The landscape of DNA methylation-mediated regulation of long non-coding RNAs in breast cancer

SUPPLEMENTARY MATERIALS

Supplementary Table 1: The result of cis-regulatory function analysis of IncRNAs. See Supplementary_Table_1

Supplementary Table 2: The result of function enrichment analysis and hallmarks in four ceRNA networks. See Supplementary_Table_2

Supplementary Table 3: The result of miRNA enrichment analysis in four ceRNA networks. The yellow color stands for the breast cancer. See Supplementary_Table_3