

Table S4. Hazard ratio for the critical- and neighbor-gene models.

Critical-gene model	Hazard Ratio	95% CI - lower	95% CI - upper
GSE2658	3.034518	1.955761	4.708297
GSE15695	1.875834	1.321967	2.661754
GSE9782	1.892779	1.375461	2.604663

Neighbor-gene model			
GSE2658	1.905190	1.158813	3.13230
GSE15695	1.657579	1.186769	2.315166
GSE9782	1.581617	1.114935	2.243639