Survival curve (p=6.423e-03)

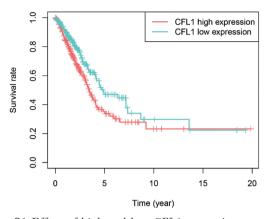


Figure S1 Effect of high and low CFL1 expression on patient survival.

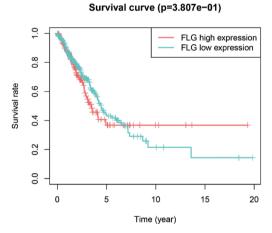


Figure S2 Effect of high and low *FLG* expression on patient survival.

Survival curve (p=3.007e-03)

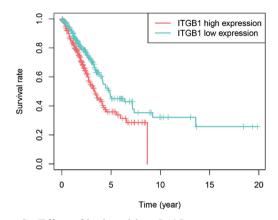


Figure S3 Effect of high and low *ITGB1* expression on patient survival.

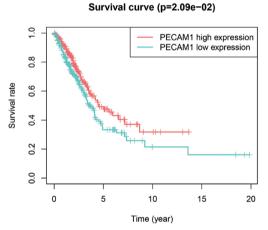


Figure S4 Effect of high and low *PECAM1* expression on patient survival.

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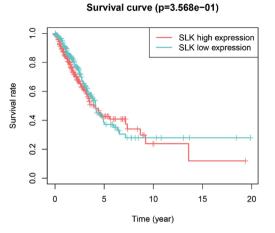


Figure S5 Effect of high and low *SLK* expression on patient survival.

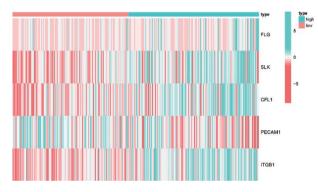


Figure S6 Heat map for high and low expression levels of prognosis-related membrane tension genes.

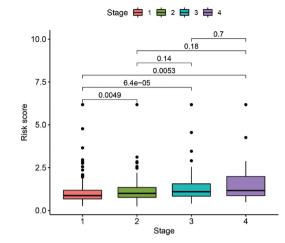
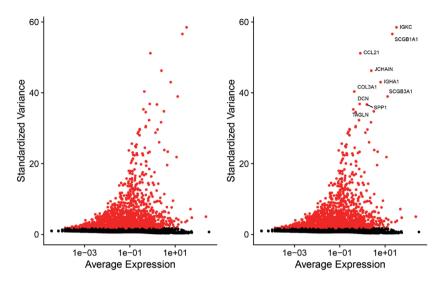


Figure S7 Correlation of risk score with TNM stage of LUAD.



Non-variable count: 23587 
Variable count: 2000

Figure S8 Top 10 highly variable genes.

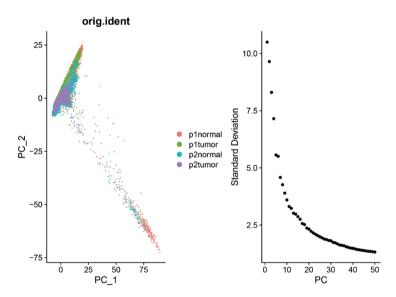


Figure S9 PCA downscaling.