

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Sequencing quality metrics for scBS-seq and scRNA-seq libraries.

File Name: Supplementary Data 2

Description: Summary statistics obtained for the cell-specific association analysis correlating methylation with expression, accessibility with expression or methylation with accessibility at individual loci.

File Name: Supplementary Data 3

Description: Genomic annotations used in this study.