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Gene symbol	parametric p-value	FDR	permutation p-value	hazards ratio
TNRC6A	0.001660	0.202	< 1e-07	0.135
RPL13	0.002913	0.202	0.005	0.150
NPIPL3	0.008794	0.307	0.009	0.190
ANKRD50	0.010858	0.307	0.015	0.375
GRHL1	0.012903	0.307	0.011	0.293
ENTPD4	0.014215	0.307	0.023	0.222
TTYH2	0.015468	0.307	0.013	12.601
HMGCS2	0.021933	0.310	0.023	0.543
AHNAK	0.024634	0.310	0.013	0.083
IL10RB	0.030052	0.310	0.024	0.256
CNNM2	0.031078	0.310	0.031	12.668
DGKB	0.031314	0.310	0.029	5.870
C1orf226	0.033464	0.310	0.031	7.390
MAP2K4	0.033824	0.310	0.035	0.341
FAM105A	0.034988	0.310	0.035	3.471
LOC144481	0.035639	0.310	0.039	0.414
DGAT2	0.042059	0.344	0.033	4.079
RCAN1	0.045748	0.353	0.061	0.271

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