X99459	H.sapiens mRNA for sigma 3B protein /cds=(30,611) /gb=	-0.021	0.164	4 consistent probesets
U25975	HSU25975 Human serine kinase (hPAK65) mRNA, p	-0.021	0.169	4 consistent probesets
U94747	Human WD repeat protein HAN11 mRNA, complete cds /cd	-0.021	0.173	4 consistent probesets
Y18863	Homo sapiens mRNA for ribonuclease P protein subunit P	-0.021	0.188	4 consistent probesets
U31346	Human calpastatin mRNA, partial cds, long 3UTR /cds=(0	-0.021	0.203	4 consistent probesets
AF091090	Homo sapiens clone 669 unknown mRNA, complete sequer	-0.021	0.208	4 consistent probesets
X55503	H.sapiens pseudogene for metallothionein and AG/CT rep	-0.022	0.108	3 consistent probesets
AI040379	ox16c05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	-0.022	0.419	3 consistent probesets
U08316	HSU08316 Human insulin-stimulated protein kin	-0.023	0.051	8 inconsistent probesets
D38524	HUM5N Human mRNA for 5 -nucleotidase	-0.023	0.115	8 inconsistent probesets
L24804	Human (p23) mRNA, complete cds /cds=(232,714) /gb=L24	-0.025	0.082	4 consistent probesets
M59499	Human lipoprotein-associated coagulation inhibitor (LA	-0.025	0.093	4 consistent probesets
U28963	Human Gps2 (GPS2) mRNA, complete cds /cds=(90,1073)	-0.025	0.103	4 consistent probesets
AF047442	Homo sapiens vesicle trafficking protein sec22b mRNA,	-0.025	0.104	4 consistent probesets
AI148772	qc69h01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-17	-0.025	0.127	4 consistent probesets
AJ224741	Homo sapiens mRNA for matrilin-3 /cds=(17,1477) /gb=AJ	-0.025	0.136	4 consistent probesets
AF034956	Homo sapiens RAD51D mRNA, complete cds /cds=(124,99	-0.025	0.203	4 consistent probesets
AL035306	H.sapiens gene from PAC 426I6, similar to syntaxin 7 /	-0.025	0.206	4 consistent probesets
U57094	HSU57094 Human small GTP-binding protein mRNA	-0.025	0.228	4 consistent probesets
U57693	Human TFIID subunits TAF20 and TAF15 mRNA, complete	-0.025	0.232	4 consistent probesets
D87451	Human mRNA for KIAA0262 gene, complete cds /cds=(698	-0.025	0.363	2 consistent probesets
L25107	Human LIS mRNA /cds=UNKNOWN /gb=L25107 /gi=60245	-0.025	0.382	2 consistent probesets
AI817618	wk39f01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	-0.025	0.436	2 consistent probesets
X79563	H.sapiens 8.2kDa differentiation factor mRNA /cds=(60,	-0.025	0.489	2 consistent probesets
D37984	HUMDHQ1 Human mRNA for DNA helicase Q1, parti	-0.025	0.495	2 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
Z11692	H.sapiens mRNA for elongation factor 2 /cds=(0,2576) /	-0.025	0.045	4 consistent probesets
D31628	Human gene for 4-hydroxyphenylpyruvic acid dioxygenase	-0.025	0.055	4 consistent probesets
AF010312	Homo sapiens Pig7 (PIG7) mRNA, complete cds /cds=(79,7	-0.025	0.061	4 inconsistent probesets
D15057	Human mRNA for DAD-1, complete cds /cds=(66,407) /gb=	-0.025	0.064	4 consistent probesets
M34079	HUMTBP1 Human immunodeficiency virus tat tran	-0.025	0.066	4 consistent probesets
L33842	Homo sapiens (clone FFE-7) type II inosine monophospha	-0.025	0.067	4 consistent probesets
W26480	30b8 Homo sapiens cDNA /gb=W26480 /gi=1307179 /	-0.025	0.068	4 consistent probesets
U15131	Human p126 (ST5) mRNA, complete cds /cds=(114,3527) /	-0.025	0.071	2 consistent probesets
X74262	HSRBAP48 H.sapiens RbAp48 mRNA encoding retin	-0.025	0.087	8 inconsistent probesets
M13755	HUMIFN15K Human interferon-induced 17-kDa/15-	-0.025	0.095	4 inconsistent probesets
X89214	H.sapiens mRNA for haptoglobin related protein /cds=(1	-0.025	0.102	4 inconsistent probesets
U23942	Human lanosterol 14-demethylase cytochrome P450 (CYP5	-0.025	0.102	4 consistent probesets
U19822	Human acetyl-CoA carboxylase mRNA, complete cds /cds=	-0.025	0.108	4 consistent probesets
AA001791	zh86c04.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-428	-0.025	0.122	4 consistent probesets
X00442	Human mRNA for haptoglobin alpha(2FS)-beta precursor /	-0.025	0.127	4 consistent probesets
X60673	Human AK3 mRNA for adenylate kinase 3 /cds=UNKNOWN	-0.025	0.127	4 consistent probesets
Z48579	H.sapiens mRNA for disintegrin-metalloprotease (partia	-0.025	0.128	4 consistent probesets
AF001903	Human 3-hydroxyacyl-CoA dehydrogenase, isoform 2 mRN	-0.025	0.135	4 consistent probesets
U40391	Human serotonin N-acetyltransferase gene, complete cds	-0.025	0.135	4 consistent probesets
U66063	Homo sapiens calcium/calmodulin-dependent protein kina	-0.025	0.151	2 consistent probesets
Y08999	H.sapiens mRNA for Sop2p-like protein /cds=(33,1145) /	-0.025	0.164	4 consistent probesets
AL049962	Homo sapiens mRNA; cDNA DKFZp564P0823 (from clone	-0.025	0.167	4 consistent probesets
AL049980	Homo sapiens mRNA; cDNA DKFZp564C152 (from clone D	-0.025	0.176	4 consistent probesets
AI660225	we68f05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-234	-0.025	0.182	2 consistent probesets
AF041434	Homo sapiens potentially prenylated protein tyrosine p	-0.025	0.183	2 inconsistent probesets
AB002559	Homo sapiens mRNA for hunc18b2, complete cds /cds=(24	-0.025	0.194	4 consistent probesets
AL080172	Homo sapiens mRNA; cDNA DKFZp434G231 (from clone D	-0.025	0.203	2 consistent probesets
Z48570	H.sapiens Sp17 gene /cds=(1210,1665) /gb=Z48570 /gi=69	-0.025	0.256	4 consistent probesets
AB017019	Homo sapiens mRNA for JKTBP2, complete cds /cds=(535	-0.025	0.271	2 consistent probesets
U92436	HSU92436 Human mutated in multiple advanced c	-0.025	0.340	4 consistent probesets
AJ001699	Homo sapiens mRNA for Brachyury (T) protein /cds=(159,	-0.025	0.454	2 consistent probesets
