Table S1. Different K values and values of probably approximately correct (PAC)

K	Value of PAC
2	0.042750
3	0.460665
4	0.408903
5	0.286367
6	0.258136
7	0.216269
8	0.208763
9	0.194634
10	0.187596

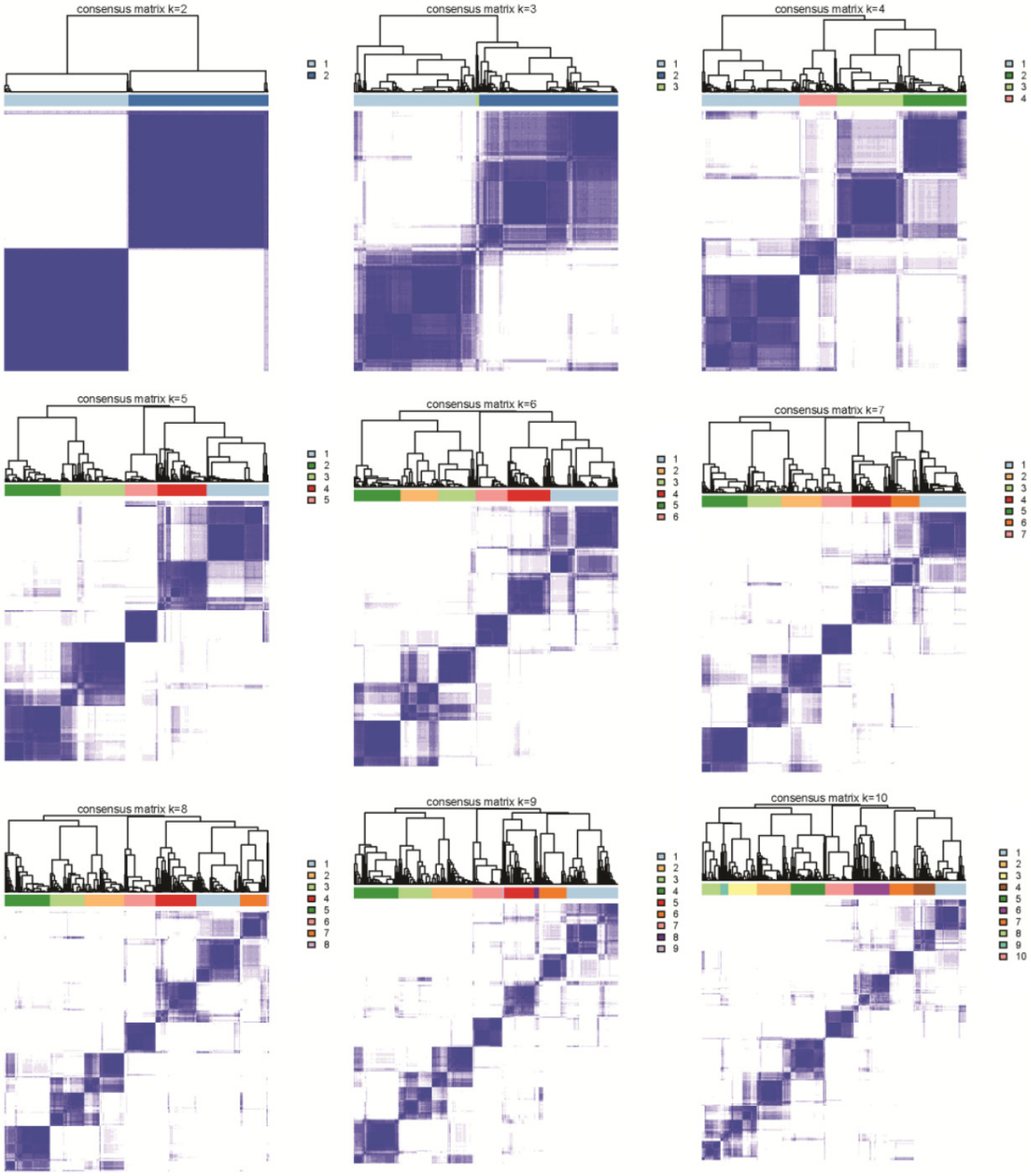

Figure S1. Consensus matrix for clustering.

