

<b>Affymetrix Probe Set</b>	<b>Gene Name / Probe Set</b>	<b>Chromosome Map Position</b>	<b>SAM Score( + higher in CD-1)</b>
216438_s_at	<i>TMSB4X</i>	Xq21.3-q22	4.91
200849_s_at	<i>AHCYL1</i>	1p13.2	4.40
223851_s_at	<i>TNFRSF18</i>	1p36.3	4.37
239533_at	<i>GPR155</i>	2q31.1	4.36
210401_at	<i>P2RX1</i>	17p13.3	4.27
220313_at	<i>GPR88</i>	1p21.3	4.27
206759_at	<i>FCER2</i>	19p13.3	4.18
204912_at	<i>IL10RA</i>	11q23	4.11
219667_s_at	<i>BANK1</i>	4q24	3.99
223562_at	<i>PARVG</i>	22q13.2-q13	3.95
233059_at	<i>KCNJ3</i>	2q24.1	3.93
209210_s_at	<i>PLEKHC1</i>	14q22.1	3.85
228583_at	<i>C14orf46</i>	14q24.3	3.80
230466_s_at	<i>230466_s_at</i>	chr12	3.79
205124_at	<i>MEF2B</i>	19p12	3.74
219109_at	<i>SPAG16</i>	2q34	3.73
204174_at	<i>ALOX5AP</i>	13q12	3.73
227786_at	<i>THRAP6</i>	8q24.11	3.70
201963_at	<i>ACSL1</i>	4q34-q35	3.68
214180_at	<i>MAN1C1</i>	1p35	3.56
205773_at	<i>CPEB3</i>	10q23.32	3.53
225166_at	<i>ARHGAP18</i>	6q22.33	3.43
227167_s_at	<i>227167_s_at</i>	chr12	3.34
235518_at	<i>SLC8A1</i>	2p23-p22	3.29
213304_at	<i>KIAA0423</i>	14q21.3	3.29
235473_at	<i>235473_at</i>	chr14	3.25
212636_at	<i>QKI</i>	6q26-27	3.22
218224_at	<i>PNMA1</i>	14q24.3	3.21
229512_at	<i>229512_at</i>	chr15	3.18
213251_at	<i>213251_at</i>	4q31.21	3.18
223228_at	<i>LDOC1L</i>	22q13.31	3.14
1560156_at	<i>1560156_at</i>	chr12	3.11
1563209_a_at	<i>C20orf133</i>	20p12.1	3.11
231944_at	<i>ERO1LB</i>	1q42.2-q43	3.08
209035_at	<i>MDK</i>	11p11.2	3.07
213340_s_at	<i>KIAA0495</i>	1p36.32	3.07
213400_s_at	<i>TBL1X</i>	Xp22.3	3.03
226003_at	<i>KIF21A</i>	12q12	3.02
235802_at	<i>PLD4</i>	14q32.33	3.02
225733_at	<i>B3GALT6</i>	1p36.33	3.01
202561_at	<i>TNKS</i>	8p23.1	2.97
230154_at	<i>WAC</i>	chr10	2.97
222982_x_at	<i>SLC38A2</i>	12q	2.95
230233_at	<i>RASGEF1B</i>	4q21.3	2.94
226126_at	<i>MGC16169</i>	4q24	2.93
229018_at	<i>FLJ22789</i>	12q21.31	2.92
203302_at	<i>DCK</i>	4q13.3-q21.1	2.91
1566433_at	<i>C14orf175</i>	14q32.33	2.88
219559_at	<i>C20orf59</i>	20q13.33	2.86
209431_s_at	<i>ZNF278</i>	22q12.2	2.86
228254_at	<i>STAM2</i>	2q23.3	2.84
240572_s_at	<i>LOC374443</i>	12p13.31	2.80
1557558_s_at	<i>MATN1</i>	1p35	2.79
225053_at	<i>CNOT7</i>	8p22-p21.3	2.79
227627_at	<i>SGK3</i>	8q12.3-8q13.1	2.78
221588_x_at	<i>ALDH6A1</i>	14q24.3	2.77
228135_at	<i>C1orf52</i>	1p22.3	2.72
224830_at	<i>NUDT21</i>	16q12.2	2.70
203117_s_at	<i>USP52</i>	12q13.2-q13.3	2.68
200902_at	<i>200902_at</i>	1p31	2.67
205702_at	<i>PHTF1</i>	1p13	2.67

