

Figure S1. Overview of our study.

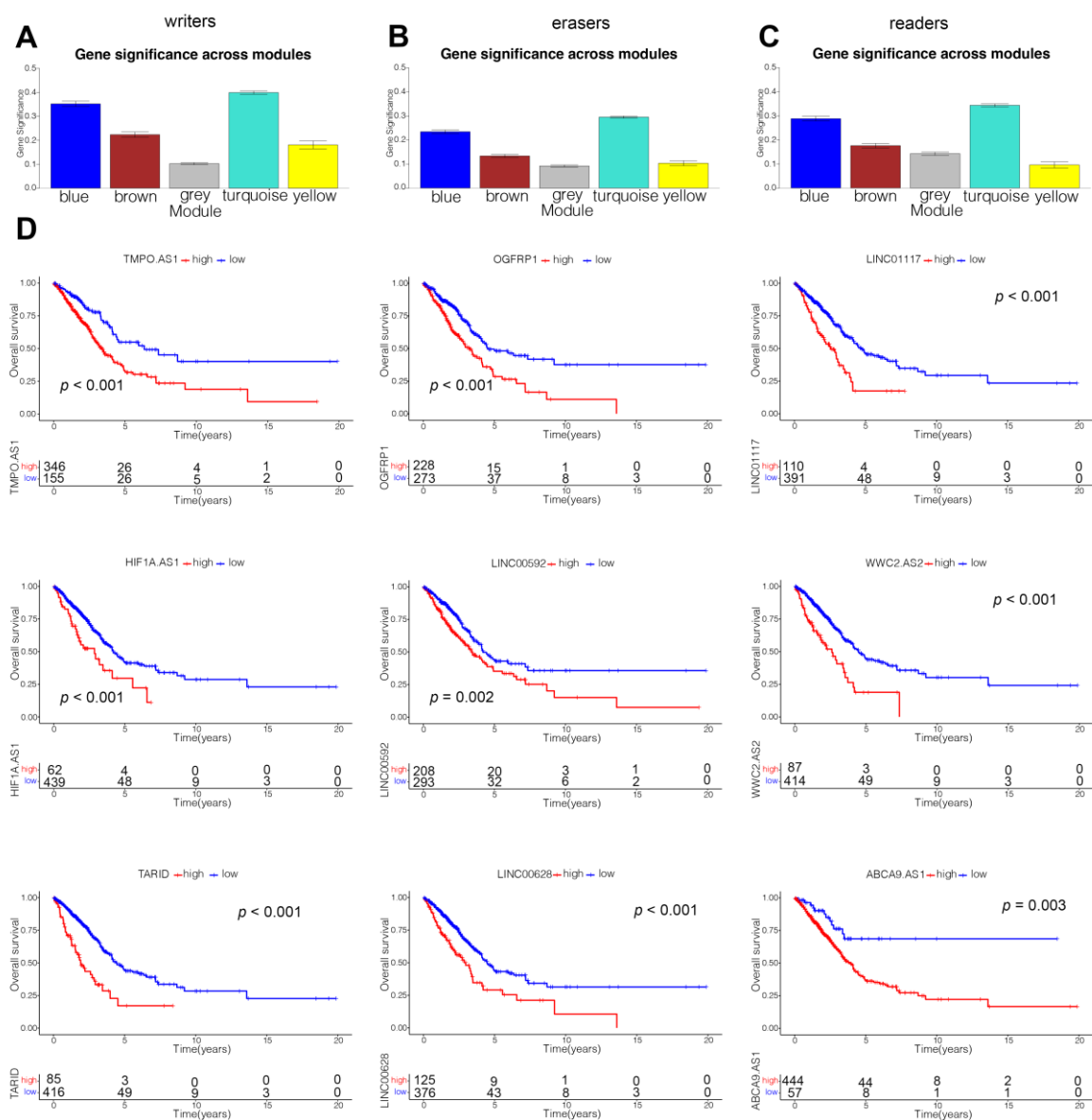


Figure S2. Nine lncRNAs were highly correlated with the survival of LUAD patients.

