

Affymetrix Probe Set	Gene Name / Probe Set	Chromosome Map Position	SAM Score	Subgroup
206640_x_at	<i>GAGE2</i>	Xp11.23	8.18	PR
202503_s_at	<i>KIAA0101</i>	15q22.31	7.84	PR
202705_at	<i>CCNB2</i>	15q22.2	7.79	PR
208235_x_at	<i>GAGE5</i>	Xp11.4-p11.2	7.78	PR
202589_at	<i>TYMS</i>	18p11.32	7.4	PR
213008_at	<i>FLJ10719</i>	15q25-q26	7.39	PR
225834_at	<i>FAM72A</i>	1p11.2	7.37	PR
223381_at	<i>CDCA1</i>	1q23.3	7.22	PR
226980_at	<i>DEPDC1B</i>	5q12.1	7.20	PR
208155_x_at	<i>GAGE4</i>	Xp11.4-p11.2	7.19	PR
201890_at	<i>RRM2</i>	2p25-p24	7.15	PR
223229_at	<i>UBE2T</i>	1q32.1	7.09	PR
207739_s_at	<i>GAGE1</i>	Xp11.4-p11.2	7.08	PR
211762_s_at	<i>KPNA2</i>	17q23.1-q23.3	7.06	PR
214612_x_at	<i>MAGEA6</i>	Xq28	7.06	PR
204023_at	<i>RFC4</i>	3q27	6.96	PR
209942_x_at	<i>MAGEA3</i>	Xq28	6.94	PR
218039_at	<i>NUSAP1</i>	15q15.1	6.87	PR
203755_at	<i>BUB1B</i>	15q15	6.85	PR
202095_s_at	<i>BIRC5</i>	17q25	6.83	PR
220651_s_at	<i>MCM10</i>	10p13	6.82	PR
218156_s_at	<i>FLJ10534</i>	17p13.3	6.81	PR
236641_at	<i>KIF14</i>	1pter-q31.3	6.74	PR
210052_s_at	<i>TPX2</i>	20q11.2	6.74	PR
222077_s_at	<i>RACGAP1</i>	12q13.12	6.72	PR
219990_at	<i>E2F8</i>	11p15.1	6.67	PR
222606_at	<i>222606_at</i>	15q22.31	6.50	PR
235572_at	<i>SPBC24</i>	19p13.2	6.49	PR
201292_at	<i>TOP2A</i>	17q21-q22	6.47	PR
208079_s_at	<i>STK6</i>	20q13.2-q13.3	6.47	PR
209891_at	<i>SPBC25</i>	2q24.3	6.47	PR
214710_s_at	<i>CCNB1</i>	5q12	6.42	PR
228273_at	<i>FLJ11029</i>	17q23.2	6.41	PR
219918_s_at	<i>ASPM</i>	1q31	6.38	PR
202580_x_at	<i>FOXM1</i>	12p13	6.38	PR
226287_at	<i>NY-REN-41</i>	11p14.1	6.35	PR
211576_s_at	<i>SLC19A1</i>	21q22.3	6.32	PR
203213_at	<i>CDC2</i>	10q21.1	6.22	PR
204033_at	<i>TRIP13</i>	5p15.33	6.21	PR
204822_at	<i>TTK</i>	6q13-q21	6.20	PR
207325_x_at	<i>MAGEA1</i>	Xq28	6.19	PR
202427_s_at	<i>BRP44</i>	1q24	6.18	PR
203358_s_at	<i>EZH2</i>	7q35-q36	6.17	PR
1554271_a_at	<i>C1orf155</i>	1q25.1	6.11	PR
203764_at	<i>DLG7</i>	14q22.3	6.11	PR
218009_s_at	<i>PRC1</i>	15q26.1	6.09	PR
219000_s_at	<i>DCC1</i>	8q24.12	6.08	PR
204641_at	<i>NEK2</i>	1q32.2-q41	6.06	PR
202954_at	<i>UBE2C</i>	20q13.12	6.03	PR
217640_x_at	<i>C18orf24</i>	18q21.1	6.00	PR
227949_at	<i>PHACTR3</i>	20q13.33	9.46	LB
205590_at	<i>RASGRP1</i>	15q15	6.56	LB
217738_at	<i>PBEF1</i>	7q22.3	6.55	LB
221538_s_at	<i>PLXNA1</i>	3q21.3	6.44	LB
1553380_at	<i>PARP15</i>	3q21.1	5.95	LB
