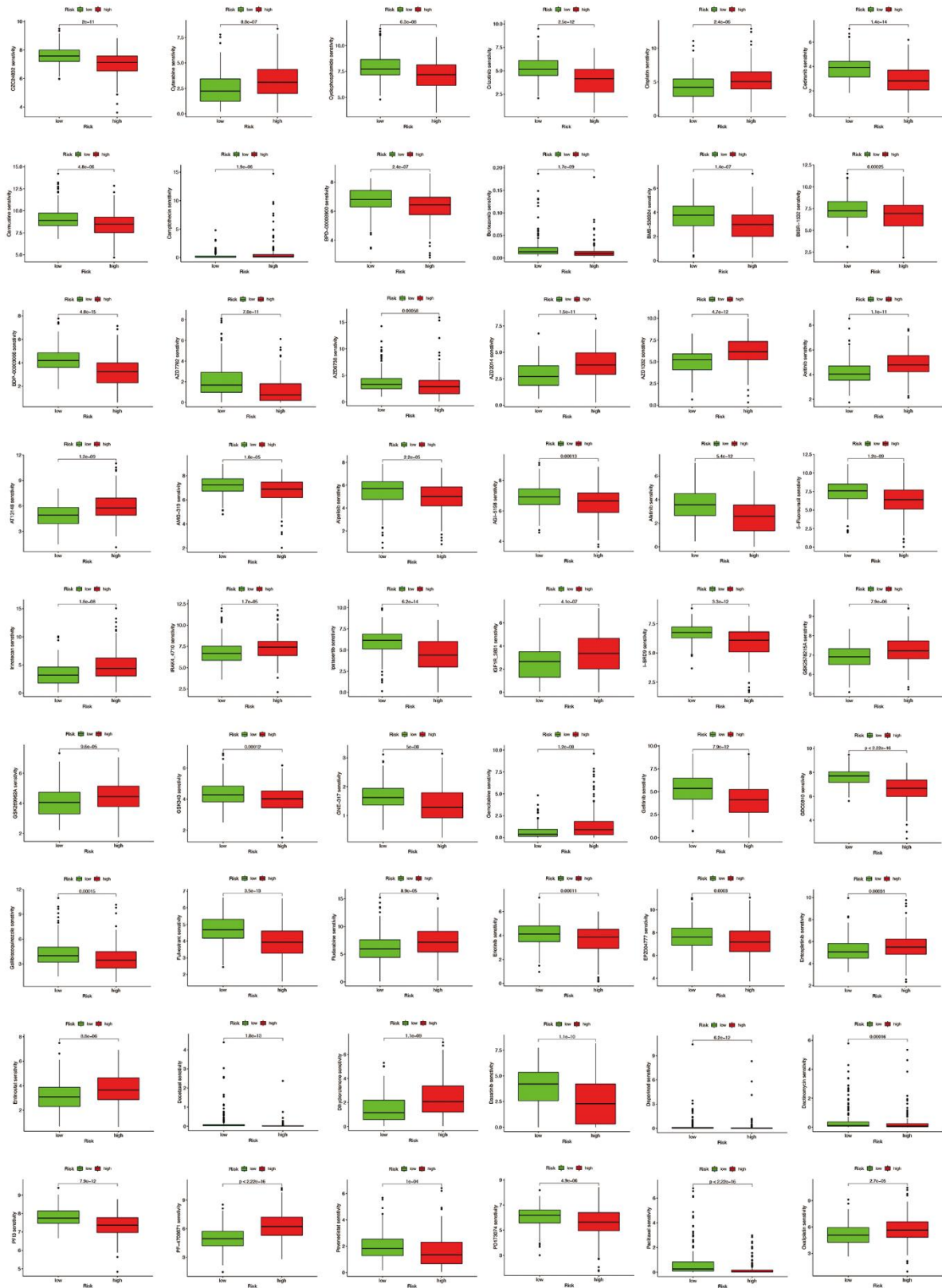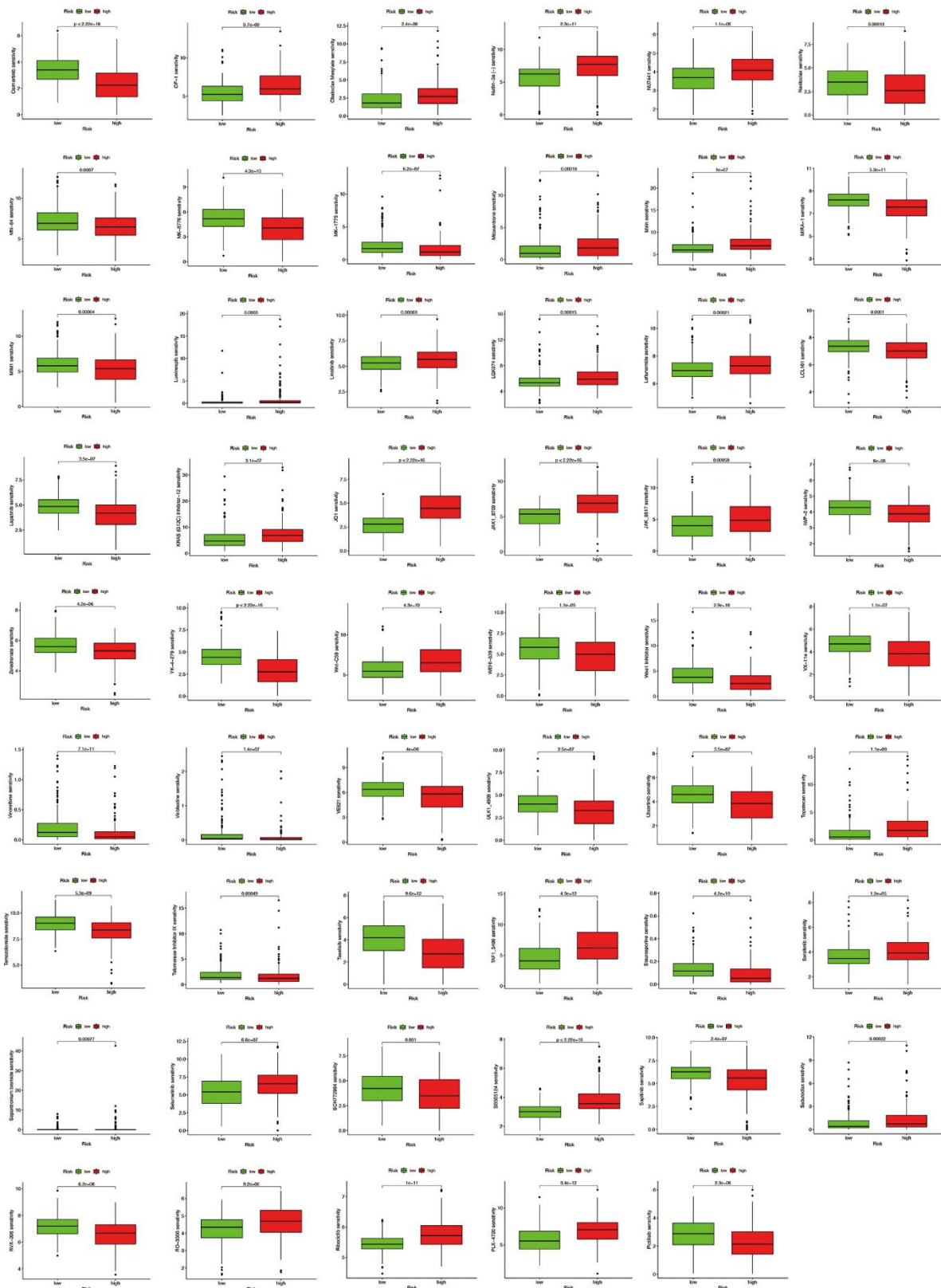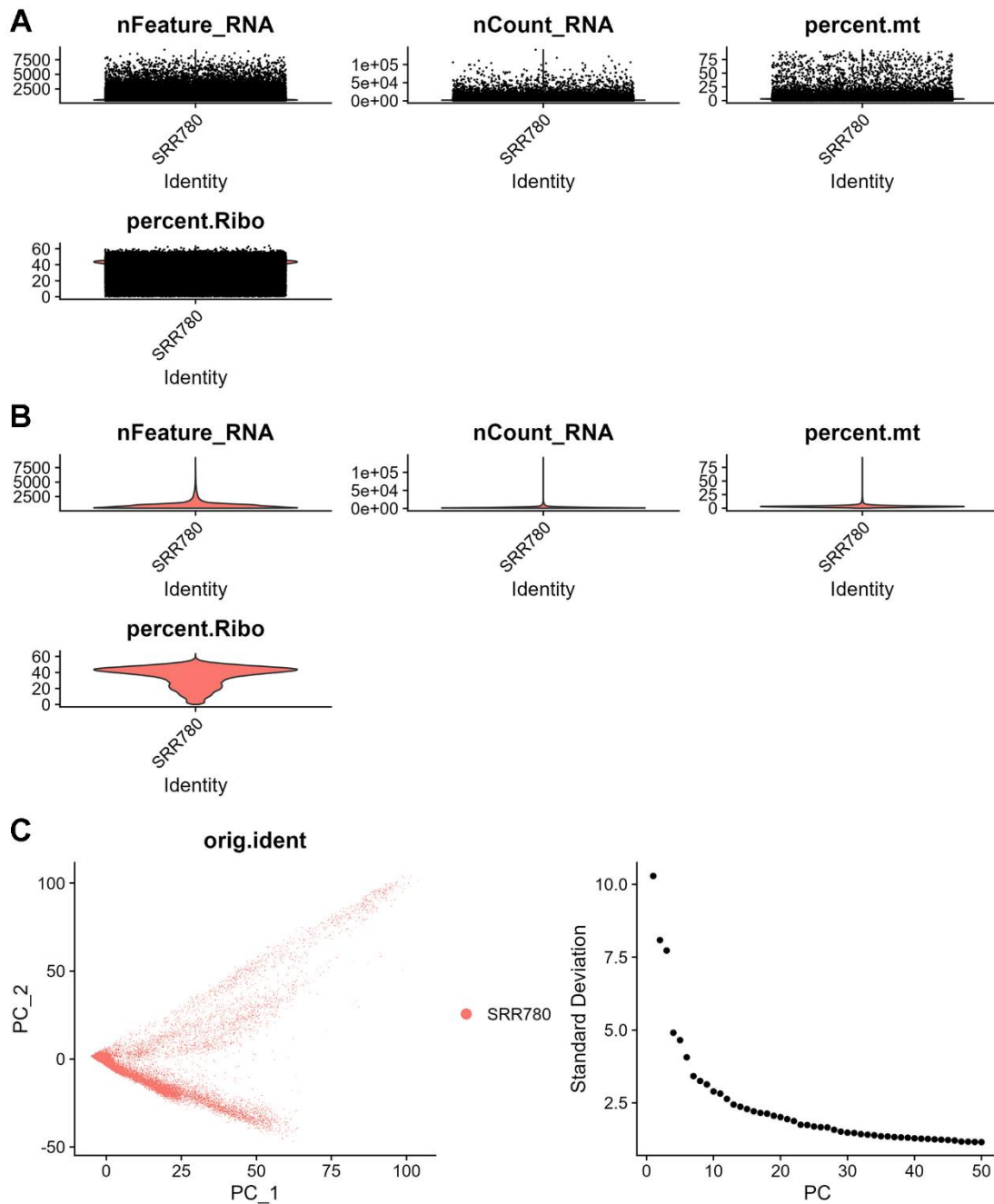# SUPPLEMENTARY FIGURES

Supplementary Figure 1. Comparison of the sensitivity of various chemotherapeutic agents in distinct risk groups of the TCGA-LIHC cohort.

Supplementary Figure 2. Single-cell RNA-seq profiling in hepatocellular carcinoma. (**A**, **B**) The Seurat package is used to filter data from individual cells. (**C**) Utilizing dimensionality reduction and the Seurat package, single-cell data can be clustered.