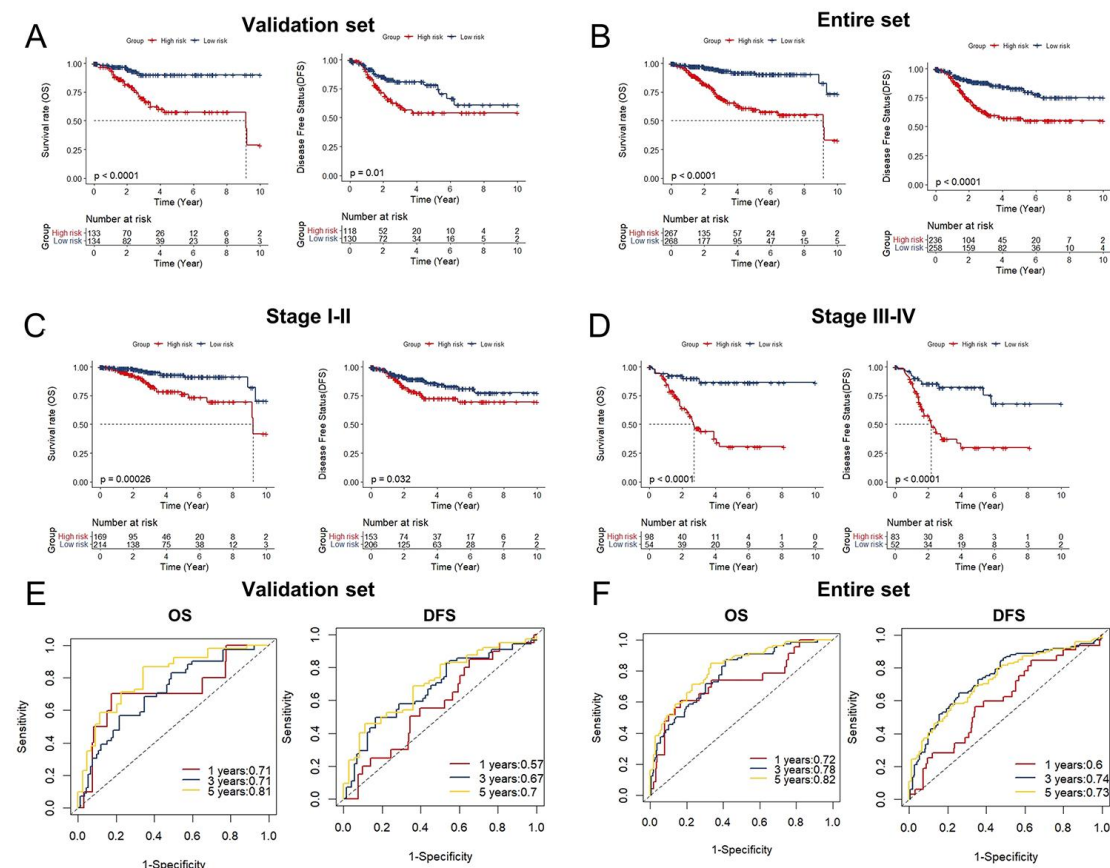


SUPPLEMENTAL FIGURES



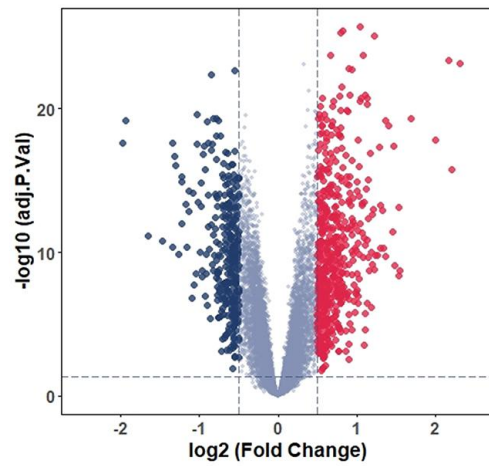
Supplement 1. Survival analyses of the low- and high-risk groups patients.

A, B. Kaplan–Meier survival analyses of the low- and high-risk groups patients in the validation set (A) and the entire set (B).

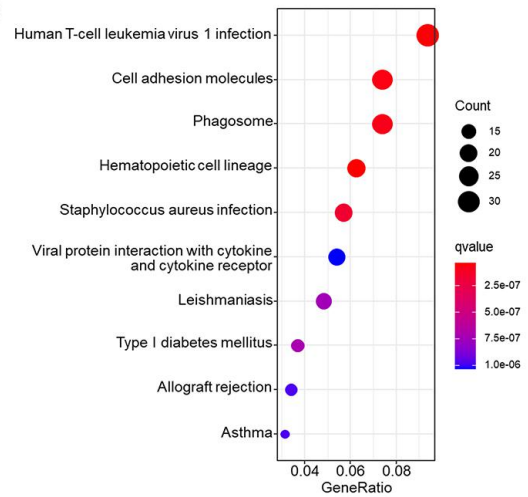
C, D. Kaplan–Meier survival analyses of the low- and high-risk groups patients at stage I–II (C) and stage III–IV (D).

E, F. ROC curve analyses according to the 1, 3, 5-year survival of the area under the AUC value in the validation set(E) and the entire set(F).

A



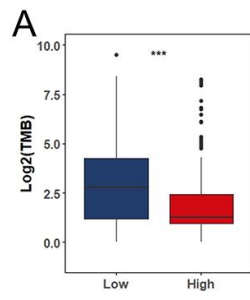
B



Supplement 2. Comprehensive analyses of enriched pathways in different risk groups.

A. Volcano plot representing 504 risk-related DEGs between low- and high-risk groups.

B. KEGG analysis of 504 risk-related DEGs. The size of the balls indicating the number of genes and color indicating q value.



Supplement 3. Comparison of genomic alterations in different risk groups

A. The tumor mutation burden in the low- and high-risk groups.