228450_at	<i>PLEKHA7</i>	11p15.1	5.81	LB
231740_at	<i>KCNJ11</i>	11p15.1	5.75	LB

230781_at	230781_at	4q21.21	5.70	LB
213622_at	COL9A2	1p33-p32	5.57	LB
222802_at	EDN1	6p24.1	5.57	LB
242417_at	LOC283278	11p15.1	5.51	LB
205159_at	CSF2RB	22q13.1	5.51	LB
222453_at	CYBRD1	2q31.1	5.47	LB
209731_at	NTHL1	16p13.3	5.34	LB
218311_at	MAP4K3	2p22.1	5.06	LB
227006_at	PPP1R14A	19q13.1	4.81	LB
227935_s_at	PCGF5	10q23.32	4.8	LB
1552388_at	FLJ30901	22q13.31	4.79	LB
227473_at	CTTN	11q13	4.79	LB
1558397_at	1558397_at	17q23.3	4.79	LB
226430_at	LOC253981	4p14	4.73	LB
225613_at	MAST4	5q12.3	4.64	LB
230930_at	LOC338620	10p11.23	4.53	LB
205937_at	CGREF1	2p23.3	4.53	LB
210448_s_at	P2RX5	17p13.3	4.52	LB
208614_s_at	FLNB	3p14.3	4.45	LB
229359_at	229359_at	2p23.3	4.44	LB
225990_at	BOC	3q13.2	4.43	LB
205780_at	BIK	22q13.31	4.35	LB
229067_at	229067_at	1	4.31	LB
230083_at	USP53	4q26	4.30	LB
225282_at	SMAP1L	1p35.3-p34.1	4.28	LB
239230_at	HES5	1p36.32	4.28	LB
202088_at	SLC39A6	18q12.2	4.25	LB
228360_at	LOC130576	2q23.3	4.24	LB
202484_s_at	MBD2	18q21	4.23	LB
34726_at	CACNB3	12q13	4.23	LB
213424_at	KIAA0895	7p14.1	4.21	LB
219561_at	COPZ2	17q21.32	4.18	LB
235427_at	235427_at	2q33.1	4.17	LB
205190_at	PLS1	3q23	4.17	LB
212614_at	ARID5B	10q21.2	4.16	LB
236782_at	SAMD3	6q23.1	4.15	LB
227019_at	LOC400793	1q23.3	4.10	LB
229552_at	LOC283454	12q24.23	4.09	LB
1568955_at	1568955_at	1	4.08	LB
213411_at	ADAM22	7q21	4.02	LB
215734_at	C19orf36	19p13.3	4.01	LB
202986_at	ARNT2	15q24	4.01	LB
229070_at	C6orf105	6p24.1	3.99	LB
222777_s_at	WHSC1	4p16.3	23.19	MS
204379_s_at	FGFR3	4p16.3	18.32	MS
217867_x_at	BACE2	21q22.3	11.94	MS
217901_at	DSG2	18q12.1	11.61	MS
211709_s_at	CLEC11A	19q13.3	10.53	MS
204066_s_at	CENTG2	2p24.3-p24.1	10.19	MS
214156_at	MYRIP	3p22.2-p22.1	9.26	MS
210220_at	FZD2	17q21.1	8.99	MS
212190_at	SERPINE2	2q33-q35	8.93	MS
204749_at	NAP1L3	Xq21.3-q22	8.87	MS
212771_at	C10orf38	10p13	8.62	MS
212813_at	JAM3	11q25	8.42	MS
241703_at	RPIB9	7q21.12	8.24	MS
232235_at	C18orf4	18q22.1	8.09	MS
217963_s_at	NGFRAP1	Xq22.2	7.99	MS
215047_at	TRIM58	1q44	7.97	MS

224325_at	<i>FZD8</i>	10p11.21	7.88	MS
212148_at	<i>PBX1</i>	1q23	7.66	MS
205559_s_at	<i>PCSK5</i>	9q21.3	7.56	MS
205830_at	<i>CLGN</i>	4q28.3-q31.1	7.55	MS
219440_at	<i>RAI2</i>	Xp22	7.50	MS
218345_at	<i>HCA112</i>	7q36.1	7.47	MS
212686_at	<i>PPM1H</i>	12q14.1-q14.2	7.44	MS
