

Figure SX: Percentage of Total (all genes), gbM, TE-like, and unmethylated genes with known presence absences variation. A two-sided Fisher's Exact Test was used to test for depletion or enrichment of PAVs amongst each category of genic DNA methylation. *FDR corrected p-value < 0.05, **FDR corrected p-value < 0.01, *** FDR corrected p-value < 0.001, NS = Not significantly different.

