

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Summary statistics of all sequencing datasets used in this study.

File Name: Supplementary Data 2

Description: Normalized chromatin accessibility and gene expression level of EGA genes at each stage.

File Name: Supplementary Data 3

Description: Enrichment of TF motifs at each stage.

File Name: Supplementary Data 4

Description: Normalized chromatin accessibility and RNA expression level of retrotransposons at each stage.