



Supplemental Figure 8. Gene expression changes is correlated with both t-DMRs and c-DMRs at CGIs and CGI shores. (A) t-DMRs between normal liver and lung (B) liver c-DMRs (C) lung c-DMRs. Y-axis indicates log₂ ratio of gene expression (measured by FPKM) between tissues. DMRs that are located within 2 kb from TSSs of genes were used for plotting. Block circles represent CGI shore-associated DMRs, while red circles represent CGI-associated DMRs.