

71	221764_at	chromosome 19 open reading frame 22 Transcribed locus, weakly similar to XP_001111709.1 similar to beta globin	C19orf22	0.0379812	1.02865
72	217572_at	[Mac	---	0.00855195	1.04219
73	216565_x_at	interferon induced transmembrane protein pseudogene	LOC391020	0.00037369	1.05037
74	219863_at	hect domain and RLD 5	HERC5	0.00294819	1.05242
75	210873_x_at	apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 3A	APOBEC3A	0.00859484	1.07536
76	201285_at	makorin, ring finger protein, 1	MKRN1	0.0114255	1.08247
77	204187_at	guanosine monophosphate reductase	GMPR	0.029472	1.0959
78	201161_s_at	cold shock domain protein A	CSDA	0.0152193	1.11688
79	212203_x_at	interferon induced transmembrane protein 3 (1-8U)	IFITM3	0.0001597	1.18928
80	221748_s_at	tensin 1	TNS1	0.0188447	1.20426
81	44673_at	sialic acid binding Ig-like lectin 1, sialoadhesin	SIGLEC1	0.00048734	1.21366
82	221479_s_at	BCL2/adenovirus E1B 19kDa interacting protein 3-like	BNIP3L	0.0108822	1.21449
83	202219_at	solute carrier family 6 (neurotransmitter transporter, creatine), member 8	SLC6A8	0.0222675	1.24964
84	218136_s_at	solute carrier family 25, member 37	SLC25A37	0.017885	1.26406
85	205660_at	2'-5'-oligoadenylate synthetase-like	OASL	0.00115211	1.29764
86	207827_x_at	synuclein, alpha (non A4 component of amyloid precursor)	SNCA	0.0208582	1.30208
87	203153_at	interferon-induced protein with tetratricopeptide repeats 1	IFIT1	0.00654798	1.30861
88	203116_s_at	ferrochelatase (porphyrin)	FECH	0.0441259	1.33516
89	204972_at	2'-5'-oligoadenylate synthetase 2, 69/71kDa	OAS2	0.00178432	1.40057
90	202145_at	lymphocyte antigen 6 complex, locus E	LY6E	0.00489014	1.42123
91	217202_s_at	glutamate-ammonia ligase (glutamine synthetase)	GLUL	0.00361312	1.43465
92	221920_s_at	solute carrier family 25, member 37	SLC25A37	0.00517058	1.49802
93	204415_at	interferon, alpha-inducible protein 6	IFI6	0.00127746	1.53927
94	212907_at	solute carrier family 30 (zinc transporter), member 1 myxovirus (influenza virus) resistance 1, interferon-inducible protein p78	SLC30A1	0.00338279	1.547
95	202086_at	(mous	MX1	0.00035515	1.56271
96	214433_s_at	selenium binding protein 1	SELENBP1	0.0243698	1.61176
97	205483_s_at	ISG15 ubiquitin-like modifier	ISG15	0.00365418	1.62692
98	200923_at	lectin, galactoside-binding, soluble, 3 binding protein	LGALS3BP	0.00010564	1.74982
99	219519_s_at	sialic acid binding Ig-like lectin 1, sialoadhesin	SIGLEC1	0.00138396	1.8583
100	216336_x_at	metallothionein 1A /// metallothionein 1M /// metallothionein 1 pseudogene 2	MT1A /// MT1	0.00132438	1.8976
101	206461_x_at	metallothionein 1H /// metallothionein 1 pseudogene 2	MT1H /// MT1	0.0009704	1.91368
102	214453_s_at	interferon-induced protein 44	IFI44	0.00021587	1.94236
103	219672_at	thyroid associated factor	ERAF	0.0104699	1.9915
104	205592_at	interleukin 8	IL8	0.00675229	1.9978
105	211456_x_at	metallothionein 1 pseudogene 2	MT1P2	0.0003371	2.02327
106	217165_x_at	metallothionein 1F	MT1F	0.00057925	2.02345
107	212185_x_at	metallothionein 2A	MT2A	0.00042827	2.03254
108	204745_x_at	metallothionein 1G	MT1G	0.00158058	2.04646
109	213629_x_at	metallothionein 1F	MT1F	0.00025608	2.07399
110	205950_s_at	carbonic anhydrase I	CA1	0.00675552	2.14908
111	204466_s_at	synuclein, alpha (non A4 component of amyloid precursor)	SNCA	0.0140767	2.23058
112	208581_at	metallothionein 1X	MT1X	0.00009042	2.27717
113	212859_x_at	metallothionein 1E	MT1E	0.00070366	2.3156
114	204326_x_at	metallothionein 1X	MT1X	0.00109143	2.56709
115	204439_at	interferon-induced protein 44-like	IFI44L	0.00025503	2.66117
116	211560_s_at	aminolevulinate, delta-, synthase 2 (sideroblastic/hypochromic anemia)	ALAS2	0.00289969	2.92339
117	202411_at	interferon, alpha-inducible protein 27	IFI27	1.97E-06	5.25733