AF003001	Homo sapiens TTAGGG repeat binding factor 1 (hTRF1-AS	-0.025	0.640	2 consistent probesets
M81118	Human alcohol dehydrogenase chi polypeptide (ADH5) gen	-0.027	0.072	8 consistent probesets
U01923	Human BTK region clone ftp-3 mRNA /cds=UNKNOWN /gb	-0.027	0.110	8 consistent probesets
M86934	Human GS1 (protein of unknown function) mRNA, complete	-0.029	0.069	4 consistent probesets
AJ001047	HSMATRIL3 Homo sapiens mRNA for matrilin-3	-0.029	0.070	4 inconsistent probesets
AF021819	Homo sapiens RNA-binding protein regulatory subunit mR	-0.029	0.072	4 consistent probesets
X54486	Human gene for C1-inhibitor /cds=(60,1562) /gb=X54486	-0.029	0.075	4 consistent probesets
Y08890	H.sapiens mRNA for Ran_GTP binding protein 5 /cds=(236	-0.029	0.078	4 consistent probesets
Z78308	HSZ78308 Homo sapiens cDNA /clone=1.47-(CEPH) /gb=Z	-0.029	0.084	4 inconsistent probesets
D78514	D78514 Homo sapiens mRNA for ubiquitin-conjug	-0.029	0.100	4 inconsistent probesets
M14219	Human chondroitin/dermatan sulfate proteoglycan (PG40)	-0.029	0.116	4 consistent probesets
X66365	HSSTHPKF H.sapiens mRNA PLSTIRE for serine/th	-0.029	0.152	4 consistent probesets
U75697	Human transcription regulator RPD3-2B mRNA, complete c	-0.029	0.165	4 consistent probesets
U17032	Human p190-B (p190-B) mRNA, complete cds /cds=(302,48	-0.029	0.180	4 inconsistent probesets
U40571	Human alpha1-syntrophin (SNT A1) mRNA, complete cds /	-0.029	0.192	4 consistent probesets
U52521	Human arfaptin 1, putative target protein of ADP-ribos	-0.029	0.233	4 consistent probesets
D88587	Homo sapiens mRNA for Hakata antigen, complete cds /cd	-0.029	0.241	4 consistent probesets
AI017382	ou92e06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-16	-0.029	0.345	4 consistent probesets
T75292	yc89b05.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-232	-0.031	0.099	8 consistent probesets
K03515	Human neuroleukin mRNA, complete cds /cds=(15,1691) /g	-0.033	0.041	4 inconsistent probesets
U43077	HSU43077 Human CDC37 homolog mRNA, complete c	-0.033	0.061	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
M14630	Human prothymosin alpha mRNA, complete cds /cds=UNK	-0.033	0.068	8 consistent probesets
AB020693	Homo sapiens mRNA for KIAA0886 protein, complete cds /	-0.033	0.070	4 consistent probesets
D23660	Human mRNA for ribosomal protein, complete cds /cds=(5	-0.033	0.076	4 consistent probesets
Y17448	Homo sapiens CCBL1 gene, last two exons /cds=(0,146) /	-0.033	0.087	3 consistent probesets
L46590	Homo sapiens very long chain acyl-CoA dehydrogenase ge	-0.033	0.088	4 inconsistent probesets
AI951946	wx39f10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	-0.033	0.091	4 consistent probesets
L13687	Homo sapiens ADP-ribosylation factor-like protein 2 (A	-0.033	0.094	4 consistent probesets
X87176	H.sapiens mRNA for 17-beta-hydroxysteroid dehydrogenas	-0.033	0.097	4 consistent probesets
AF040253	Homo sapiens transcription factor Tat-CT1 mRNA, comple	-0.033	0.098	3 inconsistent probesets
AF081280	Homo sapiens nucleoplasmin-3 (NPM3) mRNA, complete c	-0.033	0.101	4 consistent probesets
AF068706	Homo sapiens gamma2-adaptin (G2AD) mRNA, complete c	-0.033	0.109	3 consistent probesets
AF060543	Homo sapiens importin alpha 7 subunit mRNA, complete c	-0.033	0.119	3 consistent probesets
M13142	Human factor XI (blood coagulation factor) mRNA, compl	-0.033	0.129	3 consistent probesets
M29273	Human myelin-associated glycoprotein (MAG) mRNA, com	-0.033	0.135	4 consistent probesets
J02958	Human MET proto-oncogene mRNA, complete cds /cds=(1	-0.033	0.136	4 consistent probesets
L31529	Human beta1-syntrophin (SNT B1) gene, complete cds /cd	-0.033	0.139	4 inconsistent probesets
U63717	HSU63717 Homo sapiens osteoclast stimulating	-0.033	0.149	4 consistent probesets
Y10275	H.sapiens mRNA for L-3-phosphoserine phosphatase /cds=	-0.033	0.150	4 consistent probesets
Z97029	Homo sapiens mRNA for ribonuclease H I large subunit /	-0.033	0.167	3 consistent probesets
AF034957	Homo sapiens secreted T cell activation protein Attrac	-0.033	0.169	3 consistent probesets
M65214	Human (HeLa) helix-loop-helix protein HE47 (E2A) mRNA,	-0.033	0.170	4 consistent probesets
S48220	type I 5 iodothyronine deiodinase [human, mRNA, 2222 n	-0.033	0.171	4 consistent probesets
U75285	Homo sapiens apoptosis inhibitor survivin gene, comple	-0.033	0.186	3 consistent probesets
AJ011915	Homo sapiens mRNA for synaptosome associated protein c	-0.033	0.198	4 consistent probesets
AF052183	Homo sapiens clone 24804 mRNA sequence /cds=UNKNO	-0.033	0.199	4 consistent probesets
U25138	Human MaxiK potassium channel beta subunit mRNA, com	-0.033	0.202	1 consistent probesets
AB001928	Homo sapiens mRNA for cathepsin V, complete cds /cds=(	-0.033	0.205	3 consistent probesets
U24153	HSU24153 Human p21-activated protein kinase (	-0.033	0.211	4 consistent probesets
U29615	Human chitotriosidase precursor mRNA, complete cds /cd	-0.033	0.235	1 consistent probesets
M13792	HUMADAG Human adenosine deaminase (ADA) gene,	-0.033	0.262	3 consistent probesets
AF013249	Homo sapiens leukocyte-associated Ig-like receptor-1 (	-0.033	0.266	2 consistent probesets
D31764	Human mRNA for KIAA0064 gene, complete cds /cds=(222	-0.033	0.271	3 consistent probesets
AI391564	tg16b02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-210	-0.033	0.350	3 consistent probesets
AI743606	wg51f08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	-0.038	0.060	4 inconsistent probesets
