

Affymetrix ID	Gene Symbol	Gene Name	Fold Change
		Post Translational Modification	
215577_at	UBE2E1	UBIQUITIN-CONJUGATING ENZYME E2E 1 (UBC4/5 HOMOLOG, YEAST)	-1.6
203305_at	F13A1	COAGULATION FACTOR XIII, A1 POLYPEPTIDE	2.6
229383_at	MARCH--1	MEMBRANE-ASSOCIATED RING FINGER (C3HC4) 1	2.3
240971_x_at	CUL4A	CULLIN 4A	-2.2
210635_s_at	KLHL20	KELCH-LIKE 20 (DROSOPHILA)	-1.9
204079_at	TPST2	TYROSYLPROTEIN SULFOTRANSFERASE 2	2
		Phosphorylation	
236023_at	CDK9	CYCLIN-DEPENDENT KINASE 9 (CDC2-RELATED KINASE)	-1.8
231111_at	SCYL3	SCY1-LIKE 3 (S. CEREVISIAE)	1.6
212172_at	AK2	ADENYLATE KINASE 2	2
243829_at	BRAF	V-RAF MURINE SARCOMA VIRAL ONCOGENE HOMOLOG B1	-1.6
1554719_at	NDUFA10	NADH DEHYDROGENASE (UBIQUINONE) 1 ALPHA SUBCOMPLEX, 10, 42KDA	-2.5
229383_at	MARCH--1	MEMBRANE-ASSOCIATED RING FINGER (C3HC4) 1	2.3
240971_x_at	CUL4A	CULLIN 4A	-2.2
		Zinc Binding	
1569191_at	ZNF737	SIMILAR TO ZINC FINGER PROTEIN 93 (ZINC FINGER PROTEIN HTF34)	1.8
229383_at	MARCH--1	MEMBRANE-ASSOCIATED RING FINGER (C3HC4) 1	2.3
236533_at	DDEF1	DEVELOPMENT AND DIFFERENTIATION ENHANCING FACTOR 1	-1.5
244640_at	LOC342892	HYPOTHETICAL PROTEIN LOC342892	2.2
1555392_at	TES	TESTIS DERIVED TRANSCRIPT (3 LIM DOMAINS)	-1.9
		Nucleotide Binding	
212172_at	AK2	ADENYLATE KINASE 2	2
222479_s_at	DYNC1L1	DYNEIN, CYTOPLASMIC 1, LIGHT INTERMEDIATE CHAIN 1	-1.5
223200_s_at	LSG1	LARGE SUBUNIT GTPASE 1 HOMOLOG (S. CEREVISIAE)	-1.6
		Transmembrane	
223939_at	SUCNR1	SUCCINATE RECEPTOR 1	2.7
209147_s_at	PPAP2A	PHOSPHATIDIC ACID PHOSPHATASE TYPE 2A	1.7
229383_at	MARCH--1	MEMBRANE-ASSOCIATED RING FINGER (C3HC4) 1	2.3
214467_at	GPR65	G PROTEIN-COUPLED RECEPTOR 65	2.2
57588_at	SLC24A3	SOLUTE CARRIER FAMILY 24 (SODIUM/POTASSIUM/CALCIUM EXCHANGER), MEMBER	2.1
212014_x_at	CD44	CD44 ANTIGEN (INDIAN BLOOD GROUP)	-1.8
219666_at, 223922_x_at	MS4A6A	CD20-LIKE PRECURSOR	2.9
242108_at	TPCN1	TWO PORE SEGMENT CHANNEL 1	1.7
		Regulation of Transcription	
236023_at	CDK9	CYCLIN-DEPENDENT KINASE 9 (CDC2-RELATED KINASE)	-1.8
206118_at	STAT4	SIGNAL TRANSDUCER AND ACTIVATOR OF TRANSCRIPTION 4	1.6
209357_at	CITED2	CBP/P300-INTERACTING TRANSACTIVATOR	2
206240_s_at	ZNF136	ZINC FINGER PROTEIN 136 (CLONE PHZ-20)	-1.7
231085_s_at	PCBD2	COFACTOR OF HEPATOCYTE NUCLEAR FACTOR 1 ALPHA (TCF1) 2	-1.9
244640_at	LOC342892	HYPOTHETICAL PROTEIN LOC342892	2.2
1569191_at	ZNF737	SIMILAR TO ZINC FINGER PROTEIN 93 (ZINC FINGER PROTEIN HTF34)	1.8
		Unclassified	
244546_at	CYCS	CYTOCHROME C, SOMATIC	-1.6
215813_s_at	PTGS1	PROSTAGLANDIN-ENDOPEROXIDE SYNTHASE 1	1.9
205603_s_at	PTGS1	DIAPHANOUS HOMOLOG 2 (DROSOPHILA)	-1.6
205756_s_at	F8	COAGULATION FACTOR VIII, PROCOAGULANT COMPONENT (HEMOPHILIA A)	-1.6
217521_at	HAL	HISTIDINE AMMONIA-LYASE	1.7
243206_at	CCDC45	COILED-COIL DOMAIN CONTAINING 45	-1.6
208791_at, 208792_s_at	CLU	CLUSTERIN	2.4
225762_x_at, 225767_at	LOC284801	HYPOTHETICAL PROTEIN LOC284801	-2.2
211734_s_at	FCER1A	FC FRAGMENT OF IGE, HIGH AFFINITY I, RECEPTOR FOR; ALPHA POLYPEPTIDE	2.5
1554348_s_at	CDKN2AIPNL	SIMILAR TO RIKEN CDNA A430101B06 GENE	-1.6
202729_s_at	LTBP1	LATENT TRANSFORMING GROWTH FACTOR BETA BINDING PROTEIN 1	3
243918_at	KLHL24	KELCH-LIKE 24 (DROSOPHILA)	-1.6
238715_at	LOC400614	HYPOTHETICAL LOC400614	-1.5
211743_s_at	PRG2	PROTEOGLYCAN 2	7.5
205768_s_at, 205769_at	SLC27A2	FATTY-ACID-COENZYME A LIGASE, VERY LONG-CHAIN 1	1.7
238498_at	N/A	CONSENSUS INCLUDES GB:AA583038	-1.6
1558711_at	FAM13A1OS	FAMILY WITH SEQUENCE SIMILARITY 13, MEMBER A1 OPPOSITE STRAND	-1.6
206111_at	hCG_1998957	MAJOR HISTOCOMPATIBILITY COMPLEX	1.8
242051_at	N/A	CONSENSUS INCLUDES GB:A1695695	-1.8
210623_at	LOC51035	UNKNOWN PROTEIN LOC51035	-1.6
244515_at	PSMD7	PROTEASOME (PROSOME, MACROPAIN) 26S SUBUNIT, NON-ATPASE, 7 (MOV34 HOM	-1.9
222557_at	STMN3	STATHMIN-LIKE 3	-1.5
207067_s_at	HDC	HISTIDINE DECARBOXYLASE	3.8
202481_at	DHRS3	DEHYDROGENASE/REDUCTASE (SDR FAMILY) MEMBER 3	1.6
223809_at	RG518	REGULATOR OF G-PROTEIN SIGNALLING 18	4.4
221735_at	WDR48	WD REPEAT DOMAIN 48	-1.8
202149_at	NEDD9	NEURAL PRECURSOR CELL EXPRESSED, DEVELOPMENTALLY DOWN-REGULATED 9	2.2