

Figure S2

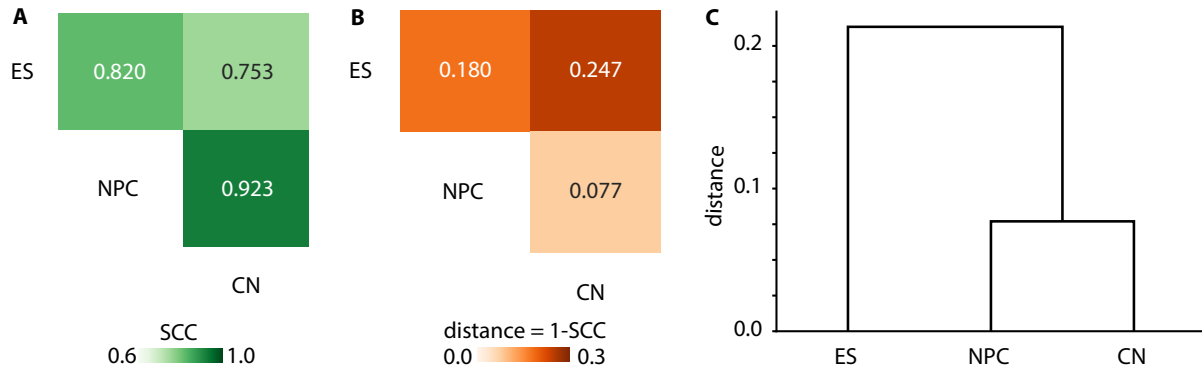


Figure S2. Inferring a tree structure between input conditions based on the similarity of the input matrices. Here we use mouse chr19 intra-chromosomal matrices (25kb resolution) from neural differentiation data with 3 timepoints (ES, NPC, CN) as an example. **(A)** Similarity between input matrices for each pair of timepoints, measured by stratum-adjusted correlation coefficient (SCC). **(B)** Similarity converted to distance = $1 - \text{SCC}$. **(C)** Dendrogram of hierarchy constructed using the distance and average linkage.