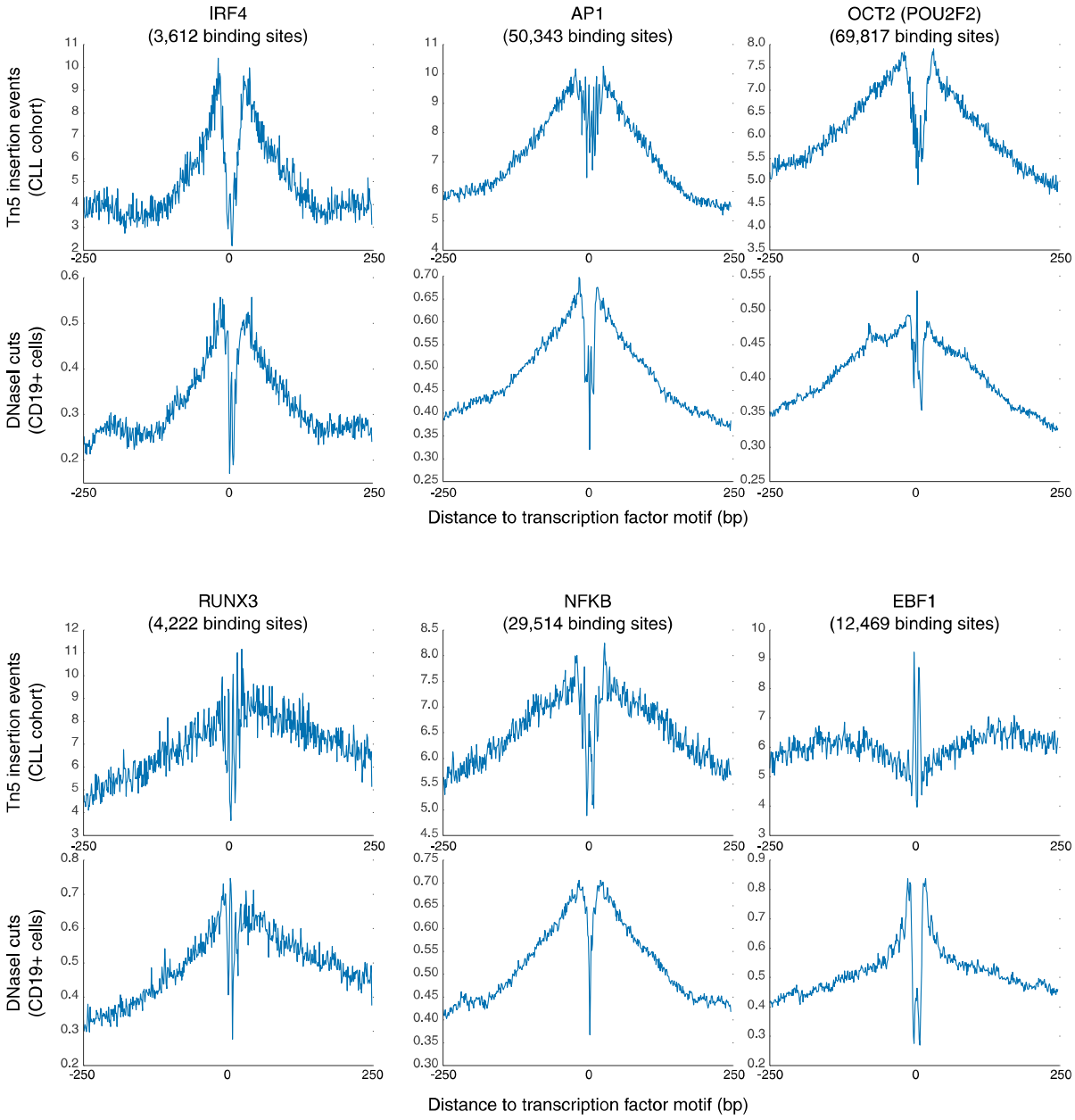

Supplementary Figure 16

Transcription factor footprints for ATAC-seq and DNase-seq are similar.

Footprinting diagrams showing the frequency of Tn5 transposase insertion events (for ATAC-seq) and DNase I cutting sites (for DNase-seq, based on data for CD19+ B cells from the Roadmap Epigenomics project) across a 500 basepair window around DNA binding motifs of transcription factors involved in B cell development.