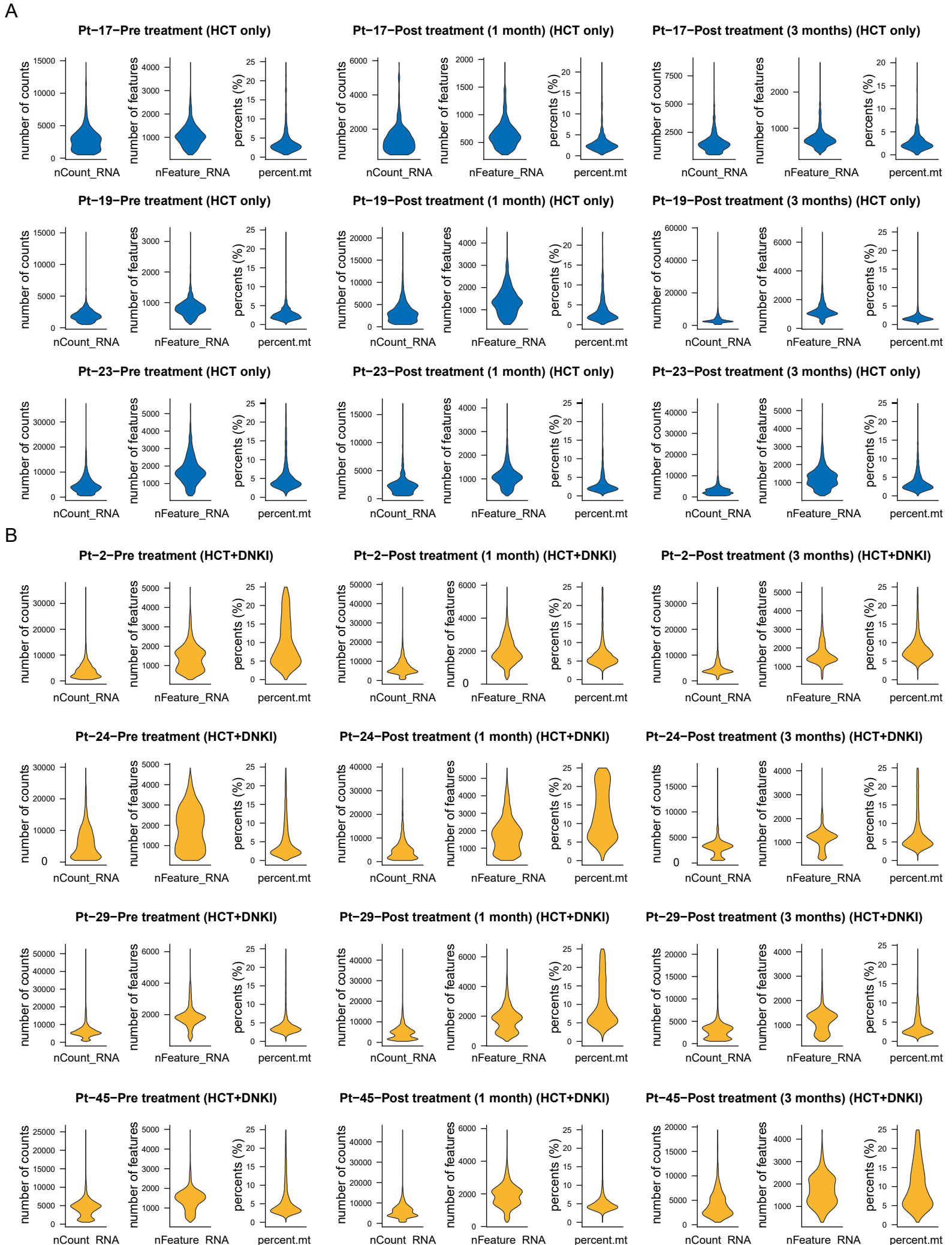




Supplementary Figure S1. QC metrics of the PBMC scRNA-seq data for the study patients.

(A, B) The violin plots showing the quality of cell data used in the present study. The number of genes per cell was over 200 but less than 7,000. In addition, mitochondrial genes was less than 15%. (A) Three patients treated with HCT only.

(B) Four patients treated with HCT and DNKI.