Description of Additional Supplementary Files

File name: Supplementary Data 1

Description: Paired RNA-seq and ATAC-seq data used for construction of hReg-CNCC.

File name: Supplementary Data 2

Description: The upstream TFs and REs regulating ALX1.

File name: Supplementary Data 3

Description: List of human paired gene expression and chromatin accessibility data used in

regression of interaction strength.

File name: Supplementary Data 4

Description: 27 tissues used for control of fold change enrichment of face GWAS.

File name: Supplementary Data 5

Description: Source data underlying the graphs and charts presented in the main figures.