

Figure S8

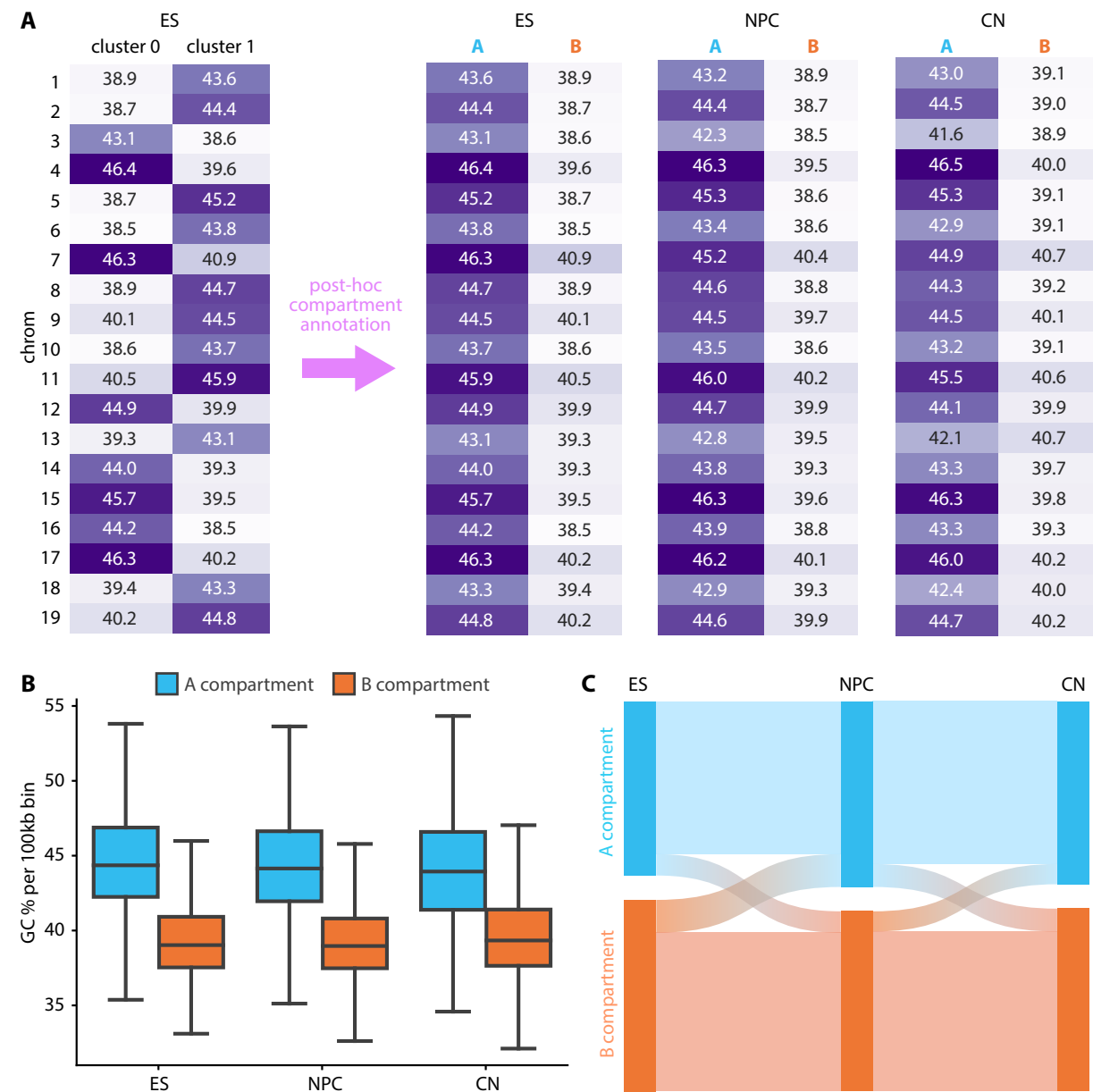


Figure S8. Post-hoc annotation of TGIF-DC clusters into A and B compartments in mouse neural differentiation data. **(A)** Within each chromosome, the mean GC percentage is measured for regions in each TGIF-DC cluster for ES. Cluster with higher mean GC content is assigned to compartment A; the other to compartment B across all timepoints. **(B)** Genome-wide distribution of GC percentage in each 100kb bin by compartment assignment and timepoint. **(C)** Dynamic compartment assignment patterns for all genomic regions from ES to NPC to CN state.