

Additional file 5

File format: DOC

Title: Genes selected by supervised principal component procedure

Description:

<b>Gene</b>	<b>p-value</b>	<b>Hazard Ratio</b>	<b>Lower 95% CL</b>	<b>Upper 95% CL</b>
MELK*	<.0001	2.32	1.55	3.46
UBE2C	0.0004	2.17	1.41	3.33
DIAPH3*	0.0002	2.04	1.41	2.95
PKMYT1*	0.0001	2.01	1.40	2.88
CONTIG63649RC	0.0003	1.89	1.34	2.66
NUSAP1	0.0026	1.83	1.24	2.71
MYBL2*	0.0005	1.82	1.30	2.55
CENPA*	0.0005	1.81	1.29	2.52
ESPL1	0.0027	1.79	1.23	2.62
CONTIG46218RC	0.0029	1.79	1.22	2.61
SPOCD1	0.0035	1.74	1.20	2.53
FBXO31	0.0012	1.74	1.24	2.43
TK1*	0.0061	1.71	1.17	2.51
MMP9	0.0041	1.71	1.18	2.47
DEPDC1B	0.0033	1.70	1.19	2.43
PRR11*	0.0013	1.69	1.23	2.34
UBE2S*	0.0011	1.69	1.23	2.31
RFC4*	0.0010	1.68	1.23	2.28
BUB1*	0.0158	1.66	1.10	2.52
CCNE2	0.0080	1.65	1.14	2.39
SERF1A	0.0006	1.61	1.23	2.12
RHBDF2	0.0048	1.59	1.15	2.19
ORC6L*	0.0037	1.58	1.16	2.16
RACGAP1*	0.0115	1.56	1.10	2.19
KIF2C	0.0162	1.55	1.08	2.23
CENPN	0.0190	1.52	1.07	2.16
CCNB1*	0.0305	1.50	1.04	2.16
DTL	0.0246	1.49	1.05	2.10
EBF4	0.0253	1.47	1.05	2.07
DC13*	0.0136	1.46	1.08	1.98
QSOX2	0.0276	1.42	1.04	1.94
SLC2A3	0.0219	1.41	1.05	1.90
STK32B	0.0364	0.73	0.54	0.98
NAT1	0.0136	0.70	0.53	0.93
GSTM1	0.0324	0.67	0.46	0.97
NTN4	0.0049	0.64	0.46	0.87
ZNF533	0.0036	0.59	0.42	0.84

\*genes in the 14-gene model from the 37-gene model reduced by LASSO via LARS