227195_at	<i>ZNF503</i>	10q22.2	7.44	MS
201360_at	<i>CST3</i>	20p11.21	7.35	MS
218775_s_at	<i>BOMB</i>	4q35.1	7.33	MS
227084_at	<i>DTNA</i>	18q12	7.26	MS
229782_at	<i>NCRMS</i>	12q23.1	7.21	MS
219631_at	<i>LRP12</i>	8q22.2-q23.1	7.18	MS
205011_at	<i>LOH11CR2A</i>	11q23	7.17	MS
227875_at	<i>KLHL13</i>	Xq23-q24	7.16	MS
241535_at	<i>241535_at</i>	2p25.3	7.13	MS
226066_at	<i>MITF</i>	3p14.2-p14.1	7.12	MS
229874_x_at	<i>LOC400741</i>	1p36.13	7.09	MS
220991_s_at	<i>RNF32</i>	7q36.3	7.03	MS
227290_at	<i>227290_at</i>	16q23.2	7.00	MS
226677_at	<i>ZNF521</i>	18q11.2	6.97	MS
227742_at	<i>CLIC6</i>	21q22.12	6.93	MS
202599_s_at	<i>NRIP1</i>	21q11.2	6.84	MS
204011_at	<i>SPRY2</i>	13q31.1	6.69	MS
211276_at	<i>TCEAL2</i>	Xq22.1-q22.3	6.66	MS
218675_at	<i>SLC22A17</i>	14q11.2	6.65	MS
201976_s_at	<i>MYO10</i>	5p15.1-p14.3	6.64	MS
200897_s_at	<i>KIAA0992</i>	4q32.3	6.63	MS
224955_at	<i>TEAD1</i>	11p15.4	6.61	MS
221047_s_at	<i>MARK1</i>	1q41	6.60	MS
227341_at	<i>C10orf30</i>	10p13	6.60	MS
220532_s_at	<i>LR8</i>	7q36.1	6.58	MS
236565_s_at	<i>LARP6</i>	15q23	6.54	MS
225474_at	<i>MAGI1</i>	3p14.1	6.48	MS
204115_at	<i>GNG11</i>	7q31-q32	12.48	HP
232352_at	<i>ISL2</i>	15q23	12.00	HP
202688_at	<i>TNFSF10</i>	3q26	11.93	HP
225698_at	<i>TIGA1</i>	5q21-q22	10.19	HP
224961_at	<i>SCYL2</i>	12q23.1	10.18	HP
209595_at	<i>GTF2F2</i>	13q14	10.00	HP
230479_at	<i>230479_at</i>	11p15.4	9.35	HP
224767_at	<i>RPL37</i>	5p13	9.29	HP
203620_s_at	<i>FCHSD2</i>	11q13.4	9.15	HP
211434_s_at	<i>CCRL2</i>	3p21	9.03	HP
226507_at	<i>PAK1</i>	11q13-q14	9.03	HP
219734_at	<i>SIDT1</i>	3q13.2	9.02	HP
227180_at	<i>ELOVL7</i>	5q12.1	8.92	HP
203989_x_at	<i>F2R</i>	5q13	8.91	HP
208745_at	<i>ATP5L</i>	11q23.3	8.86	HP
244632_at	<i>CNTN5</i>	11q21-q22.2	8.81	HP
227056_at	<i>KIAA0141</i>	5q31.3	8.74	HP
203698_s_at	<i>FRZB</i>	2qter	8.71	HP
230425_at	<i>EPHB1</i>	3q21-q23	8.70	HP
222943_at	<i>GBA3</i>	4p15.31	8.63	HP
229050_s_at	<i>MGC16037</i>	9q34.3	8.63	HP
219122_s_at	<i>ICF45</i>	5q33.3	8.62	HP
227979_at	<i>RBM30</i>	11q13	8.59	HP
222793_at	<i>DDX58</i>	9p12	8.53	HP
234339_s_at	<i>GLTSCR2</i>	19q13.3	8.48	HP

224610_at	224610_at	11q12.3	8.47	HP
204236_at	<i>FLI1</i>	11q24.1-q24.3	8.38	HP
235228_at	<i>KIAA1912</i>	2p16.1	8.30	HP
238039_at	238039_at	5q14.1	8.28	HP
216348_at	<i>RPS17</i>	15q	8.27	HP
217969_at	<i>C11orf2</i>	11q13	8.24	HP
210813_s_at	<i>XRCC4</i>	5q13-q14	8.23	HP
