

Seven LncRNA-mRNA based risk score predicts the survival of head and neck squamous cell carcinoma

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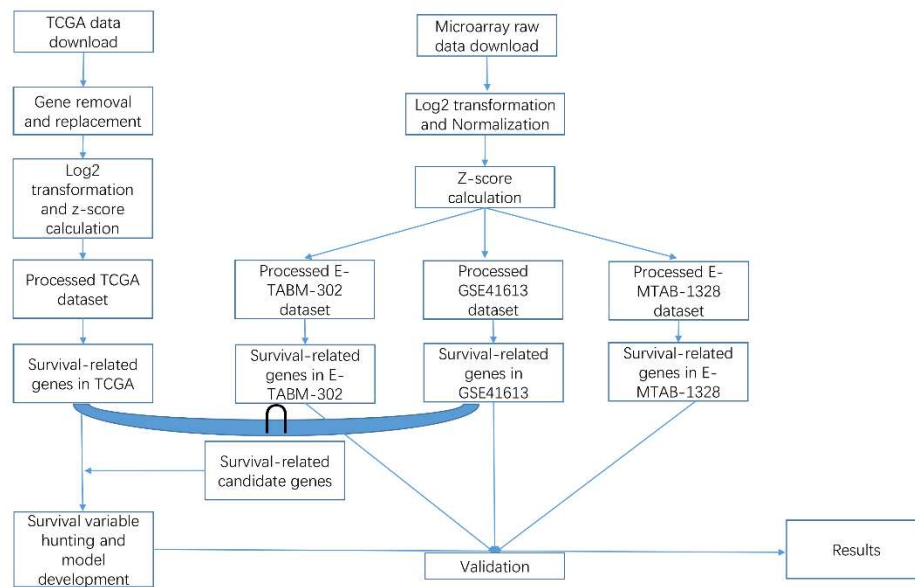


Figure S1. The workflow of this article.

80% resampling p values distribution

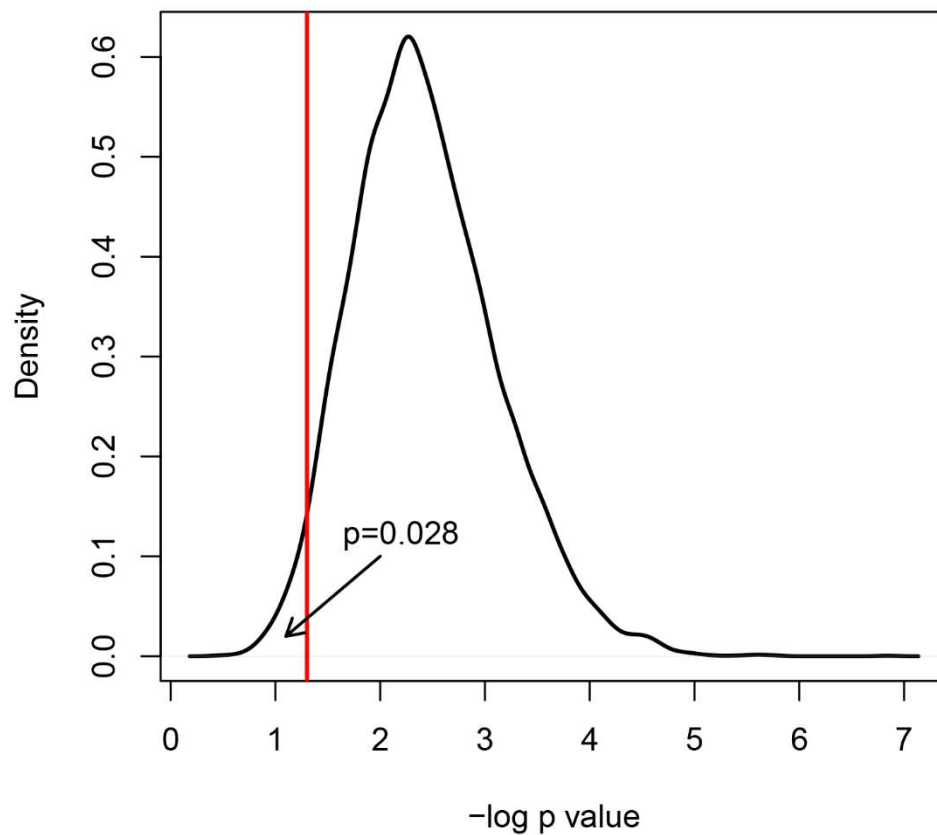


Figure S2. The distribution of p values of 10000 times 80% resampling. The x axis indicates the log 10 transformed p values, and y axis indicates the density of p value distribution. The red line is $\log_{10}0.05$.

Rank	TCGA (train)	E- MTAB302	GSE41613	E- TABM- 1328	number_significant
1	2.40E-07	0.52	0.037	0.031	3
2	0.01	0.37	0.032	0.22	1
3	2.50E-05	0.18	0.33	0.15	1
4	0.00077	0.12	0.00063	0.22	2
5	1.40E-02	0.18	0.004	0.46	2
6	1.20E-06	0.0013	0.16	0.19	2
7	2.30E-03	0.08	0.07	0.88	3
8	9.80E-06	0.12	0.016	0.053	3
9	3.40E-03	0.65	0.044	0.16	2
10	5.60E-04	0.08	0.0097	0.031	3

Table S1. The p values of ten times of randomly selected 7-gene combinations.