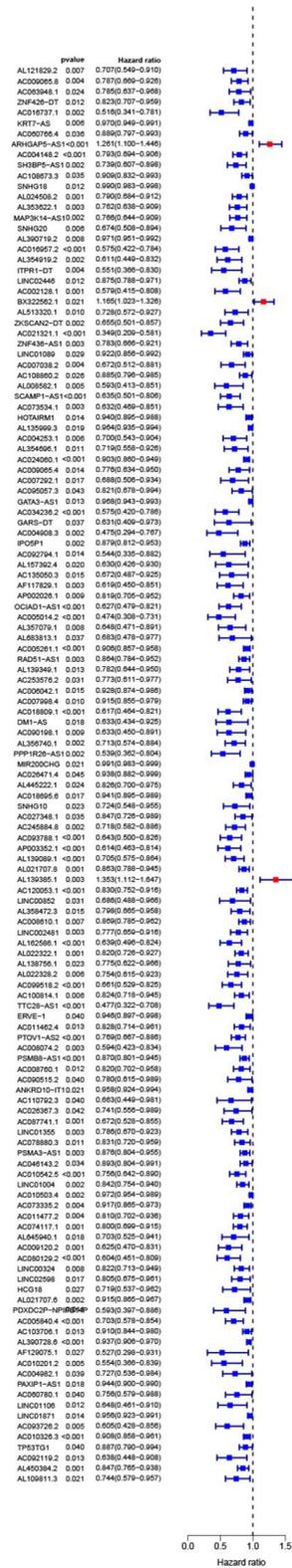
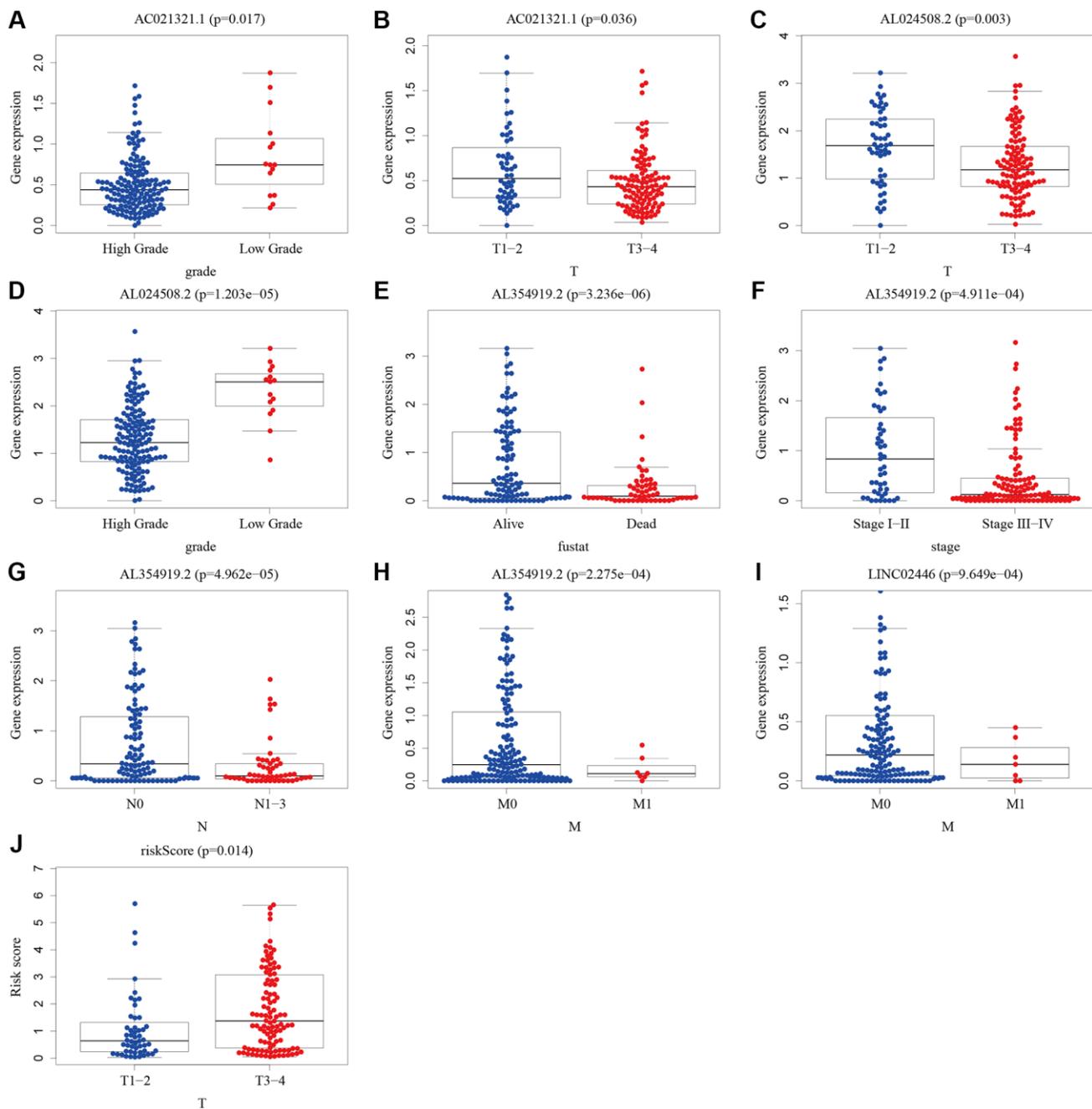


SUPPLEMENTARY FIGURES



Supplementary Figure 1. A total of 135 lncRNAs analyzed by Univariate Cox regression analysis.



Supplementary Figure 2. The correlations between risk scores/8 cuproptosis-related lncRNAs and clinicopathological variables. (A, B) Correlation between AC021321.1 expression level and grade and T stage. (C, D) Correlation between AL024508.2 expression level and grade and T stage. (E–H) Correlation between AL354919.2 expression level and fustat, stage N stage and M stage. (I) Correlation between LINC02446 and M stage. (J) Correlation between risk scores expression level and T stage. Abbreviations: lncRNAs: long noncoding RNAs; T: tumor; N: lymph node; M: metastasis.