U14394	HSU14394 Human tissue inhibitor of metallopro	-0.038	0.066	8 consistent probesets
D87002	D87002 Homo sapiens immunoglobulin lambda gen	-0.038	0.070	4 inconsistent probesets
U96074	Human translation initiation factor eIF3 p44 subunit m	-0.038	0.080	4 consistent probesets
W32483	zc67e07.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-327	-0.038	0.081	4 consistent probesets
X12901	Human mRNA for villin /cds=(24,2507) /gb=X12901 /gi=37	-0.038	0.082	4 consistent probesets
L22474	HUMBAXB Human Bax beta mRNA, complete cds"	-0.038	0.083	4 consistent probesets
L10284	Homo sapiens integral membrane protein, calnexin, (IP9	-0.038	0.087	4 consistent probesets
M68895	Human alcohol dehydrogenase 6 gene, complete cds /cds=	-0.038	0.096	4 inconsistent probesets
AF035531	Homo sapiens syntaxin 10 mRNA, complete cds /cds=(0,74	-0.038	0.098	4 inconsistent probesets
D59253	Human mRNA for NCBP interacting protein 1, complete cd	-0.038	0.098	4 consistent probesets
M33146	Human cysteine-rich peptide mRNA, complete cds /cds=(3	-0.038	0.098	4 consistent probesets
AA977136	oq24f02.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-158	-0.038	0.108	4 inconsistent probesets
AL079298	Homo sapiens mRNA full length insert cDNA clone EUROIM	-0.038	0.108	4 inconsistent probesets
U43368	Human VEGF related factor isoform VRF186 precursor (VR	-0.038	0.122	4 consistent probesets
AF030555	Homo sapiens acyl-CoA synthetase 4 (ACS4) mRNA, comp	-0.038	0.145	4 consistent probesets
AF058954	Homo sapiens GTP-specific succinyl-CoA synthetase beta	-0.038	0.152	4 inconsistent probesets
AA151922	zo30d07.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-588	-0.038	0.168	4 inconsistent probesets
W27675	36b3 Homo sapiens cDNA /gb=W27675 /gi=1307623 /	-0.038	0.175	4 inconsistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AF043733	Homo sapiens death effector domain-containing testicul	-0.038	0.186	4 consistent probesets
X52221	HSERCC25 H.sapiens ERCC2 gene, exons 1 & 2 (p	-0.038	0.125	7 consistent probesets
U79528	Human SR31747 binding protein 1 mRNA, complete cds /c	-0.042	0.075	4 consistent probesets
AF023676	Homo sapiens lamin B receptor homolog TM7SF2 (TM7SF	-0.042	0.085	4 inconsistent probesets
L12711	Homo sapiens transketolase (tk) mRNA, complete cds /cd	-0.042	0.087	4 consistent probesets
Z47553	H.sapiens mRNA for flavin-containing monooxygenase 5 (	-0.042	0.091	4 inconsistent probesets
M13699	Human ceruloplasmin (ferroxidase) mRNA, complete cds /	-0.042	0.091	4 consistent probesets
U41843	Human Dr1-associated corepressor (DRAP1) mRNA, comp	-0.042	0.095	4 consistent probesets
X63359	H.sapiens UGT2BIO mRNA for udp glucuronosyltransferase	-0.042	0.116	4 consistent probesets
AF070537	Homo sapiens clone 24606 mRNA sequence /cds=UNKNO	-0.042	0.124	4 consistent probesets
M63175	Human autocrine motility factor receptor mRNA /cds=(17	-0.042	0.138	4 consistent probesets
U69127	Human FUSE binding protein 3 (FBP3) mRNA, partial cds	-0.042	0.138	4 consistent probesets
D13413	HUMTA120 Human mRNA for tumor-associated 120	-0.042	0.142	4 inconsistent probesets
U01160	Human transmembrane 4 superfamily protein (SAS) mRNA	-0.042	0.148	4 consistent probesets
AB014536	Homo sapiens mRNA for KIAA0636 protein, complete cds /	-0.042	0.162	4 consistent probesets
AF051321	Homo sapiens Sam68-like phosphotyrosine protein alpha	-0.042	0.200	4 consistent probesets
U00928	Human clone CE29 4.1 (CAC)n/(GTG)n repeat-containing r	-0.042	0.489	2 consistent probesets
H12054	ym11b12.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-47	-0.044	0.194	3 consistent probesets
AF095288	Homo sapiens pituitary tumor transforming gene protein	-0.044	0.321	3 consistent probesets
X05236	Human fibroblast mRNA for aldolase A /cds=(146,1240) /	-0.046	0.058	4 consistent probesets
N25547	yx76e06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-26	-0.046	0.089	4 inconsistent probesets
L22009	Human hnRNP H mRNA, complete cds /cds=(72,1421) /gb=	-0.046	0.102	4 consistent probesets
AL096741	Homo sapiens mRNA; cDNA DKFZp586O0223 (from clone	-0.046	0.133	4 consistent probesets
W25866	14c12 Homo sapiens cDNA /gb=W25866 /gi=1305989 /	-0.046	0.152	4 consistent probesets
U13395	Human oxidoreductase (HHCMA56) mRNA, complete cds /	-0.046	0.159	4 consistent probesets
U66676	HSU66676 Homo sapiens cDNA /gb=U66676 /gi=1906561 /	-0.046	0.159	4 consistent probesets
N63574	yy63f05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-278	-0.046	0.165	4 consistent probesets
AF091077	Homo sapiens clone 558 unknown mRNA, complete sequer	-0.046	0.166	4 consistent probesets
X79780	H.sapiens YPT3 mRNA /cds=(6,662) /gb=X79780 /gi=7631	-0.046	0.169	4 consistent probesets
M87338	HUMA1SBU Human replication factor C, 40-kDa s	-0.046	0.185	4 consistent probesets
U57057	Human WD protein IR10 mRNA, complete cds /cds=(0,157	-0.050	0.158	4 consistent probesets
AL050019	Homo sapiens mRNA; cDNA DKFZp564C186 (from clone D	-0.050	0.161	2 consistent probesets
X71440	H.sapiens mRNA for peroxisomal acyl-CoA oxidase /cds=(	-0.050	0.161	4 consistent probesets
AA045160	zk63e03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-487	-0.050	0.294	2 consistent probesets
U58090	Human Hs-cul-4A mRNA, partial cds /cds=(0,1266) /gb=U5	-0.050	0.615	2 consistent probesets
M20867	Human glutamate dehydrogenase (GDH) mRNA, complete	-0.050	0.071	4 consistent probesets
M16342	Human nuclear ribonucleoprotein particle (hnRNP) C pro	-0.050	0.079	4 consistent probesets
J00194	human hla-dr antigen alpha-chain mrna & ivs fragments	-0.050	0.101	1 consistent probesets
