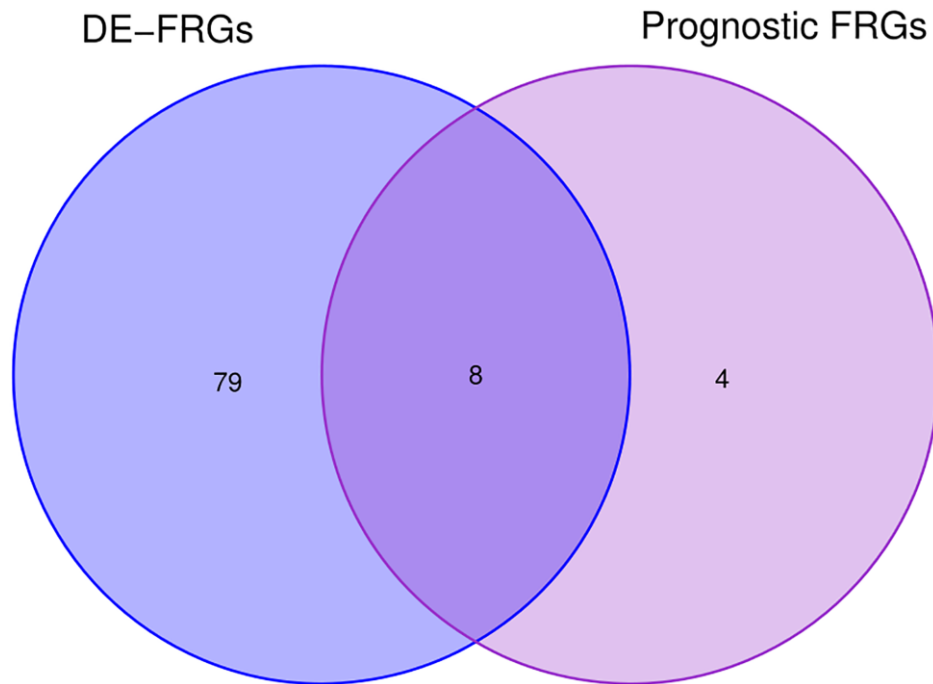
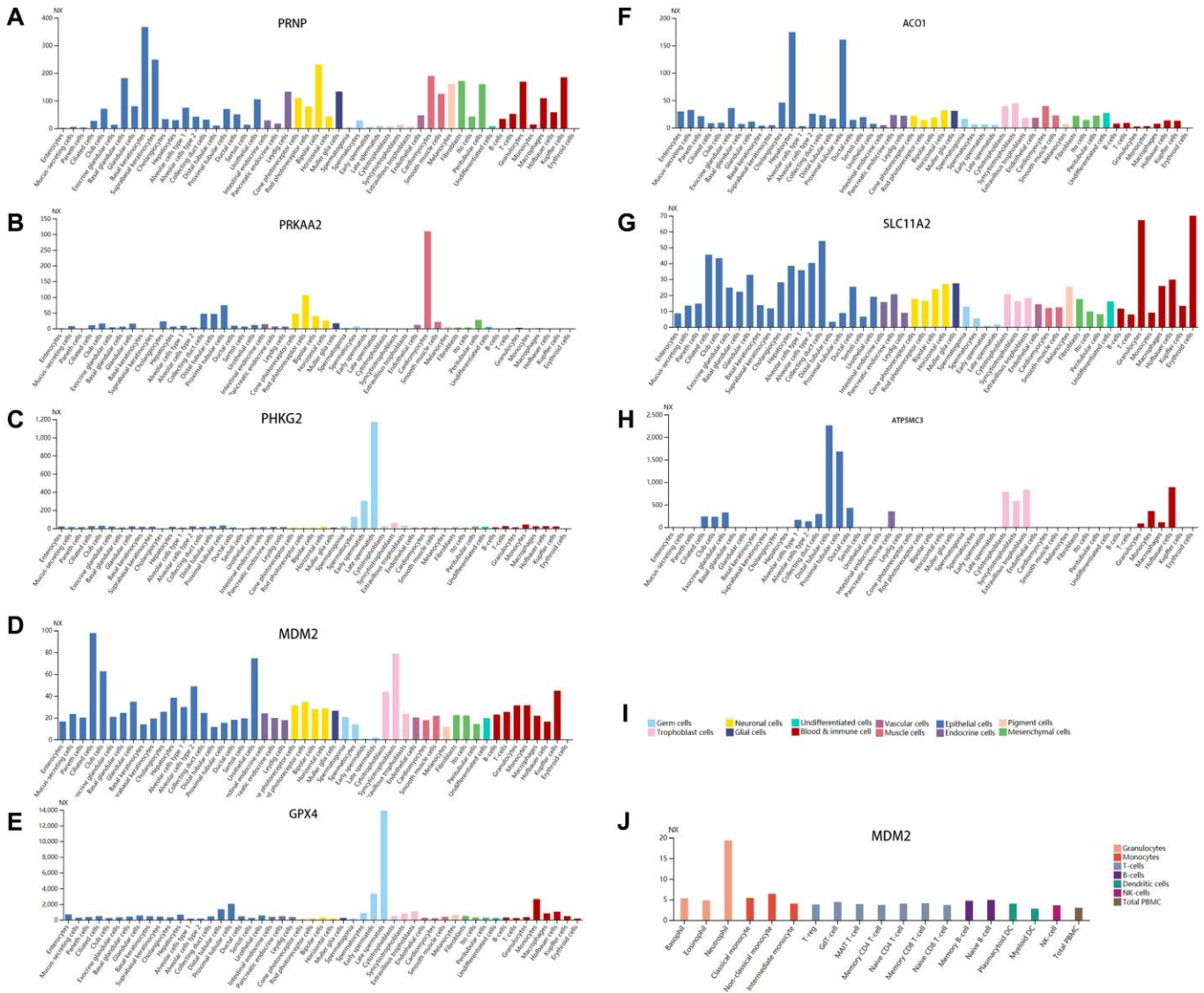


