



Research article

Identification of an endoplasmic reticulum stress-related gene signature to predict prognosis and potential drugs of uterine corpus endometrial cancer

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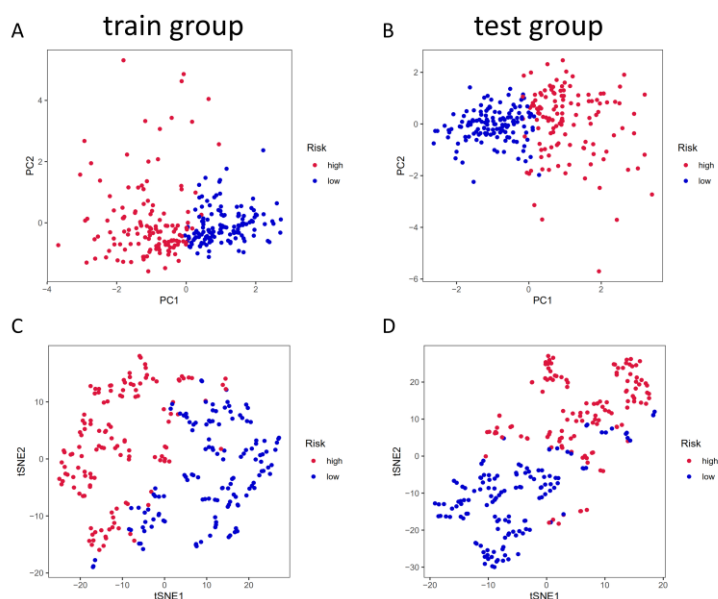


Figure S1. The principal component analysis (PCA) (A and B) and t-distributed stochastic neighbor embedding (t-SNE) (C and D) plots of the train and test groups.

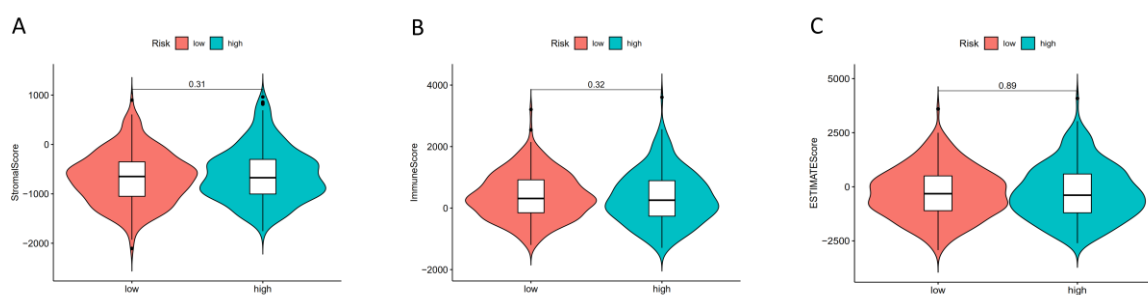


Figure S2. Comparison of tumor microenvironment score between high and low risk groups in UCEC, including (A) stromal score, (B) immune score, and (C) ESTIMATE score.