U68487	Human 5-hydroxytryptamine7 receptor isoform b mRNA, co	-0.050	0.101	1 consistent probesets
M94856	Human fatty acid binding protein homologue (PA-FABP) m	-0.050	0.106	4 inconsistent probesets
AF000231	Homo sapiens rab11a GTPase mRNA, complete cds /cds=(	-0.050	0.122	4 consistent probesets
X52730	Human gene for phenylethanolamine N-methylase (PNMT)	-0.050	0.126	2 consistent probesets
W26677	11f7 Homo sapiens cDNA /gb=W26677 /gi=1305788 /	-0.050	0.127	3 consistent probesets
X74331	HSPRIM2 H.sapiens mRNA for DNA primase (subun	-0.050	0.130	7 consistent probesets
D63880	Human mRNA for KIAA0159 gene, complete cds /cds=(799	-0.050	0.137	4 consistent probesets
X78627	H.sapiens mRNA for translin /cds=(81,767) /gb=X78627 /	-0.050	0.137	4 consistent probesets
W27611	35b9 Homo sapiens cDNA /gb=W27611 /gi=1307559 /	-0.050	0.191	4 consistent probesets
L14565	Human peripherin (PRPH) gene exons 1-9, complete cds /	-0.050	0.202	1 consistent probesets
AJ012590	Homo sapiens mRNA for glucose 1-dehydrogenase /cds=(2	-0.050	0.244	1 consistent probesets
D30742	HUMCDPKIV Human mRNA for calmodulin-dependent	-0.050	0.252	1 consistent probesets
X84746	H.sapiens Histo-blood group AB0 gene, exon 1 /cds=(0,1	-0.050	0.252	1 consistent probesets
AB018310	Homo sapiens mRNA for KIAA0767 protein, partial cds /c	-0.050	0.259	2 consistent probesets

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AF038440	Homo sapiens phospholipase D2 (PLD2) mRNA, splice vari	-0.050	0.263	3 consistent probesets
AF029761	Homo sapiens decoy receptor 2 mRNA, complete cds /cds=	-0.050	0.270	3 consistent probesets
X97674	H.sapiens mRNA for transcriptional intermediary factor	-0.050	0.273	2 consistent probesets
M64925	Human palmitoylated erythrocyte membrane protein (MPP1	-0.050	0.278	3 consistent probesets
M90355	Human BTF3 protein homologue gene, complete cds /cds=(	-0.050	0.293	1 consistent probesets
AA975427	oq28g02.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-15	-0.050	0.302	1 consistent probesets
AI200373	qf98c03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-175	-0.050	0.302	1 consistent probesets
AB029821	Homo sapiens mRNA for phosphatidylethanolamine N-meth	-0.050	0.302	1 consistent probesets
L11066	Human mRNA sequence /cds=UNKNOWN /gb=L11066 /gi=	-0.050	0.325	4 consistent probesets
X58822	Human IFN-omega 1 gene for interferon-omega 1 /cds=(57	-0.050	0.353	1 consistent probesets
U48730	HSU48730 Homo sapiens transcription factor St	-0.050	0.403	1 consistent probesets
X02910	HSTNFA Human gene for tumor necrosis factor (	-0.050	0.403	1 consistent probesets
AF092563	Homo sapiens chromosome-associated protein-E (hCAP-E)	-0.050	0.403	1 consistent probesets
AB014557	Homo sapiens mRNA for KIAA0657 protein, partial cds /c	-0.050	0.403	1 consistent probesets
W28830	52c8 Homo sapiens cDNA /gb=W28830 /gi=1308778 /	-0.050	0.403	1 consistent probesets
X66867	HSMAXG H.sapiens max gene	-0.050	0.403	1 consistent probesets
N36267	N36267 yx98b09.r1 Soares melanocyte 2NbHM Hom	-0.050	0.410	3 consistent probesets
L13329	Homo sapiens iduronate-2-sulfatase (IDS) gene /cds=(33	-0.050	0.454	1 consistent probesets
AF052137	Homo sapiens clone 23918 mRNA sequence /cds=UNKNO	-0.050	0.454	1 consistent probesets
U03495	Human transcription factor LSF-ID mRNA, complete cds /	-0.050	0.454	1 consistent probesets
X12496	Human mRNA for erythrocyte membrane sialoglycoprotein	-0.050	0.504	1 consistent probesets
U89344	Human acetyl-CoA carboxylase (ACC2) mRNA, complete c	-0.050	0.504	1 consistent probesets
AL079294	Homo sapiens mRNA full length insert cDNA clone EUROIN	-0.050	0.524	2 consistent probesets
D87470	Human mRNA for KIAA0280 gene, partial cds /cds=(0,876)	-0.050	0.554	1 consistent probesets
AF070618	Homo sapiens clone 24627 mRNA sequence /cds=UNKNO	-0.050	0.554	1 consistent probesets
T83979	yd66a11.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-11	-0.050	0.655	1 consistent probesets
AB018304	Homo sapiens mRNA for KIAA0761 protein, partial cds /c	-0.050	0.756	1 consistent probesets
AB018282	Homo sapiens mRNA for KIAA0739 protein, partial cds /c	-0.050	0.756	1 consistent probesets
AF094521	Homo sapiens MSE55-related protein (UB1) mRNA, comple	-0.050	0.806	1 consistent probesets
M96954	Homo sapiens nucleolysin TIAR mRNA, complete cds /cds=	-0.050	1.008	1 consistent probesets
X59834	Human rearranged mRNA for glutamine synthase /cds=(109	-0.054	0.077	4 consistent probesets
U80764	Human EST clone 122887 mariner transposon Hsmar1 seq	-0.054	0.087	4 inconsistent probesets
AL049261	Homo sapiens mRNA; cDNA DKFZp564E053 (from clone D	-0.054	0.090	4 inconsistent probesets
L32961	Human 4-aminobutyrate aminotransferase (GABAT) mRNA	-0.054	0.095	4 consistent probesets
AJ009985	Homo sapiens mRNA for annexin 31 /cds=(436,1452) /gb=A	-0.054	0.125	4 consistent probesets
U46569	Human aquaporin-5 (AQP5) gene /cds=(0,797) /gb=U46569	-0.054	0.137	4 inconsistent probesets
AA224768	nc12d09.r1 Homo sapiens cDNA /clone=IMAGE-1007921 /g	-0.054	0.156	4 consistent probesets
L27841	Human autoantigen pericentriol material 1 (PCM-1) mRNA	-0.056	0.298	3 consistent probesets
X69086	H.sapiens mRNA for utrophin /cds=(0,10301) /gb=X69086	-0.058	0.102	4 inconsistent probesets
AF047448	Homo sapiens TLS-associated protein TASR mRNA, compl	-0.058	0.136	4 inconsistent probesets
X61094	H.sapiens RNA for GM2-activator protein (clone pGM2A)	-0.058	0.164	4 consistent probesets
