

Table S2 Univariate Cox proportional hazards regression analyses of TRIM family genes in TCGA cohort

Gene	HR	95%CI	P value
TRIM16	1.27	1.06 – 1.52	0.009
TRIM2	0.81	0.70 – 0.94	0.005
TRIM22	0.81	0.69 – 0.95	0.010
TRIM27	0.68	0.47 – 0.97	0.034
TRIM29	1.17	1.07 – 1.29	0.001
TRIM38	0.58	0.41 – 0.80	0.001
TRIM39	0.58	0.39 – 0.87	0.007
TRIM58	1.41	1.03 – 1.93	0.030
TRIM59	1.48	1.09 – 2.00	0.012
TRIM6	1.43	1.18 – 1.73	<0.001
TRIM62	0.58	0.40 – 0.83	0.003
TRIM7	1.77	1.31 – 2.38	<0.001

HR: Hazard ratio; CI: Confidence interval