

Table S5. Multivariate analysis comparing both the critical- and the neighbor-gene models with other prognostic gene models in GSE15695, GSE9782, GSE2658 (TT2).

Gene signature	Multivariate Cox regression p-value		
	GSE15695	GSE9782	GSE2658 (TT2)
Critical-gene model	0.037	0.039	7.9e-03
Neighbor-gene model	0.017	0.034	0.39
IFM 15-gene	0.68	0.140	0.36
UAMS 70-gene	0.2	0.024	2.1e-06
6-gene signature	9.3E-04	0.140	0.62