AA009569	zi04h03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-429	-0.058	0.222	4 consistent probesets
Y10148	H.sapiens mRNA for NTR2 receptor /cds=(36,1268) /gb=Y1	-0.061	0.176	3 consistent probesets
U26209	Human renal sodium/dicarboxylate cotransporter (NADC1)	-0.061	0.220	3 consistent probesets
M64231	HUMSPERSYN Human spermidine synthase gene, co	-0.062	0.140	7 inconsistent probesets
M83216	Human aorta caldesmon mRNA, complete cds /cds=(229,26	-0.063	0.130	4 inconsistent probesets
X99142	H.sapiens mRNA for hair keratin, hHb6 /cds=(0,1451) /g	-0.063	0.145	4 consistent probesets
U72518	Human destrin-2 pseudogene mRNA, complete cds /cds=(2	-0.063	0.157	4 consistent probesets
U55980	HSU55980 Homo sapiens cDNA, 3 end /clone=25453 /clon	-0.067	0.093	4 consistent probesets
AF046798	untitled /cds=(113,2656) /gb=AF046798 /gi=3170406 /	-0.067	0.093	4 consistent probesets
D30036	HUMPITPA Human mRNA for phosphatidylinositol	-0.067	0.113	3 consistent probesets
AL050126	Homo sapiens mRNA; cDNA DKFZp586G011 (from clone D	-0.067	0.120	4 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
U38980	U38980 Human PMS2 related (hPMSR6) mRNA, com	-0.067	0.130	4 consistent probesets
U13061	HSDHEASTD6 Human dehydroepiandrosterone sulfo	-0.067	0.139	4 consistent probesets
AA079018	zm94e12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-54	-0.067	0.199	3 consistent probesets
AI671905	wb41d12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	-0.067	0.226	3 consistent probesets
Y13622	Homo sapiens mRNA for latent transforming growth facto	-0.067	0.234	3 consistent probesets
AB018566	Homo sapiens gene for Proline synthetase associated, c	-0.067	0.258	3 consistent probesets
U43203	Human thyroid transcription factor 1 (TTF-1) mRNA, com	-0.067	0.337	4 consistent probesets
AL049219	Homo sapiens mRNA; cDNA DKFZp564J1616 (from clone	-0.067	0.406	3 consistent probesets
M16592	HUMHCKB Human hemopoietic cell protein-tyrosi	-0.067	0.538	1 consistent probesets
X81637	H.sapiens clathrin light chain b gene /cds=UNKNOWN /gb	-0.070	0.215	5 consistent probesets
AB009282	Homo sapiens mRNA for cytochrome b5, partial cds /cds=	-0.071	0.057	4 inconsistent probesets
U39067	Homo sapiens translation initiation factor eIF3 p36 su	-0.071	0.077	4 inconsistent probesets
U19765	Human nucleic acid binding protein gene, complete cds	-0.071	0.082	4 inconsistent probesets
X97671	HSERYTHR H.sapiens mRNA for erythropoietin re	-0.071	0.082	4 inconsistent probesets
L76528	HUMPS1A11 Homo sapiens presenilin (PS1;S182)	-0.071	0.097	4 consistent probesets
K03183	HUMCGBBA3 Human chorionic gonadotropin beta s	-0.071	0.158	4 consistent probesets
AL049675	Human gene from PAC 886K2, chromosome 1 /cds=UNKN	-0.071	0.222	4 consistent probesets
AF019612	Homo sapiens S2P mRNA, complete cds /cds=(99,1658) /g	-0.075	0.316	2 consistent probesets
Z11695	HS40KDAP H.sapiens 40 kDa protein kinase rela	-0.075	0.124	4 inconsistent probesets
U41767	Human metargidin precursor mRNA, complete cds /cds=(7,	-0.075	0.129	4 consistent probesets
X80818	H.sapiens mRNA for metabotropic glutamate receptor typ	-0.075	0.131	2 consistent probesets
M90354	Human BTF3 protein homologue gene, complete cds /cds=	-0.075	0.133	4 consistent probesets
AF054177	Homo sapiens chromodomain-helicase-DNA-binding protein	-0.075	0.177	4 consistent probesets
J00140	Human dihydrofolate reductase gene /cds=(42,605) /gb=J	-0.075	0.259	2 consistent probesets
U08854	Human UDP glucuronosyltransferase precursor (UGT2B15)	-0.075	0.259	2 consistent probesets
X83618	H.sapiens mRNA for 3-hydroxy-3-methylglutaryl coenzyme	-0.075	0.268	2 consistent probesets
AJ005694	Homo sapiens mRNA for short form of beta II spectrin,	-0.075	0.271	2 consistent probesets
U39226	Human myosin VIIA (USH1B) mRNA, complete cds /cds=(2	-0.075	0.287	2 consistent probesets
AB026891	Homo sapiens mRNA for cystine/glutamate transporter, c	-0.075	0.328	2 consistent probesets
AI302176	qn58c03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-19	-0.075	0.385	2 consistent probesets
AF093264	Homo sapiens homer-2b mRNA, complete cds /cds=(0,1064	-0.075	0.394	2 consistent probesets
AF047470	Homo sapiens malate dehydrogenase precursor (MDH) mR	-0.075	0.428	2 consistent probesets
Y11588	H.sapiens mRNA for apoptosis specific protein /cds=(32	-0.075	0.449	2 consistent probesets
AL050114	Homo sapiens mRNA; cDNA DKFZp586P2219 (from clone	-0.075	0.475	2 consistent probesets
X60382	H.sapiens COL10A1 gene for collagen (alpha-1 type X) /	-0.075	0.484	2 consistent probesets
X51630	HSWT1 Human Wilms tumor WT1 mRNA for zinc fin	-0.075	0.524	2 consistent probesets
AL049932	Homo sapiens mRNA; cDNA DKFZp564H2416 (from clone	-0.075	0.543	2 consistent probesets
M14764	HUMNGFR Human nerve growth factor receptor mR	-0.078	0.149	3 consistent probesets
X76770	H.sapiens PAP mRNA /cds=UNKNOWN /gb=X76770 /gi=55	-0.079	0.067	4 consistent probesets
AB007191	Homo sapiens mRNA for AMY-1, complete cds /cds=(38,34	-0.079	0.093	4 inconsistent probesets
U36764	HSU36764 Human TGF-beta receptor interacting	-0.079	0.110	4 consistent probesets
L31801	Homo sapiens monocarboxylate transporter 1 (SLC16A1) m	-0.079	0.165	4 consistent probesets
D26535	Human gene for dihydrolipoamide succinyltransferase, c	-0.079	0.179	4 consistent probesets
AF051941	Homo sapiens type 6 nucleoside diphosphate kinase NM23	-0.079	0.217	4 consistent probesets
U61538	Human calcium-binding protein chp mRNA, complete cds /	-0.079	0.219	4 inconsistent probesets
