

Additional file 3

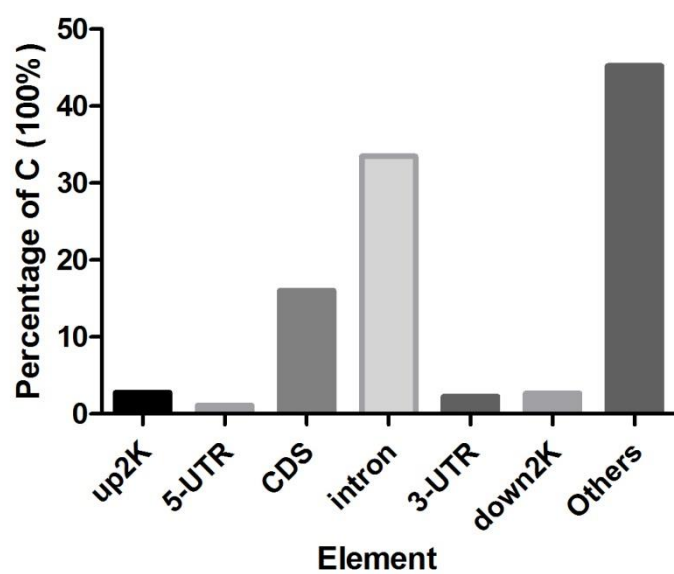


Figure S4. Distribution of differentially methylated loci in gene elements.

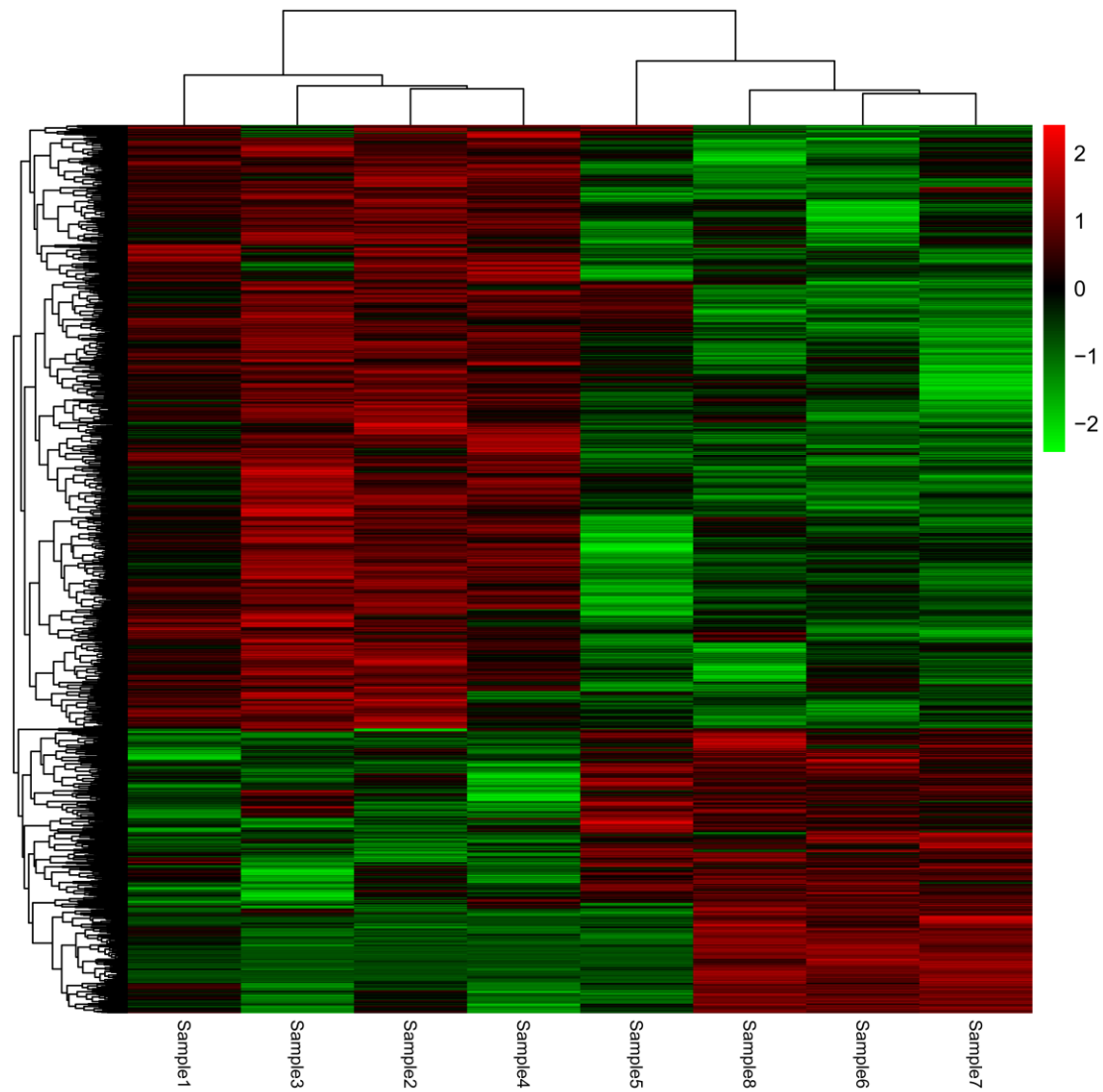


Figure S5. Hierarchical clustering presentation of RRBS sequencing in PGCs. Each horizontal row represents one differentially methylated loci, each column represents one sample. Sample1-4: control group, sample 5-8: GDM group. Increasing green intensities denote loci that was hypomethylated, and increasing red intensities denote loci that was hypermethylated in GDM samples compared with control samples.