213554_s_at	<i>H41</i>	3q22.1	8.22	HP
203528_at	<i>SEMA4D</i>	9q22-q31	8.15	HP
200715_x_at	<i>RPL13A</i>	19q13.3	8.10	HP
227867_at	<i>LOC129293</i>	2p11.2	8.09	HP
242625_at	<i>RSAD2</i>	2p25.2	8.08	HP
220201_at	<i>MNAB</i>	9q34	7.95	HP
220768_s_at	<i>CSNK1G3</i>	5q23	7.93	HP
206486_at	<i>LAG3</i>	12p13.32	7.86	HP
205632_s_at	<i>PIP5K1B</i>	9q13	7.86	HP
234875_at	<i>RPL7A</i>	9q34	7.8	HP
228607_at	<i>OAS2</i>	12q24.2	7.75	HP
206991_s_at	<i>CCR5</i>	3p21	7.72	HP
224915_x_at	<i>TALDO1</i>	11p15.5-p15.4	7.72	HP
216399_s_at	<i>ZNF291</i>	15q24	7.63	HP
200995_at	<i>IPO7</i>	11p15.4	7.63	HP
218258_at	<i>POLR1D</i>	13q12.2	7.57	HP
222108_at	<i>AMIGO2</i>	12q13.11	7.56	HP
205569_at	<i>LAMP3</i>	3q26.3-q27	7.52	HP
1553027_a_at	<i>KLHL4</i>	Xq21.3	8.21	CD-1
229309_at	<i>ADRB1</i>	10q24-q26	7.87	CD-1
210587_at	<i>INHBE</i>	12q13.3	7.74	CD-1
203423_at	<i>RBP1</i>	3q23	7.57	CD-1
233500_x_at	<i>CLEC2D</i>	12p13	7.52	CD-1
209921_at	<i>SLC7A11</i>	4q28-q32	6.92	CD-1
213524_s_at	<i>G0S2</i>	1q32.2-q41	6.65	CD-1
212713_at	<i>MFAP4</i>	17p11.2	6.59	CD-1
220993_s_at	<i>GPR63</i>	6q16.1-q16.3	6.53	CD-1
227915_at	<i>ASB2</i>	14q31-q32	6.20	CD-1
228737_at	<i>C20orf100</i>	20q13.12	5.98	CD-1
210105_s_at	<i>FYN</i>	6q21	5.93	CD-1
228367_at	<i>ALPK2</i>	18q21.31	5.88	CD-1
206631_at	<i>PTGER2</i>	14q22	5.85	CD-1
222116_s_at	<i>TBC1D16</i>	17q25.3	5.84	CD-1
217127_at	<i>CTH</i>	1p31.1	5.77	CD-1
221539_at	<i>EIF4EBP1</i>	8p12	5.69	CD-1
219892_at	<i>TM6SF1</i>	15q24-q26	5.53	CD-1
227353_at	227353_at	17q25.3	5.48	CD-1
239062_at	239062_at	17q25.3	5.48	CD-1
232054_at	<i>PCDH20</i>	13q21	5.42	CD-1
212501_at	<i>CEBPB (NF-IL6)</i>	20q13.1	5.25	CD-1
204328_at	<i>EVER1</i>	17q25.3	5.24	CD-1
201036_s_at	<i>HADHSC</i>	4q22-q26	5.21	CD-1
243816_at	<i>ZNF70</i>	22q11.2	5.20	CD-1
202847_at	<i>PCK2</i>	14q11.2	5.15	CD-1
224839_s_at	<i>GPT2</i>	16q12.1	5.10	CD-1
227134_at	<i>SYTL1</i>	1p36.11	5.10	CD-1
201397_at	<i>PHGDH</i>	1p12	5.07	CD-1
204114_at	<i>NID2</i>	14q21-q22	5.03	CD-1
227117_at	<i>XPOT</i>	12q14.2	5.02	CD-1
223195_s_at	<i>SESN2</i>	1p35.3	5.01	CD-1
238983_at	<i>FLJ14001</i>	4p14	5.01	CD-1
204999_s_at	<i>ATF5</i>	19q13.3	4.88	CD-1

206039_at	<i>RAB33A</i>	Xq25	4.80	CD-1
217763_s_at	<i>RAB31</i>	18p11.3	4.76	CD-1
235522_at	<i>OCIL</i>	12p13	4.69	CD-1
227823_at	<i>RGAG4</i>	Xq13.1	4.68	CD-1
218145_at	<i>TRIB3</i>	20p13-p12.2	4.67	CD-1