U78107	Homo sapiens gamma SNAP mRNA, complete cds /cds=(7	-0.079	0.283	4 consistent probesets
X14968	HSCAMPR2 Human testis mRNA for the RII-alpha	-0.080	0.217	5 consistent probesets
U60805	HSU60805 Human oncostatin-M specific receptor	-0.081	0.141	7 inconsistent probesets
U21551	Human ECA39 mRNA, complete cds /cds=(0,1154) /gb=U2	-0.083	0.139	4 consistent probesets
M22898	HUMP53A11 Human phosphoprotein p53 gene, exon	-0.083	0.186	3 inconsistent probesets
AJ133421	Homo sapiens mRNA for leucocyte vacuolar protein sorti	-0.083	0.241	3 consistent probesets
U28687	Human zinc finger containing protein ZNF157 (ZNF157) m	-0.083	0.282	3 consistent probesets

Accession	Brief Description	Rate	StdDev	Probe Set Information
AL031737	Human DNA sequence from clone 8B22 on chromosome 1p	-0.083	0.308	3 consistent probesets
AL050178	Homo sapiens mRNA; cDNA DKFZp586J1822 (from clone	-0.083	0.339	4 consistent probesets
U14528	Human sulfate transporter (DTD) mRNA, complete cds /cd	-0.083	0.442	3 consistent probesets
AF063020	Homo sapiens lens epithelium-derived growth factor mRN	-0.083	0.540	3 consistent probesets
U56417	Human lysophosphatidic acid acyltransferase-alpha mRNA	-0.088	0.064	4 inconsistent probesets
U03109	Human aspartyl beta-hydroxylase mRNA, complete cds /cd	-0.088	0.102	4 inconsistent probesets
X91117	H.sapiens HG NET gene exon 1 /cds=(49,1902) /gb=X9111	-0.088	0.129	4 consistent probesets
AL049430	Homo sapiens mRNA; cDNA DKFZp586H201 (from clone D	-0.088	0.141	4 consistent probesets
AB020662	Homo sapiens mRNA for KIAA0855 protein, partial cds /c	-0.088	0.148	4 consistent probesets
AF009767	AF009767 Homo sapiens cDNA /clone=C97A-12 /gb=AF00	-0.090	0.115	8 inconsistent probesets
D42073	Human mRNA for reticulocalbin, complete cds /cds=(52,1	-0.092	0.199	4 consistent probesets
M27504	HUMTOPIIX Homo sapiens topoisomerase type II	-0.096	0.141	4 inconsistent probesets
M90356	Human BTF3 protein homologue gene, complete cds /cds=(	-0.096	0.165	4 consistent probesets
AL050253	H.sapiens mRNA similar to D29763 mouse mRNA for seizu	-0.100	0.202	1 consistent probesets
AI524873	promrna-10.C03.r Homo sapiens cDNA, 5 end /clone_end=	-0.100	0.235	4 consistent probesets
AB018344	Homo sapiens mRNA for KIAA0801 protein, complete cds /	-0.100	0.280	4 consistent probesets
U88965	Human PO42 gene, complete cds /cds=(327,932) /gb=U889	-0.100	0.370	1 consistent probesets
AW043812	wy81b07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-25	-0.100	0.480	2 consistent probesets
M12886	HUMTCBYY Human T-cell receptor active beta-ch	-0.100	0.161	2 inconsistent probesets
W28907	53e12 Homo sapiens cDNA /gb=W28907 /gi=1308855 /	-0.100	0.162	4 consistent probesets
AF038185	Homo sapiens clone 23700 mRNA sequence /cds=UNKNO	-0.100	0.162	3 consistent probesets
X71661	H.sapiens ERGIC-53 mRNA /cds=(21,1553) /gb=X71661 /g	-0.100	0.187	4 consistent probesets
AJ238381	Homo sapiens pax9 gene, exons 1-2 and joined CDS /cds=	-0.100	0.193	2 consistent probesets
M55513	Human potassium channel (HPCN1) mRNA, complete cds /	-0.100	0.202	1 consistent probesets
AA873266	oh68e03.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-14	-0.100	0.202	1 consistent probesets
AA013087	ze27c09.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-360	-0.100	0.208	3 consistent probesets
S75989	gamma-aminobutyric acid transporter type 3 [human, fet	-0.100	0.232	2 consistent probesets
AI885381	wl93b01.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	-0.100	0.235	3 consistent probesets
L39059	Homo sapiens transcription factor SL1 mRNA, complete c	-0.100	0.248	2 consistent probesets
AF052186	Homo sapiens clone 24431 mRNA sequence /cds=UNKNO	-0.100	0.252	2 consistent probesets
AB011163	Homo sapiens mRNA for KIAA0591 protein, partial cds /c	-0.100	0.252	1 consistent probesets
M34455	Human interferon-gamma-inducible indoleamine 2,3-dioxy	-0.100	0.252	1 consistent probesets
AI290660	qm12b10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-18	-0.100	0.293	1 consistent probesets
S78296	neurofilament-66 [human, fetal brain, mRNA, 3197 nt] /	-0.100	0.302	1 consistent probesets
S80864	cytochrome c-like polypeptide [human, lung adenocarcin	-0.100	0.353	1 consistent probesets
D28113	Human mRNA for MOBP (myelin-associated oligodendrocy	-0.100	0.353	1 consistent probesets
W26805	13a12 Homo sapiens cDNA /gb=W26805 /gi=1305889 /	-0.100	0.403	1 consistent probesets
AB006628	Homo sapiens mRNA for KIAA0290 gene, partial cds /cds=	-0.100	0.403	1 consistent probesets
U34683	Human glutathione synthetase mRNA, complete cds /cds=(	-0.100	0.422	2 consistent probesets
AJ007014	Homo sapiens mRNA for AMMERC1 protein /cds=(168,116	-0.100	0.454	1 consistent probesets
X17360	HSHOX51 Human HOX 5.1 gene for HOX 5.1 protei	-0.100	0.454	1 consistent probesets
X78926	H.sapiens HZF3 mRNA for zinc finger protein /cds=(0,11	-0.100	0.454	1 consistent probesets
U60207	Human stress responsive serine/threonine protein kinas	-0.100	0.465	2 consistent probesets
M95767	Homo sapiens di-N-acetylchitobiase mRNA, complete cds	-0.100	0.504	1 consistent probesets
Z75190	H.sapiens mRNA for apolipoprotein E receptor 2 /cds=(0	-0.100	0.539	2 consistent probesets
X52228	Human mRNA for secreted epithelial tumour mucin antige	-0.100	0.550	2 consistent probesets
Y10746	H.sapiens mRNA for protein containing MBD 1 /cds=(139,	-0.100	0.585	1 consistent probesets
M22976	Human cytochrome b5 mRNA, 3 end /cds=UNKNOWN /gb	-0.100	0.655	1 consistent probesets
X96744	H.sapiens PAX7 gene, exon 1 (and joined CDS) /cds=(599	-0.100	0.655	1 consistent probesets
