



Figure S5. Low detection limits of MCTA-Seq.

(A) Of all 3,579 CGIs that were not detected in both WBCs replicates (MePM = 0), the proportion of the loci positively detected in both (blue) or either (red) of the technical replicates of each dilution group (MePM > 0) were shown in histograms. The amount (pg) and the haploid genomic equivalents (GE) of the spiked FMG for each dilution group were shown.

(B) Of the 3,579 CGIs, the proportion of the loci detected with different number of UMIs in technical replicates of the 7.5 pg and 15 pg dilution groups were shown in histograms.