

Entrez Gene ID	Symbol	Gene Name	Fold Change (log2(Recurrence/N onRecurrence))
332	BIRC5	baculoviral IAP repeat-containing 5	0.525534741
604	BCL6	B-cell CLL/lymphoma 6	-0.324826297
672	BRCA1	breast cancer 1, early onset	0.168514568
861	RUNX1	runt-related transcription factor 1	-0.425389522
891	CCNB1	cyclin B1	0.553332943
894	CCND2	cyclin D2	-0.279525094
983	CDC2	cell division cycle 2, G1 to S and G2 to M	0.533405133
995	CDC25C	cell division cycle 25 homolog C (S. pombe)	0.237402042
1019	CDK4	cyclin-dependent kinase 4	0.19392358
1386	ATF2	activating transcription factor 2	0.187614742
1894	ECT2	epithelial cell transforming sequence 2 oncogene	0.490921486
2002	ELK1	ELK1, member of ETS oncogene family	-0.224147538
2022	ENG	endoglin	-0.287278403
2353	FOS	v-fos FBJ murine osteosarcoma viral oncogene homolog	-0.884614147
2885	GRB2	growth factor receptor-bound protein 2	0.302956377
2887	GRB10	growth factor receptor-bound protein 10	-0.419307606
2934	GSN	gelsolin (amyloidosis, Finnish type)	-0.414611209
3066	HDAC2	histone deacetylase 2	0.336448302
3725	JUN	jun oncogene	-0.25073622
3726	JUNB	jun B proto-oncogene	-0.381292422
4242	MFNG	MFNG O-fucosylpeptide 3-beta-N-acetylglucosaminyltransferase	-0.220544272
4582	MUC1	mucin 1, cell surface associated	-0.540474233
4851	NOTCH1	Notch homolog 1, translocation-associated (Drosophila)	-0.251020211
5296	PIK3R2	phosphoinositide-3-kinase, regulatory subunit 2 (beta)	-0.170435853
5310	PKD1	polycystic kidney disease 1 (autosomal dominant)	-0.265291721
5347	PLK1	polo-like kinase 1 (Drosophila)	0.132883555
5371	PML	promyelocytic leukemia; similar to promyelocytic leukemia protein isoform 1	-0.190408636
5468	PPARG	peroxisome proliferator-activated receptor gamma	-0.467009362
5594	MAPK1	mitogen-activated protein kinase 1	0.310605668
5603	MAPK13	mitogen-activated protein kinase 13	0.390618994
5685	PSMA4	proteasome (prosome, macropain) subunit, alpha type, 4	0.199878791
5713	PSMD7	proteasome (prosome, macropain) 26S subunit, non-ATPase, 7	0.333535736
5716	PSMD10	proteasome (prosome, macropain) 26S subunit, non-ATPase, 10	0.249477702
5984	RFC4	replication factor C (activator 1) 4, 37kDa	0.303129653

6777	STAT5B	signal transducer and activator of transcription 5B	-0.215670749
6790	AURKA	aurora kinase A; aurora kinase A pseudogene 1 similar to G/T mismatch-specific thymine DNA	0.586894665
6996	TDG	glycosylase; thymine-DNA glycosylase	0.30849646
7040	TGFB1	transforming growth factor, beta 1	-0.287566795
7048	TGFBR2	transforming growth factor, beta receptor II (70/80kDa)	-0.245458396
7431	VIM	vimentin	-0.200376664
7535	ZAP70	zeta-chain (TCR) associated protein kinase 70kDa	-0.199687398
8318	CDC45L	CDC45 cell division cycle 45-like (<i>S. cerevisiae</i>) mitogen-activated protein kinase-activated protein	0.374948834
8550	MAPKAPK5	kinase 5	0.185703912
8805	TRIM24	tripartite motif-containing 24	0.304387027
9133	CCNB2	cyclin B2	0.441986629
9612	NCOR2	nuclear receptor co-repressor 2	-0.115717718
10482	NXF1	nuclear RNA export factor 1	-0.240313133
10617	STAMBP	STAM binding protein	0.306671217
10856	RUVBL2	RuvB-like 2 (<i>E. coli</i>)	0.25485086
11171	STRAP	serine/threonine kinase receptor associated protein	0.273783403
22927	HABP4	hyaluronan binding protein 4	-0.221994708
23043	TNIK	TRAF2 and NCK interacting kinase	-0.426386363
23225	NUP210	nucleoporin 210kDa	-0.412124352
55165	CEP55	centrosomal protein 55kDa	0.460341582
55568	GALNT10	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 10 (GalNAc-T10)	-0.519407666
55759	WDR12	WD repeat domain 12	0.227374836
55872	PBK	PDZ binding kinase	0.545601185
