

Figure S1 Exploring the genetic variations in the high- and low-CRS subgroups. (A,B) The overall mutation status with the most mutated genes in the CRS subgroups. (C) Comparison of the TMB between the two risk-score subgroups. (D) Correlation between the CRS and TMB. (E,F) Kaplan-Meier plots showing the prognostic effect of TMB and the combination of TMB and the CRS. CRS, cuproptosis-related score; TMB, tumor mutational burden.

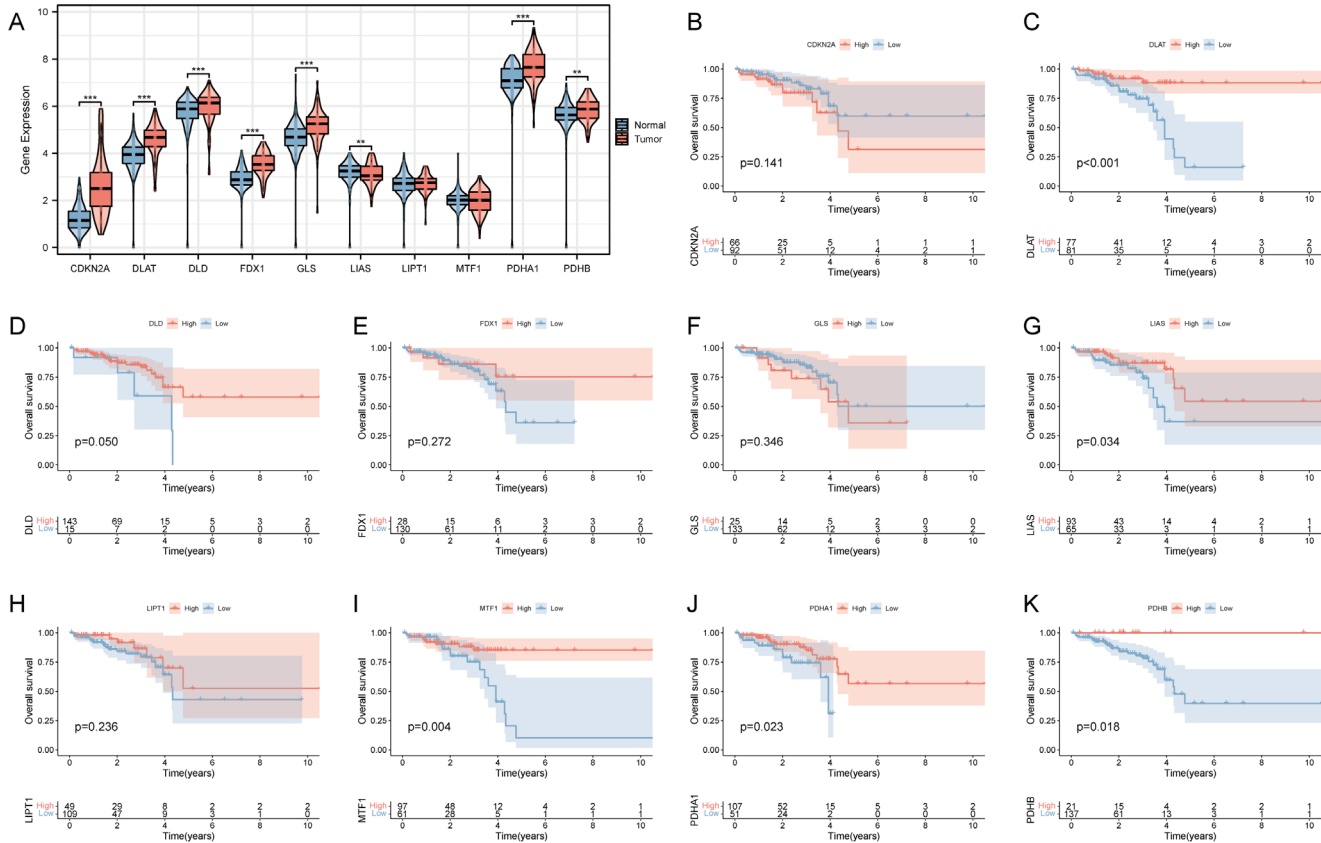


Figure S2 The prognostic value of the 10 CRGs. (A) Expression differences in the 10 CRGs between the normal and tumor tissues. (B-K) Kaplan-Meier plots showing the prognostic value of the 10 CRGs. **, $P \leq 0.01$; ***, $P \leq 0.001$. CRGs, cuproptosis-related genes.