

 Supplemental Fig S1. MeD-seq gene body and CpG island correlation analysis.
a) Pearson correlation analysis of MeD-seq read counts of GpG islands for LpnPI-MspJI (left), LpnPI-FspEI (middle) and MspJI-FspEI (right) comparisons.
b) Pearson correlation analysis of MeD-seq read counts of CpG islands for technical replicates digested with LpnPI (left) and MspJI-(right).
c) Pearson correlation analysis of MeD-seq read counts of CpG islands for technical replicates digested with LpnPI (left) and MspJI (right). a) b) c) d) e)

As in (c) but now for gene bodies. Significant DMRs found between LpnPI-MspJI, LpnPI-FspEI and MspJI-FspEI comparisons. The number of hypermethylated DMRs in each comparison is shown.