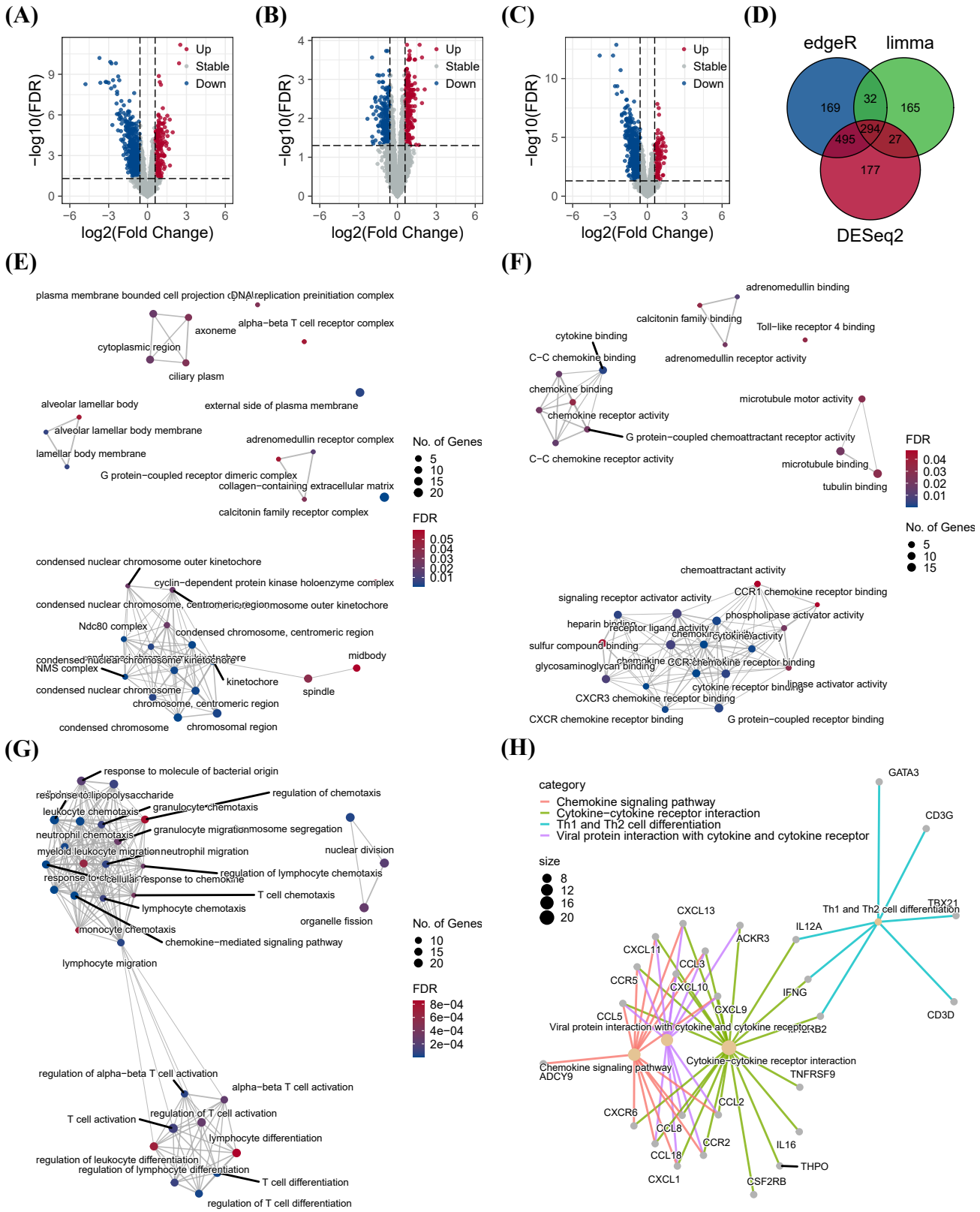

Supplementary Figure 2. Differential expression gene (DEG) analysis between patients who achieved pCR and those who did not by edgeR (A), DESeq2 (B), and limma (C). (D) overlapping DEGs. Over-representation analysis of DEGs for cellular component (E), molecular function (F), biological process (G), and KEGG pathways (H).