228328_at	228328_at	chr14	2.65
213410_at	C10orf137	10q26.13	2.64
236235_at	ITCH	20q11.22	2.64
223823_at	KCNMB2	3q26.2-q27.1	2.63
201425_at	ALDH2	12q24.2	2.63
235698_at	ZFP90	16q22.1	2.63
209085_x_at	RFC1	4p14-p13	2.61
215629_s_at	DLEU2	13q14.3	2.60
224886_at	LOC339123	16p13.3	2.60
223569_at	PPAPDC1B	8p12	2.58
223275_at	HRMT1L6	1p13.3	2.58
201969_at	NASP	1p34.1	2.56
229614_at	ZFPL	19q13.41	2.56
221675_s_at	CHPT1	12q	2.56
210792_x_at	SIVA	14q32.33	2.54
242239_at	NSUN6	10p12.31	2.54
204568_at	KIAA0831	14q22.2	2.53
213677_s_at	PMS1	2q31-q33 2q31.1	2.53
214259_s_at	AKR7A2	1p35.1-p36.23	2.53
212930_at	ATP2B1	12q21.3	2.52
221788_at	PGM3	6q14.1-q15	2.51
228630_at	ZNF84	12q24.33	2.50
204224_s_at	GCH1	14q22.1-q22.2	2.50
227095_at	LEPR	1p31	2.48
221478_at	BNIP3L	8p21	2.48
212462_at	MYST4	10q22.2	2.46
209682_at	CBLB	3q13.11	2.46
229145_at	C10orf104	10q22.1	2.45
205296_at	RBL1	20q11.2	2.45
1554155_at	MCPH1	8p23.1	2.43
228603_at	ACTR3	2q14.1	2.41
219920_s_at	GMPPB	3p21.31	2.40
1557433_at	1557433_at		2.40
211208_s_at	CASK	Xp11.4	2.35
212331_at	RBL2	16q12.2	2.35
218176_at	MAGEF1	3q13	2.34
1559975_at	BTG1	12q22	2.33
221031_s_at	DKFZP434F0318	12p13.2	2.33
216080_s_at	FADS3	11q12-q13.1	2.31
235251_at	235251_at	chr03	-2.44
208837_at	TMED3	15q24-q25	-2.44
225973_at	TAP2	6p21.3	-2.51
241781_at	C9orf41	9q21.13	-2.56
211763_s_at	UBE2B	5q23-q31	-2.58
209065_at	UQCRB	8q22	-2.60
218501_at	ARHGEF3	3p21-p13	-2.60
1557915_s_at	GSTO1	10q25.1	-2.61
205230_at	RPH3A	12q24.13	-2.61
202431_s_at	MYC	8q24.12-q24.13	-2.65
227492_at	227492_at	chr05	-2.66
225863_s_at	C19orf12	19q12	-2.71
219532_at	ELOVL4	6q14	-2.71
215071_s_at	HIST1H2AC	6p21.3	-2.72
1552277_a_at	MGC17337	9q31.1	-2.73
234725_s_at	SEMA4B	15q25	-2.74
212820_at	DMXL2	15q21.2	-2.76
207996_s_at	C18orf1	18p11.2	-2.83
214290_s_at	HIST2H2AA	1q21.2	-2.93
202329_at	CSK	15q23-q25	-2.95
222538_s_at	APPL	3p21.1-p14.3	-3.04
202604_x_at	ADAM10	15q2 15q22	-3.08
223193_x_at	E2IG5	3q21.1	-3.08
225634_at	ZC3HAV1	7q34	-3.10

221004_s_at	<i>ITM2C</i>	2q37	-3.12
210448_s_at	<i>P2RX5</i>	17p13.3	-3.17
1560661_x_at	<i>LOC150759</i>	2q11.2	-3.21
205590_at	<i>RASGRP1</i>	15q15	-3.26
1554834_a_at	<i>RASSF5</i>	1q32.1	-3.29
211043_s_at	<i>CLTB</i>	4q2-q3 5q35	-3.30
201051_at	<i>ANP32A</i>	15q22.3-q23	-3.32
211676_s_at	<i>IFNGR1</i>	6q23-q24	-3.33
235581_at	<i>235581_at</i>	chr07	-3.53
207677_s_at	<i>NCF4</i>	22q13.1	-3.54
223704_s_at	<i>DMRT2</i>	9p24.3	-3.62
217865_at	<i>RNF130</i>	5q35.3	-3.65
238681_at	<i>GDPD1</i>	17q23.2	-3.65
206472_s_at	<i>TLE3</i>	15q22	-3.66
63825_at	<i>ABHD2</i>	15q26.1	-3.71
204779_s_at	<i>HOXB7</i>	17q21.3	-3.72
232429_at	<i>232429_at</i>	chr15	-3.73
205352_at	<i>SERPINI1</i>	3q26.1	-3.77
208788_at	<i>ELOVL5</i>	6p21.1-p12.1	-3.77
202732_at	<i>PKIG</i>	20q12-q13.1	-3.85
227099_s_at	<i>LOC387763</i>	11p11.2	-3.93
213523_at	<i>CCNE1</i>	19q12	-3.94
232181_at	<i>LOC153346</i>	5q33.1	-3.94
213552_at	<i>GLCE</i>	15q23	-4.00
208669_s_at	<i>CRI1</i>	15q21.1-q21.2	-4.06
217489_s_at	<i>IL6R</i>	1q21	-4.16
224851_at	<i>CDK6</i>	7q21-q22	-4.34
221973_at	<i>221973_at</i>	chr02	-4.41
224948_at	<i>MRPS24</i>	7p14	-4.45
217196_s_at	<i>CAMSAP1L1</i>	1q32.1	-4.73
226961_at	<i>LOC222171</i>	7p15.1	-4.80
212386_at	<i>TCF4</i>	18q21.1	-5.12
205159_at	<i>CSF2RB</i>	22q13.1	-5.47
213578_at	<i>BMPR1A</i>	10q22.3	-6.08