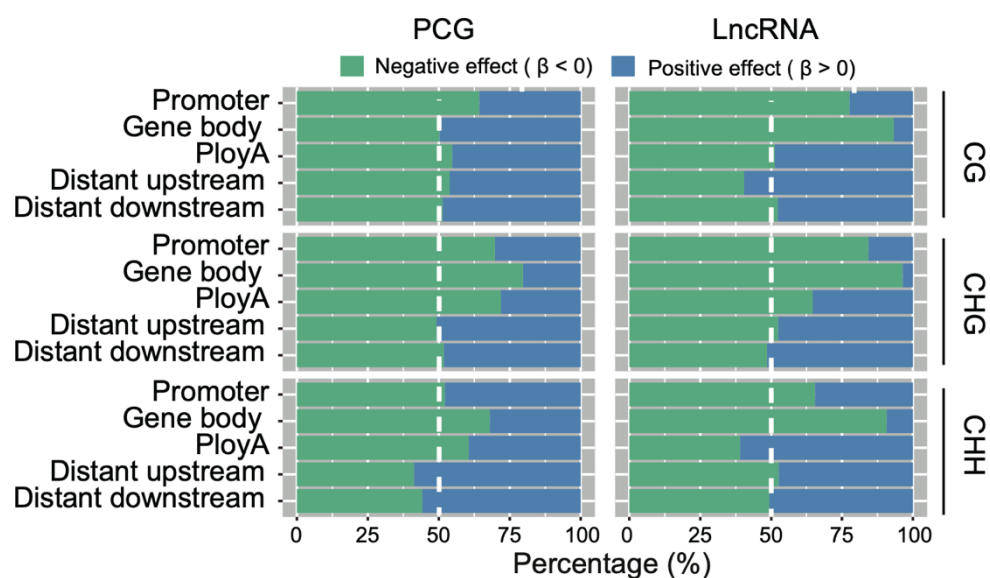




**Supplementary information, Fig. S7. The impact of DNA methylation on gene expression.** The effect of *cis*-eQTM across different genomic feature (Promoter, Gene body, Poly (A), distant upstream (> -2000 bp), and distant downstream (> 2000 bp). The effect of each *cis*-eQTM were grouped into negative ( $\beta < 0$ ) and positive ( $\beta > 0$ ).