206956_at	<i>BGLAP</i>	1q25-q31	4.61	CD-1
220358_at	<i>SNFT</i>	1q32.3	4.58	CD-1
228024_at	<i>228024_at</i>	8p22	4.56	CD-1
239045_at	<i>ERN1</i>	17q24.2	4.5	CD-1
212767_at	<i>GTPBP7</i>	10q26.3	4.5	CD-1
203438_at	<i>STC2</i>	5q35.1	4.49	CD-1
213475_s_at	<i>ITGAL</i>	16p11.2	4.44	CD-1
227630_at	<i>PPP2R5E</i>	14q23.1	4.43	CD-1
204971_at	<i>CSTA</i>	3q21	4.39	CD-1
236554_x_at	<i>EVER2</i>	17q25.3	4.39	CD-1
240094_at	<i>DJ971N18.2</i>	20p12	4.37	CD-1
228592_at	<i>MS4A1</i>	11q12	10.95	CD-2
220068_at	<i>VPREB3</i>	22q11.23	10.83	CD-2
227336_at	<i>DTX1</i>	12q24.13	10.4	CD-2
1558185_at	<i>CLU1</i>	12q22	8.93	CD-2
236065_at	<i>236065_at</i>	12q12	8.57	CD-2
212807_s_at	<i>SORT1</i>	1p21.3-p13.1	8.45	CD-2
1556583_a_at	<i>1556583_a_at</i>	2p22.1	7.85	CD-2
221969_at	<i>PAX5</i>	9p13	7.72	CD-2
213324_at	<i>SRC</i>	20q12-q13	6.94	CD-2
212158_at	<i>SDC2</i>	8q22-q23	6.92	CD-2
225955_at	<i>METRNL</i>	17q25.3	6.89	CD-2
219670_at	<i>C1orf165</i>	1p33	6.78	CD-2
213436_at	<i>CNR1</i>	6q14-q15	6.75	CD-2
208703_s_at	<i>APLP2</i>	11q23-q25	6.67	CD-2
238452_at	<i>FCRLM2</i>	1q23.3	6.66	CD-2
212678_at	<i>NF1</i>	17q11.2	6.62	CD-2
205901_at	<i>PNOC</i>	8p21	6.58	CD-2
206150_at	<i>TNFRSF7 (CD27L)</i>	12p13	6.57	CD-2
212226_s_at	<i>PPAP2B</i>	1pter-p22.1	6.55	CD-2
225897_at	<i>MARCKS</i>	6q22.2	6.51	CD-2
216840_s_at	<i>LAMA2</i>	6q22-q23	6.45	CD-2
208890_s_at	<i>PLXNB2</i>	22q13.33	6.44	CD-2
207511_s_at	<i>C2orf24</i>	2q35	6.38	CD-2
243780_at	<i>243780_at</i>	22q13.2	6.36	CD-2
205049_s_at	<i>CD79A</i>	19q13.2	6.35	CD-2
204759_at	<i>RCBTB2</i>	13q14.3	6.33	CD-2
213407_at	<i>PHLPPL</i>	16q22.2	6.15	CD-2
1560225_at	<i>1560225_at</i>	6q15	6.14	CD-2
238893_at	<i>LOC338758</i>	12q21.33	6.09	CD-2
204208_at	<i>RNGTT</i>	6q16	6.02	CD-2
238282_at	<i>238282_at</i>	3q26.32	5.95	CD-2
227396_at	<i>227396_at</i>	11p11.2	5.91	CD-2
235764_at	<i>235764_at</i>	4q27	5.83	CD-2
202969_at	<i>DYRK2</i>	12q15	5.82	CD-2
211812_s_at	<i>B3GALT3</i>	3q25	5.77	CD-2
232693_s_at	<i>ZNF395</i>	8p21.1	5.76	CD-2
204352_at	<i>TRAF5</i>	1q32	5.76	CD-2
219888_at	<i>SPAG4</i>	20q11.21	5.75	CD-2
226325_at	<i>ADSSL1</i>	14q32.33	5.74	CD-2
226459_at	<i>PIK3AP1</i>	10q24.1	5.66	CD-2
220595_at	<i>PDZRN4</i>	12q12	5.65	CD-2
229817_at	<i>ZNF608</i>	5q23.2	5.65	CD-2
210258_at	<i>RGS13</i>	1q31.2	5.64	CD-2

203313_s_at	<i>TGIF</i>	18p11.3	5.64	CD-2
