

Description of Additional Supplementary Files

Title: Supplementary Data 1

Description: Gene ontology (GO) analysis of the differentially methylated genes

Title: Supplementary Data 2

Description: List of the common transcription factor binding sites in the promoter regions of ESR1, GATA3, and FOXA1 genes

Title: Supplementary Data 3

Description: Motif analysis of the differentially methylated DNA regions