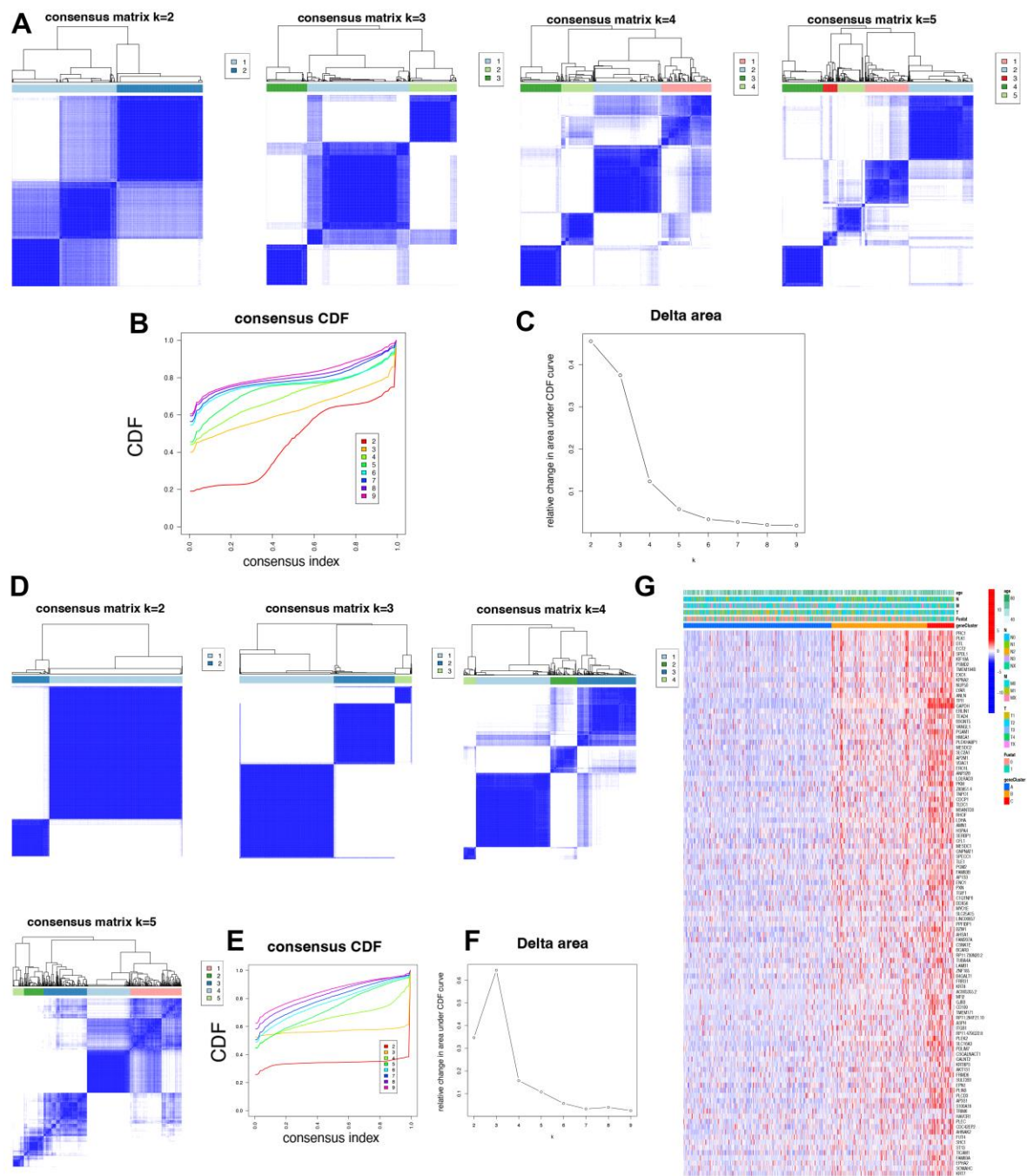


Figure S3. Unsupervised clustering of lncRNAs and DEGs; (A) Consensus matrices of TCGA-LUAD cohort for $k = 2$ to 5.

