

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: **Gene Ontology (GO) enrichment results.** The tables show Gene Ontology (GO) enrichment results of ATAC-seq peak closest genes (Figure 2d-f), VME or EPC activated bivalent genes (Figure 3e, h), stable bivalent genes (supplementary Figure 3g), and feature genes in each lineage defined by scRNA-seq (supplementary Figure 4b).