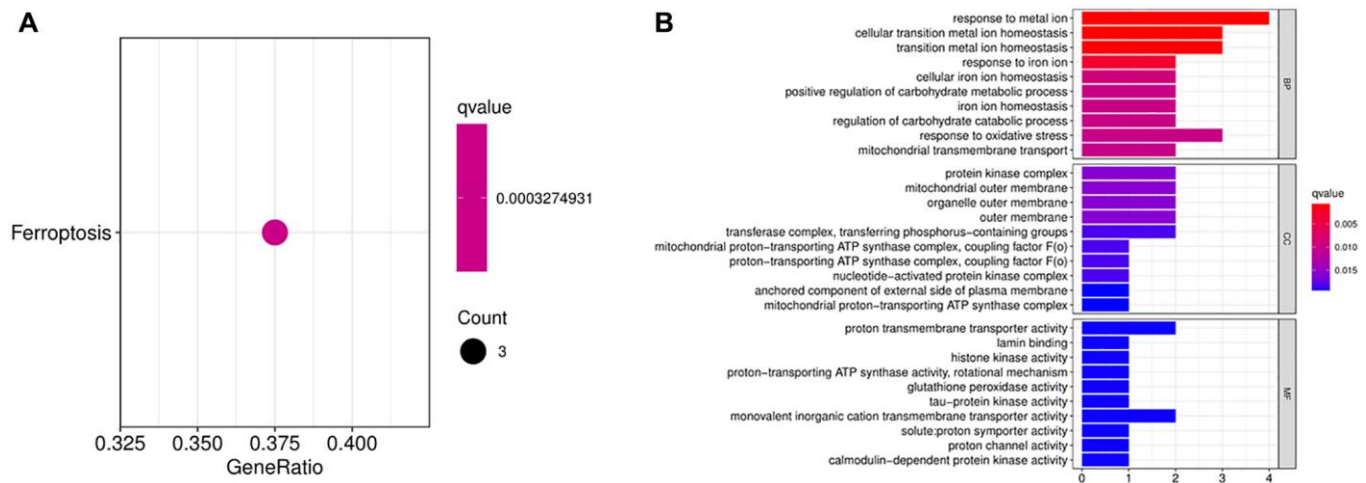
SUPPLEMENTARY FIGURES



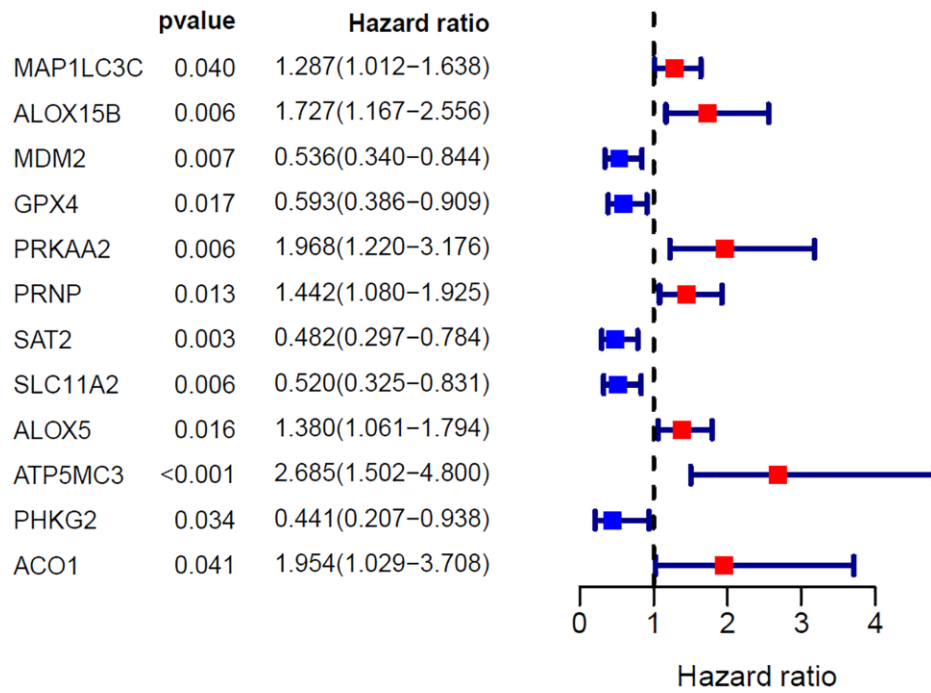
Supplementary Figure 1. Venn diagram of 87 differentially expressed ferroptosis-related genes (DE-FRGs) between normal samples and EC patient samples and 12 prognostic FRGs.



Supplementary Figure 2. (A–I) RNA single-cell analysis of 8 FRGs in normal human tissue cells. (J) MDM2 RNA single-cell analysis in blood cells.



Supplementary Figure 3. (A) KEGG analysis of the eight ferroptosis-related genes (FRGs). (B) GO analysis of the eight FRGs.



Supplementary Figure 4. Results of the univariate Cox analysis of the twelve prognostic ferroptosis-related genes in EC patients.