D84488	Homo sapiens mRNA for small GTP-binding protein, compl	-0.100	0.706	1 consistent probesets
AL050209	Homo sapiens mRNA; cDNA DKFZp586G0123 (from clone	-0.100	0.958	1 consistent probesets
AJ237839	Homo sapiens mRNA for hypothetical protein /cds=(85,69	-0.100	1.008	1 consistent probesets

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L42450	Homo sapiens pyruvate dehydrogenase kinase isoenzyme 1	-0.104	0.170	4 consistent probesets
D14582	Human mRNA for epimorphin /cds=(95,994) /gb=D14582 /g	-0.108	0.519	2 consistent probesets
Z85986	Human DNA sequence from clone 108K11 on chromosome	-0.112	0.148	7 consistent probesets
AF027515	Homo sapiens trans-golgi network glycoprotein 48 (TGN)	-0.113	0.101	4 inconsistent probesets
AJ001612	Homo sapiens mRNA for L-3-phosphoserine-phosphatase h	-0.117	0.154	4 inconsistent probesets
Y09568	Homo sapiens mRNA for SNAP23B protein, complete CDS	-0.117	0.213	4 consistent probesets
W26406	29b7 Homo sapiens cDNA /gb=W26406 /gi=1307105 /	-0.117	0.378	2 consistent probesets
S79325	SYT...SSX1 {translocation breakpoint} [human, synovial	-0.121	0.151	4 consistent probesets
D12676	Human mRNA for lysosomal sialoglycoprotein, complete c	-0.121	0.170	4 inconsistent probesets
Z12830	H.sapiens mRNA for SSR alpha subunit /cds=(29,889) /gb	-0.125	0.085	4 inconsistent probesets
U60205	Human methyl sterol oxidase (ERG25) mRNA, complete cd	-0.125	0.121	4 inconsistent probesets
S76346	AML1=AML1 {alternatively spliced, exons 5 and b} [huma	-0.125	0.221	4 consistent probesets
M62302	HUMGDF1 Human growth/differentiation factor 1	-0.125	0.225	2 consistent probesets
AL031846	dJ742C19.5 (novel Chromobox protein) /cds=(89,844) /gb	-0.125	0.232	2 consistent probesets
AF099731	Homo sapiens connexin 31.1 (GJB5) gene, complete cds /	-0.125	0.282	2 inconsistent probesets
U22028	HSU22028 Human cytochrome P450 (CYP2A13) gene	-0.125	0.318	2 consistent probesets
W87858	zh68c03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-417	-0.125	0.721	2 consistent probesets
X87838	H.sapiens mRNA for beta-catenin /cds=(214,2559) /gb=X8	-0.129	0.088	4 inconsistent probesets
AF027516	Homo sapiens trans-golgi network glycoprotein 51 (TGN)	-0.129	0.119	4 consistent probesets
X59618	H.sapiens RR2 mRNA for small subunit ribonucleotide re	-0.129	0.176	4 inconsistent probesets
AJ001810	Homo sapiens mRNA for pre-mRNA cleavage factor I subu	-0.133	0.118	4 inconsistent probesets
M22299	Human T-plastin polypeptide mRNA, complete cds, clone	-0.133	0.170	5 consistent probesets
AB020635	Homo sapiens mRNA for KIAA0828 protein, partial cds /c	-0.133	0.269	3 consistent probesets
J00139	HUMFOL5 Human dihydrofolate reductase gene, e	-0.133	0.393	3 consistent probesets
X74614	H.sapiens ODF2 (allele 2) gene for outer dense fiber p	-0.133	0.437	1 consistent probesets
U12778	Human acyl-CoA dehydrogenase mRNA, complete cds /cds	-0.142	0.100	4 inconsistent probesets
U39487	Human xanthine dehydrogenase/oxidase mRNA, complete	-0.142	0.273	2 consistent probesets
X15331	Human mRNA for phosphoribosylpyrophosphate synthetase	-0.144	0.108	3 consistent probesets
AF055023	Homo sapiens clone 24723 mRNA sequence /cds=UNKNO	-0.144	0.387	3 consistent probesets
U50648	HSIIPKR17 Human interferon-inducible RNA-depe	-0.146	0.183	4 consistent probesets
X15722	Human mRNA for glutathione reductase (EC 1.6.4.2) /cds	-0.150	0.097	4 consistent probesets
AF038897	Homo sapiens syntaxin 16 mRNA, complete cds /cds=(0,92	-0.150	0.108	3 inconsistent probesets
AF038198	Homo sapiens clone 23928 mRNA sequence /cds=UNKNO	-0.150	0.151	1 consistent probesets
D63861	D63861 Homo sapiens DNA for cyclophilin 40, c	-0.150	0.170	4 inconsistent probesets
U21931	Human fructose-1,6-biphosphatase (FBP1) gene /cds=(211	-0.150	0.182	2 consistent probesets
L01694	Homo sapiens (clone 58N-1) Ga subunit mRNA, complete c	-0.150	0.195	1 consistent probesets
U86358	Human chemokine (TECK) mRNA, complete cds /cds=(0,45	-0.150	0.202	1 consistent probesets
M32373	Human arylsulfatase B (ASB) mRNA, complete cds /cds=(5	-0.150	0.203	2 consistent probesets
U81984	Human endothelial PAS domain protein 1 (EPAS1) mRNA,	-0.150	0.241	3 consistent probesets
AF087036	Homo sapiens musculin mRNA, partial cds /cds=(0,606) /	-0.150	0.252	1 consistent probesets
AF091071	Homo sapiens clone 192 Rer1 mRNA, complete cds /cds=(	-0.150	0.268	2 consistent probesets
AB022017	Homo sapiens mRNA for AMP-activated protein kinase alp	-0.150	0.294	2 consistent probesets
AB006713	Homo sapiens mRNA for dihydropyrimidinase related prot	-0.150	0.302	1 consistent probesets
AF070596	Homo sapiens clone 24796 mRNA sequence /cds=UNKNO	-0.150	0.353	1 consistent probesets
X16867	Human mRNA for cytochrome P-450IID (clone pMP34) /cds	-0.150	0.353	1 consistent probesets
D87447	Human mRNA for KIAA0258 gene, complete cds /cds=(85,1	-0.150	0.403	1 consistent probesets
M25269	HUMELK1A Homo sapiens tyrosine kinase (ELK1)	-0.150	0.403	1 consistent probesets
AB014583	Homo sapiens mRNA for KIAA0683 protein, complete cds /	-0.150	0.403	1 consistent probesets
AF038199	Homo sapiens clone 23728 mRNA sequence /cds=UNKNO	-0.150	0.403	1 consistent probesets
M34715	Human pregnancy-specific beta-1-glycoprotein mRNA PSG	-0.150	0.454	1 consistent probesets
U68385	Human Meis1-related protein 2 (MRG2), mRNA, partial cd	-0.150	0.487	2 consistent probesets
X64624	H.sapiens mRNA for RDC-1 POU domain containing protei	-0.150	0.504	1 consistent probesets

