Entrez Gene ID	Symbol	Gene Name	Fold Change (log2(Recurrence/NonRecurrence))
142	parp1	poly (ADP-ribose) polymerase 1	0.0553
154	ADRB2	adrenergic, beta-2-, receptor, surface	-0.0641
196	ahr	aryl hydrocarbon receptor	-0.0849
367	Ar	androgen receptor	-0.12
550	Aup1	ancient ubiquitous protein 1	0.0577
596	BCL2	B-cell CLL/lymphoma 2	-0.1569
613	bcr	breakpoint cluster region	0.0484
857	cav1	caveolin 1, caveolae protein, 22kDa	-0.0778
890	CCNA2	cyclin A2	0.0968
983	Cdk1	cell division cycle 2, G1 to S and G2 to M	0.1121
994	Cdc25b	cell division cycle 25 homolog B (S. pombe)	0.1462
995	CDC25C	cell division cycle 25 homolog C (S. pombe)	0.0472
1026	CDKN1A	cyclin-dependent kinase inhibitor 1A (p21, Cip1)	0.045
1058	CENPA	centromere protein A	0.2031
1059	Cenpb	centromere protein B, 80kDa	0.0414
1213	CLTC	clathrin, heavy chain (Hc)	0.0806
1457	CSNK2A1P	casein kinase 2, alpha 1 polypeptide pseudogene; casein kinase 2, alpha 1 polypeptide	0.0325
2064	Erbb2	v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neuro/glioblastoma derived oncogene homolog (avian)	0.0868
2099	Esr1	estrogen receptor 1	-0.2198
2101	esrra	estrogen-related receptor alpha	0.0488
2157	F8	coagulation factor VIII, procoagulant component	-0.033
2247	FGF2	fibroblast growth factor 2 (basic)	-0.0569
2353	FOS	v-fos FBJ murine osteosarcoma viral oncogene homolog	-0.1296
2810	SFN	stratifin	0.0724
2870	grk6	G protein-coupled receptor kinase 6	0.044
2885	GRB2	growth factor receptor-bound protein 2	0.0721
2932	GSK3B	glycogen synthase kinase 3 beta	0.0395
3309	HSPA5	hypothetical gene supported by AF216292; NM_005347; heat shock 70kDa protein 5 (glucose-regulated protein, 78kDa)	0.0409
3667	IRS1	insulin receptor substrate 1	-0.0863
3678	ITGA5	integrin, alpha 5 (fibronectin receptor, alpha polypeptide)	0.0537
3925	STMN1	stathmin 1	0.1115
4176	MCM7	minichromosome maintenance complex component 7	0.0915
4605	MYBL2	v-myb myeloblastosis viral oncogene homolog (avian)-like 2	0.1016
4771	Nf2	neurofibromin 2 (merlin)	0.0542

5111	pcnA	proliferating cell nuclear antigen	0.0965
5241	PGR	progesterone receptor	-0.202
5300	PIN1	peptidylprolyl cis/trans isomerase, NIMA-interacting 1	0.0291
5425	Pold2	polymerase (DNA directed), delta 2, regulatory subunit 50kDa	0.0473
5585	PKN1	protein kinase N1	0.0364
5594	MAPK1	mitogen-activated protein kinase 1	0.0411
5829	PXN	paxillin	0.0334
5888	rad51	RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae)	0.0945
6128	RPL6P27	ribosomal protein L6 pseudogene 27; ribosomal protein L6 pseudogene 19; ribosomal protein L6; ribosomal protein L6 pseudogene 10	-0.0759
6159	RPL29P12	ribosomal protein L29 pseudogene 9; ribosomal protein L29 pseudogene 12; ribosomal protein L29 pseudogene 11; ribosomal protein L29; ribosomal protein L29 pseudogene 26	-0.074
6256	RXRA	retinoid X receptor, alpha	0.0535
6696	SPP1	secreted phosphoprotein 1	0.1638
6776	STAT5A	signal transducer and activator of transcription 5A	-0.0764
6790	AURKA	aurora kinase A; aurora kinase A pseudogene 1	0.1746
7083	tk1	thymidine kinase 1, soluble	0.2329
7157	tp53	tumor protein p53	-0.0092
7186	traF2	TNF receptor-associated factor 2	0.0483
7283	TUBG1	tubulin, gamma 1; similar to Tubulin, gamma 1	0.047
7327	UBE2G2	ubiquitin-conjugating enzyme E2G 2 (UBC7 homolog, yeast)	0.0289
7337	Ube3a	ubiquitin protein ligase E3A	-0.0377
7430	EZR	hypothetical protein LOC100129652; ezrin	0.0596
7529	YWHAB	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, beta polypeptide	0.0257
7531	YWHAE	similar to 14-3-3 protein epsilon (14-3-3E) (Mitochondrial import stimulation factor L subunit) (MSF L); tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide	-0.0159
7534	YWHAZ	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide	0.1085
7704	ZBTB16	zinc finger and BTB domain containing 16	-0.161
7706	TRIM25	tripartite motif-containing 25	0.0557
8318	Cdc45	CDC45 cell division cycle 45-like (S. cerevisiae)	0.1407
8648	NCOA1	nuclear receptor coactivator 1	-0.0309
9024	BRSK2	BR serine/threonine kinase 2	0.0259
9088	PKMYT1	protein kinase, membrane associated tyrosine/threonine 1	0.189

9133	CCNB2	cyclin B2	0.2096
9212	AURKB	aurora kinase B	0.144
9368	SLC9A3R1	solute carrier family 9 (sodium/hydrogen exchanger), member 3 regulator 1	0.0883
9513	FXR2	fragile X mental retardation, autosomal homolog 2	-0.001
9612	ncor2	nuclear receptor co-repressor 2	0.0565
10376	TUBA1B	hypothetical gene supported by AF081484; NM_006082; tubulin, alpha 1b	0.0673
10521	DDX17	DEAD (Asp-Glu-Ala-Asp) box polypeptide 17	-0.0832
10563	CXCL13	chemokine (C-X-C motif) ligand 13	-0.1864
10733	PLK4	polo-like kinase 4 (Drosophila)	0.0601
11188	nisch	nischarin	-0.052
23594	ORC6L	origin recognition complex, subunit 6 like (yeast)	0.1505
51053	gmnn	geminin, DNA replication inhibitor	0.046
140890	SFRS12	splicing factor, arginine/serine-rich 12	-0.0615