(a) Bar chart showing the number of segments with hypomethylation and hypermethylation across different cell lines. (HMC18, Hs578T, MCF7, MDAMB231, SKBR3, T47D).

(b) Scatter plot showing the log2 MeDIP/ Input (HMEC) against the log2 MeDIP/ Input (BCCs) for various positions.

(c) Heatmap indicating hypomethylation and hypermethylation across different chromosomes.

(d) Density plot showing the distribution of CpGs with hypermethylation and hypomethylation.

(e) Histogram showing the fraction of genes with different numbers of CpGs.