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AB011154	Homo sapiens mRNA for KIAA0582 protein, partial cds /c	-0.150	0.504	1 consistent probesets
U92014	Human clone 121711 defective mariner transposon Hsmar2	-0.150	0.504	1 consistent probesets
AF020267	Homo sapiens myosin-IXb splice variant (Myo9b) mRNA, p	-0.150	0.554	1 consistent probesets
X70297	H.sapiens mRNA for neuronal nicotinic acetylcholine re	-0.150	0.554	1 consistent probesets
X78686	H.sapiens ENA-78 mRNA /cds=(106,450) /gb=X78686 /gi=4	-0.150	0.554	1 consistent probesets
AB023182	Homo sapiens mRNA for KIAA0965 protein, partial cds /c	-0.150	0.634	1 consistent probesets
X57527	Human COL8A1 mRNA for alpha 1(VIII) collagen /cds=(0,2	-0.150	0.655	1 consistent probesets
X02751	HSNRASR Human N-ras mRNA and flanking regions	-0.162	0.150	4 inconsistent probesets
Y13647	Homo sapiens mRNA for stearoyl-CoA desaturase /cds=(20	-0.162	0.171	4 inconsistent probesets
D13540	HUMSHPTP3 Homo sapiens SH-PTP3 mRNA for prote	-0.163	0.210	4 inconsistent probesets
D89678	Homo sapiens mRNA for A+U-rich element RNA binding fa	-0.167	0.369	3 consistent probesets
S68271	CREM=cyclic AMP-responsive element modulator [human,	-0.167	0.463	3 consistent probesets
D16217	Human mRNA for calpastatin, complete cds /cds=(162,228	-0.171	0.305	4 consistent probesets
U82256	Homo sapiens arginase type II mRNA, complete cds /cds=	-0.175	0.392	2 consistent probesets
Y17979	Homo sapiens mRNA for glycoprotein 6-alpha-L-fucosyltr	-0.175	0.502	2 consistent probesets
U08997	Human glutamate dehydrogenase gene, complete cds /cds=	-0.181	0.047	8 inconsistent probesets
U28964	Homo sapiens 14-3-3 protein mRNA, complete cds /cds=(1	-0.188	0.097	4 inconsistent probesets
J03071	Human growth hormone (GH-1 and GH-2) and chorionic so	-0.194	0.136	3 inconsistent probesets
X02469	HSP53 Human mRNA for p53 cellular tumor antig	-0.200	0.153	3 consistent probesets
AL049299	Homo sapiens mRNA; cDNA DKFZp564P233 (from clone D	-0.200	0.252	1 consistent probesets
M57230	Human membrane glycoprotein gp130 mRNA, complete cd	-0.200	0.253	3 inconsistent probesets
AF060231	Homo sapiens herpesvirus entry protein C (HVEC) mRNA,	-0.200	0.353	1 consistent probesets
AF091890	Homo sapiens G-protein coupled receptor RE2 mRNA, com	-0.200	0.353	1 consistent probesets
AL042668	DKFZp434O1721_r1 Homo sapiens cDNA, 5 end /clone=D	-0.200	0.454	1 consistent probesets
Z36715	H.sapiens mRNA for Net transcription factor /cds=(279,	-0.200	0.504	1 consistent probesets
W25985	17e6 Homo sapiens cDNA /gb=W25985 /gi=1306252 /	-0.200	0.554	1 consistent probesets
AF007111	AF007111 Homo sapiens MDM2-like p53-binding p	-0.200	0.605	1 consistent probesets
L40157	Human endosome-associated protein (EEA1) mRNA, comp	-0.200	0.655	1 consistent probesets
AI798834	we93c04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-23	-0.200	0.756	1 consistent probesets
AB020640	Homo sapiens mRNA for KIAA0833 protein, partial cds /c	-0.200	1.058	1 consistent probesets
X79201	H.sapiens mRNA for SYT /cds=(3,1178) /gb=X79201 /gi=53	-0.204	0.163	4 inconsistent probesets
M11507	Human transferrin receptor mRNA, complete cds (_5, _M,	-0.208	0.075	12 inconsistent probesets
AF013591	Homo sapiens homolog of the Aspergillus nidulans sudD	-0.225	0.642	2 consistent probesets
U31382	Human G protein gamma-4 subunit mRNA, complete cds /c	-0.250	0.154	4 inconsistent probesets
AB012229	Homo sapiens gene for fructose-6-phosphate,2-kinase/fr	-0.250	0.353	1 consistent probesets
M74525	HUMHHR6B Human HHR6B (yeast RAD 6 homologue)	-0.250	0.554	1 consistent probesets
AB011083	Homo sapiens mRNA for KIAA0511 protein, partial cds /c	-0.250	0.605	1 consistent probesets
AB002366	Human mRNA for KIAA0368 gene, partial cds /cds=(0,4327	-0.250	0.655	1 consistent probesets
AF013168	Homo sapiens hamartin (TSC1) mRNA, complete cds /cds=	-0.250	0.655	1 consistent probesets
AI739548	wi23d06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-239	-0.250	0.706	1 consistent probesets
X61615	H.sapiens mRNA for leukemia inhibitory factor (LIF) re	-0.250	1.058	1 consistent probesets
W25828	13g2 Homo sapiens cDNA /gb=W25828 /gi=1305951 /	-0.275	0.636	2 consistent probesets
L19161	HUMIEF2G Human translation initiation factor	-0.290	0.065	8 inconsistent probesets
Z12173	H.sapiens GNS mRNA encoding glucosamine-6-sulphatase	-0.293	0.110	5 inconsistent probesets
D87120	Homo sapiens mRNA for GS3786, complete cds /cds=(167,	-0.294	0.374	3 consistent probesets
AA151971	zo30b03.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-588	-0.300	0.101	1 consistent probesets
D50402	Human mRNA for NRAMP1, complete cds /cds=(175,1827)	-0.300	0.252	1 consistent probesets
J04129	Human placental protein 14 (PP14) mRNA, complete cds /	-0.300	0.302	1 consistent probesets
AI857673	wk96c02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-24	-0.300	0.439	1 consistent probesets
U43083	Human G alpha-q (Gaq) mRNA, complete cds /cds=(220,12	-0.300	0.504	1 consistent probesets
AF060219	Homo sapiens RCC1-like G exchanging factor RLG mRNA,	-0.300	0.554	1 consistent probesets
D28483	Human scr3 mRNA for RNA binding protein SCR3, complet	-0.300	0.605	1 consistent probesets

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L09235	Human vacuolar ATPase (isoform VA68) mRNA, complete	-0.329	0.132	4 inconsistent probesets
W28953	54b7 Homo sapiens cDNA /gb=W28953 /gi=1308901 /	-0.350	0.403	1 consistent probesets
W25958	18h6 Homo sapiens cDNA /gb=W25958 /gi=1306360 /	-0.450	0.146	1 consistent probesets