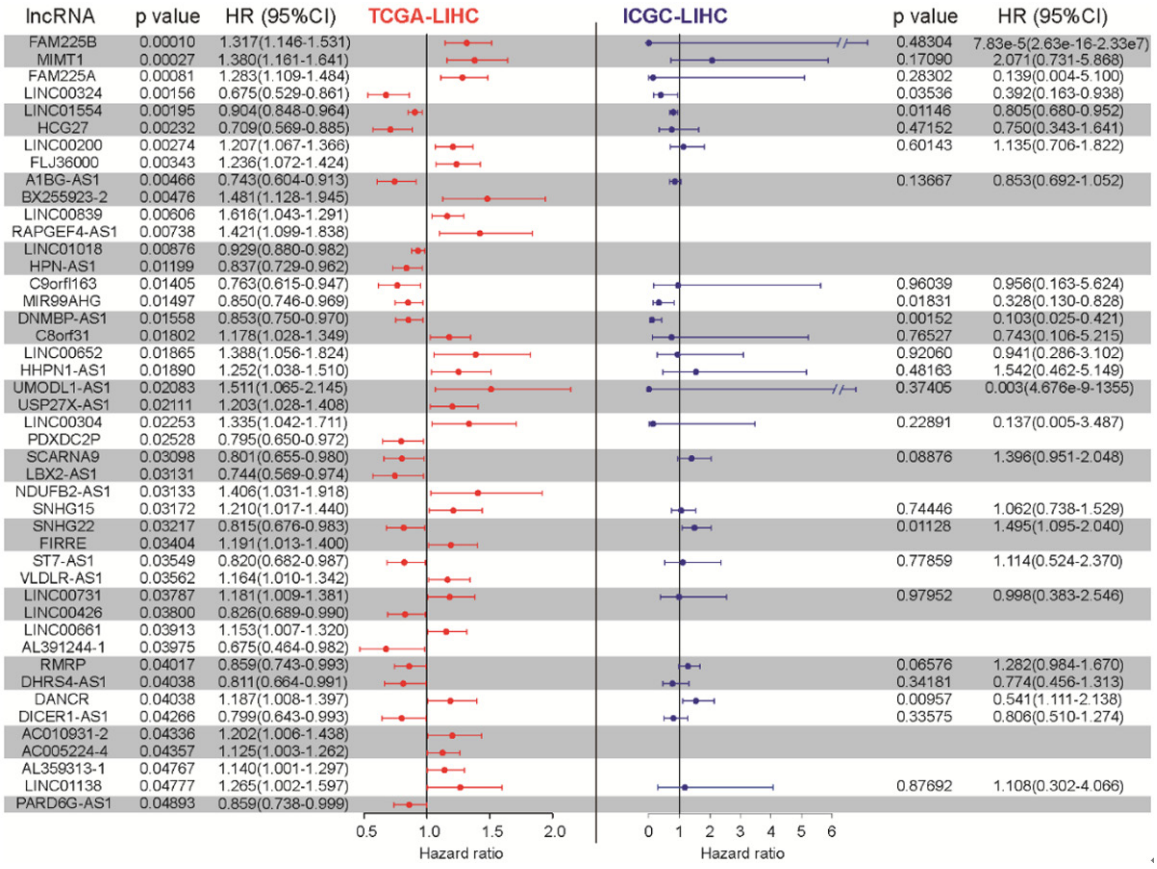

Figure S2. Survival analysis of individual lncRNA (adjusting for established risk factors) in the TCGA-LIHC (left and red) and ICGC-LIHC cohort (right and blue).