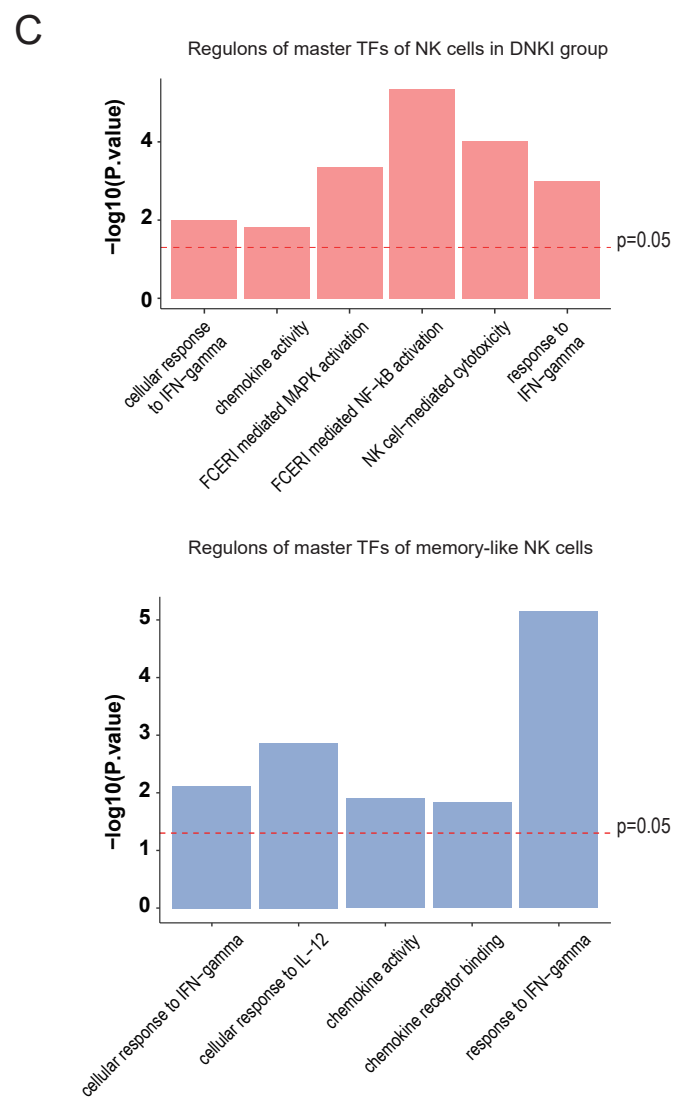
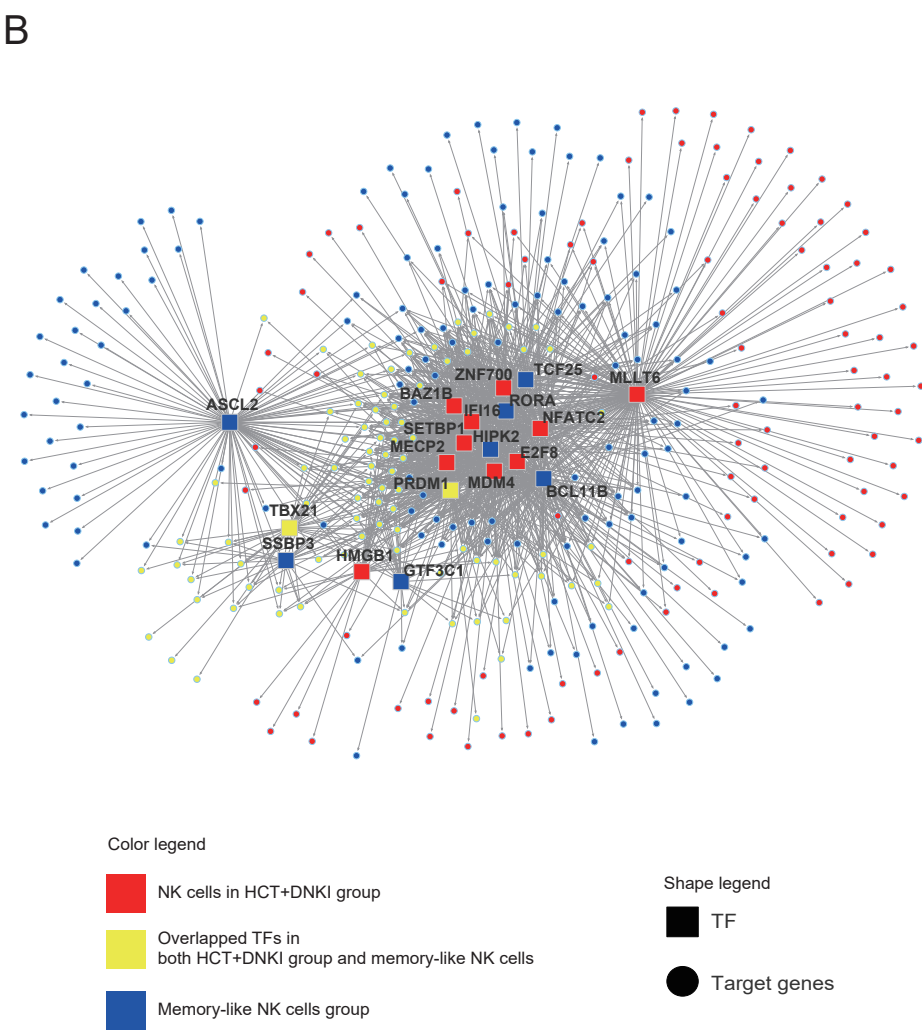
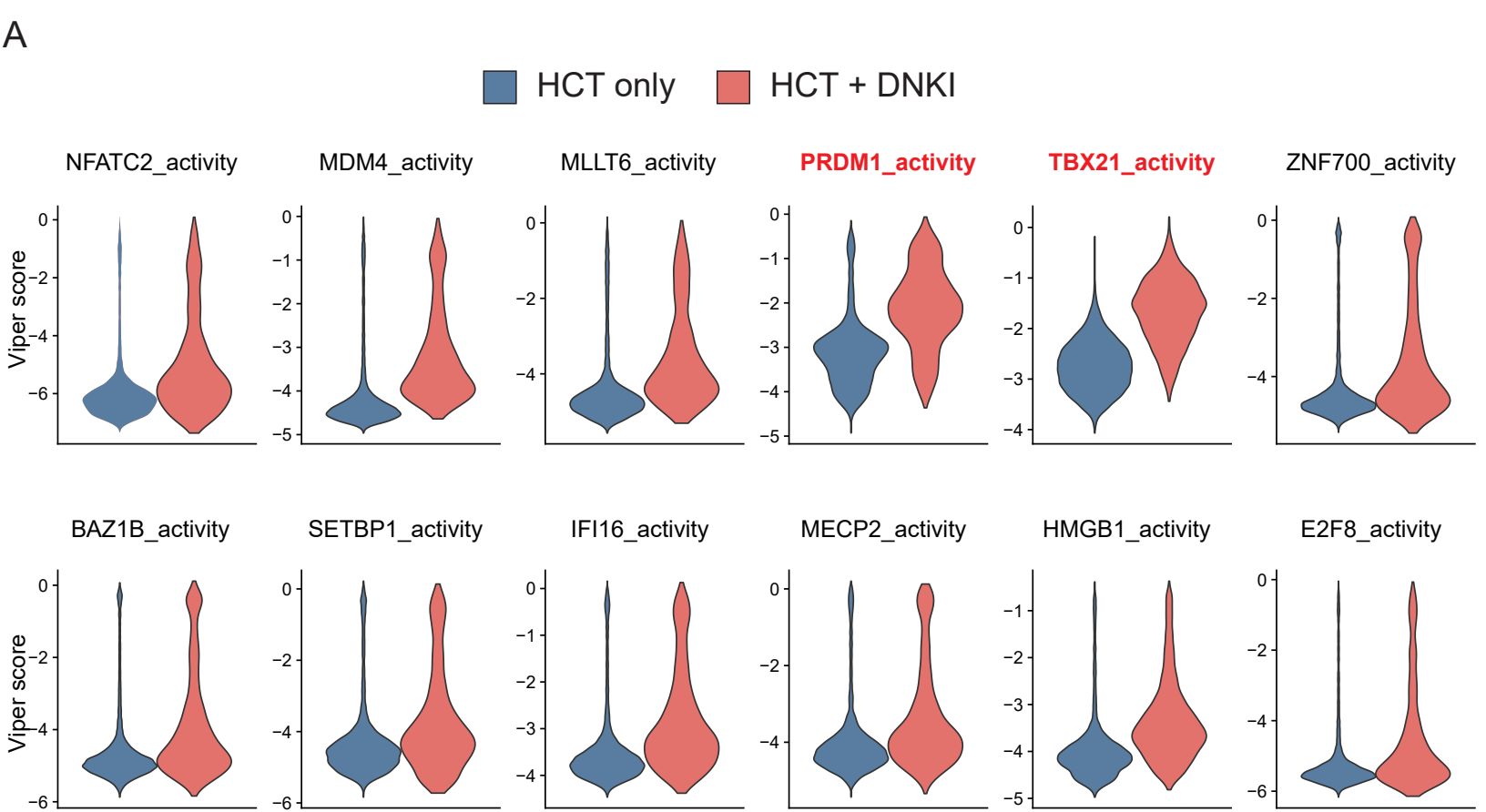




Supplementary Figure S6. PRDM1 and TBX21 participated in the differentiation of memory-like NK cells.

(A) The violin plots showing significantly different viper scores representing transcription factor (TF) activities according to the treatment groups (HCT only vs HCT plus DNKI). Among these 12 TFs, PRDM1 and TBX21 showed the most significantly different TF activities.

(B) A visual representation of gene regulatory network (GRN) of master TF candidates with statistical significance of $P < 0.01$ in VIPER analysis. The 17 TFs (square) and their estimated target genes (circle) were depicted. The regulatory links between the regulators and their inferred targets were represented in the periphery.

(C) The bar plots depicting the biological phenotypes of regulons regulated by the master TFs of NK cells in DNKI group or memory-like NK cells.