

gene	pvalue	Hazard ratio
IFNB1	0.028	0.617(0.401-0.949)
PSMA2	0.002	0.736(0.607-0.892)
SEM1	0.031	0.761(0.594-0.975)
STAT4	0.045	0.773(0.601-0.995)
CASP2	0.024	0.775(0.621-0.967)
CD38	0.001	0.776(0.667-0.904)
PSMA4	0.036	0.785(0.627-0.984)
ISG20	0.002	0.800(0.693-0.923)
DNAJA1	0.040	0.812(0.666-0.991)
GCH1	0.026	0.829(0.703-0.978)
STAT1	0.013	0.849(0.746-0.966)
ZBP1	0.023	0.863(0.760-0.980)
GZMB	0.032	0.896(0.810-0.991)
TAP1	0.027	0.896(0.813-0.988)
UBB	0.023	0.904(0.829-0.986)
TIMP3	0.016	1.119(1.021-1.227)
UNC5B	0.024	1.132(1.016-1.261)
GADD45B	0.048	1.139(1.001-1.295)
AKT2	0.050	1.168(1.000-1.363)
PIK3CD	0.035	1.171(1.011-1.356)
IGF2R	0.021	1.186(1.026-1.371)
STAT5A	0.027	1.200(1.021-1.411)
CDKN1B	0.049	1.200(1.001-1.438)
NFKB1	0.042	1.209(1.007-1.452)
PIK3CG	0.048	1.212(1.002-1.466)
ERBB2	0.012	1.219(1.044-1.423)
BBC3	0.042	1.228(1.007-1.496)
PRKAR1B	0.024	1.228(1.028-1.468)
EMP1	<0.001	1.247(1.097-1.417)
GLUD1	0.038	1.252(1.012-1.548)
PSMB1	0.036	1.276(1.016-1.604)
PPP3CA	0.007	1.299(1.074-1.572)
CHP1	0.025	1.325(1.036-1.693)
PYGB	0.003	1.334(1.105-1.611)
HGF	0.005	1.338(1.092-1.640)
APC	0.007	1.376(1.092-1.734)
ELANE	0.029	2.305(1.090-4.875)
CAMK2A	0.033	2.396(1.074-5.346)
PSMA8	0.032	2.750(1.094-6.914)
TICAM2	0.022	3.809(1.216-11.936)