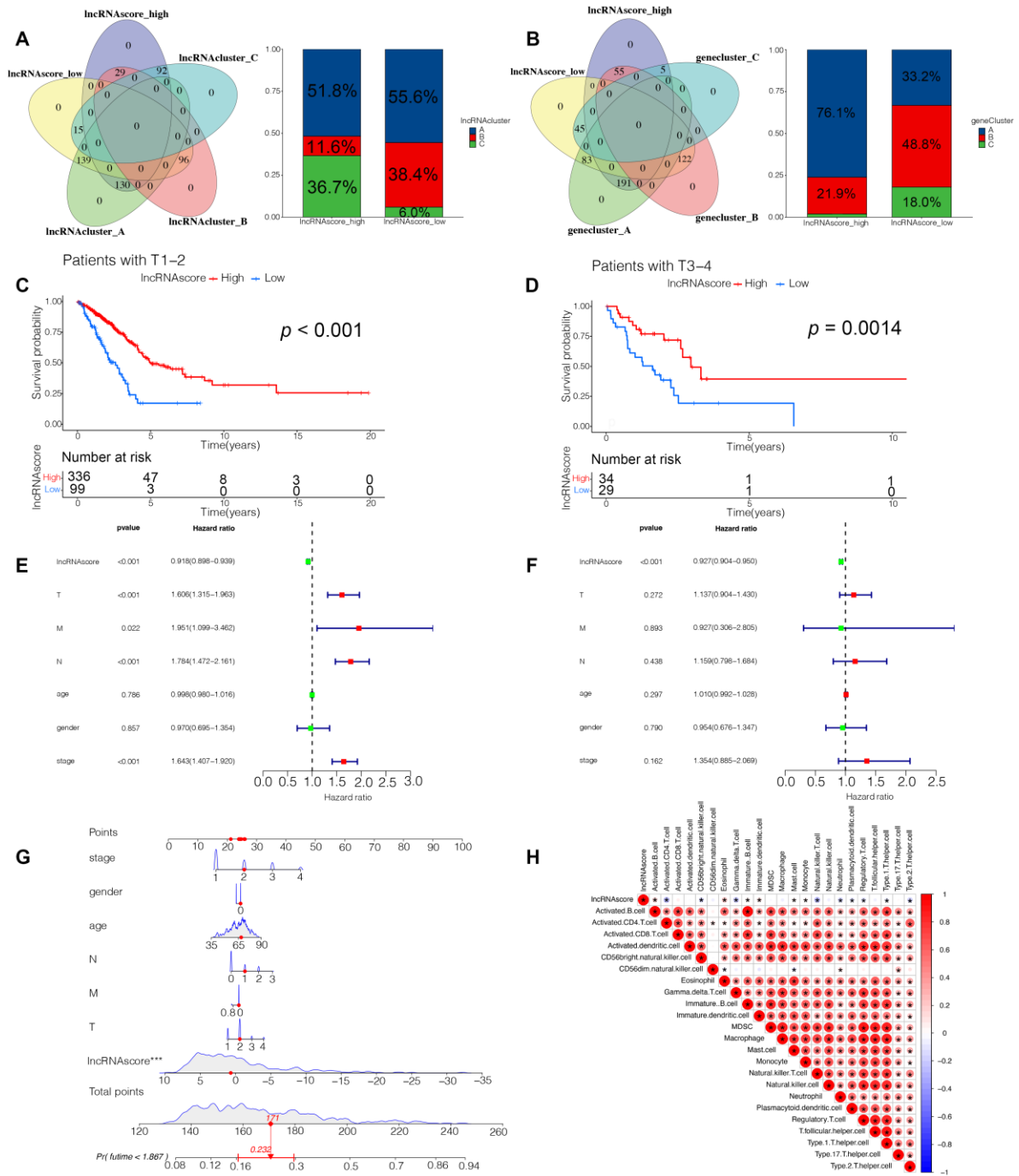


Figure S4. Prognostic value of IncRNA score and correlation between the clinicopathological features and IncRNA score.