218231_at	<i>NAGK</i>	2p13.3	5.63	CD-2
226440_at	<i>DUSP22</i>	6p25.3	5.63	CD-2
207216_at	<i>TNFSF8 (CD30L)</i>	9q33	5.63	CD-2
200940_s_at	<i>RERE</i>	1p36.1-p36.2	5.62	CD-2
227094_at	<i>DHTKD1</i>	10p14	5.61	CD-2
203723_at	<i>ITPKB</i>	1q42.13	5.57	CD-2
205898_at	<i>CX3CR1</i>	3p21	14.68	MF
1554485_s_at	<i>TMEM37</i>	2q14.2	14.14	MF
204589_at	<i>NUAK1 (ARK5)</i>	12q23.3	13.46	MF
209167_at	<i>GPM6B</i>	Xp22.2	12.32	MF
221060_s_at	<i>TLR4</i>	9q32-q33	12.25	MF
205718_at	<i>ITGB7</i>	12q13.13	11.92	MF
215294_s_at	<i>SMARCA1</i>	Xq25	11.86	MF
213138_at	<i>ARID5A</i>	2q11.2	11.75	MF
221796_at	<i>NTRK2</i>	9q22.1	10.31	MF
51158_at	<i>LOC400451</i>	15q26.1	10.27	MF
209875_s_at	<i>SPP1</i>	4q21-q25	9.99	MF
1570445_a_at	<i>1570445_a_at</i>	5q35.2	9.42	MF
1569785_at	<i>1569785_at</i>	18q12.3	9.40	MF
235740_at	<i>MCTP1</i>	5q15	9.37	MF
205862_at	<i>GREB1</i>	2p25.1	9.35	MF
207638_at	<i>PRSS7</i>	21q21	9.32	MF
200762_at	<i>DPYSL2</i>	8p22-p21	9.31	MF
224583_at	<i>COTL1</i>	16q24.1	9.09	MF
203658_at	<i>SLC25A20</i>	3p21.31	8.79	MF
203119_at	<i>MGC2574</i>	11q12.2	8.72	MF
235416_at	<i>LOC441117</i>	5q35.2	8.70	MF
207356_at	<i>DEFB4</i>	8p23.1-p22	8.59	MF
209980_s_at	<i>SHMT1</i>	17p11.2	8.48	MF
203417_at	<i>MFAP2</i>	1p36.1-p35	8.27	MF
225868_at	<i>TRIM47</i>	17q25.1	8.23	MF
221887_s_at	<i>DFNB31</i>	9q32-q34	8.13	MF
203973_s_at	<i>CEBPD</i>	8p11.2-p11.1	8.01	MF
212098_at	<i>LOC151162</i>	2q21.3	7.96	MF
226436_at	<i>RASSF4</i>	10q11.21	7.89	MF
228123_s_at	<i>C20orf22</i>	20p11.21	7.46	MF
201131_s_at	<i>CDH1 (E-cadherin)</i>	16q22.1	7.43	MF
201954_at	<i>ARPC1B</i>	7q22.1	7.42	MF
219895_at	<i>FAM70A</i>	Xq24	7.41	MF
222930_s_at	<i>AGMAT</i>	1p36.13	7.39	MF
226545_at	<i>CD109</i>	6q13	7.30	MF
206023_at	<i>NMU</i>	4q12	7.26	MF
219310_at	<i>C20orf39</i>	20p11.21	7.22	MF
214023_x_at	<i>TUBB-PARALOG</i>	6p25	7.16	MF
204858_s_at	<i>ECGF1</i>	22q13	7.15	MF
205003_at	<i>DOCK4</i>	7q31.1	7.10	MF
209687_at	<i>CXCL12 (SDF-1)</i>	10q11.1	7.08	MF
200965_s_at	<i>ABLIM1</i>	10q25	7.06	MF
206167_s_at	<i>ARHGAP6</i>	Xp22.3	7.02	MF
213566_at	<i>RNASE6</i>	14q11.2	7.02	MF
211941_s_at	<i>PBP</i>	12q24.23	7.00	MF
218608_at	<i>ATP13A2</i>	1p36	6.99	MF
210873_x_at	<i>APOBEC3A</i>	22q13.1-q13.2	6.88	MF
201481_s_at	<i>PYGB</i>	20p11.2-p11.1	6.80	MF
205269_at	<i>LCP2</i>	5q33.1-qter	6.79	MF
223121_s_at	<i>SFRP2</i>	4q31.3	6.72	MF