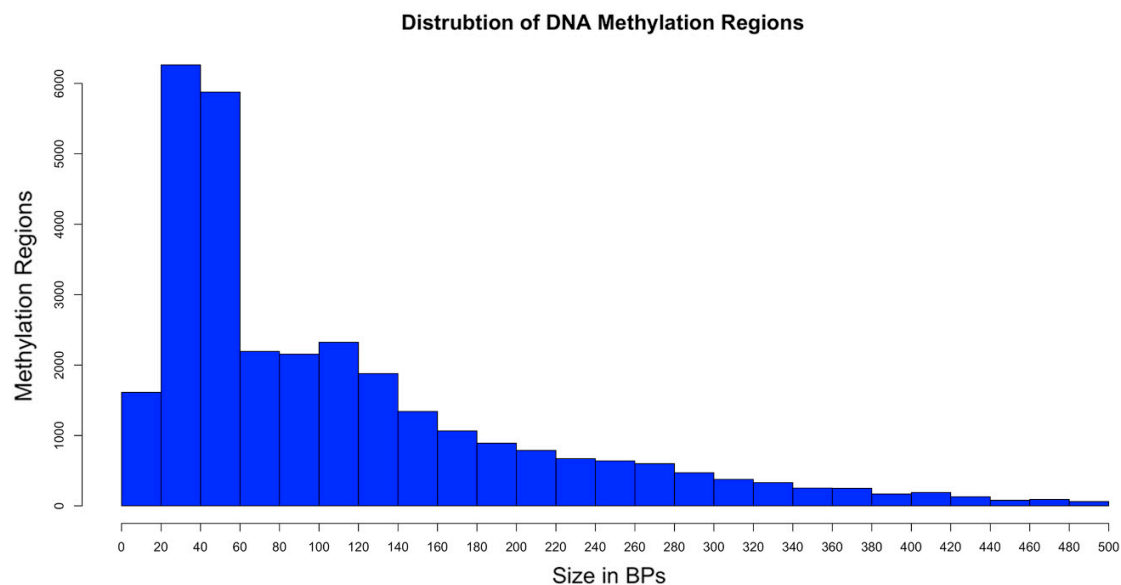


A



B

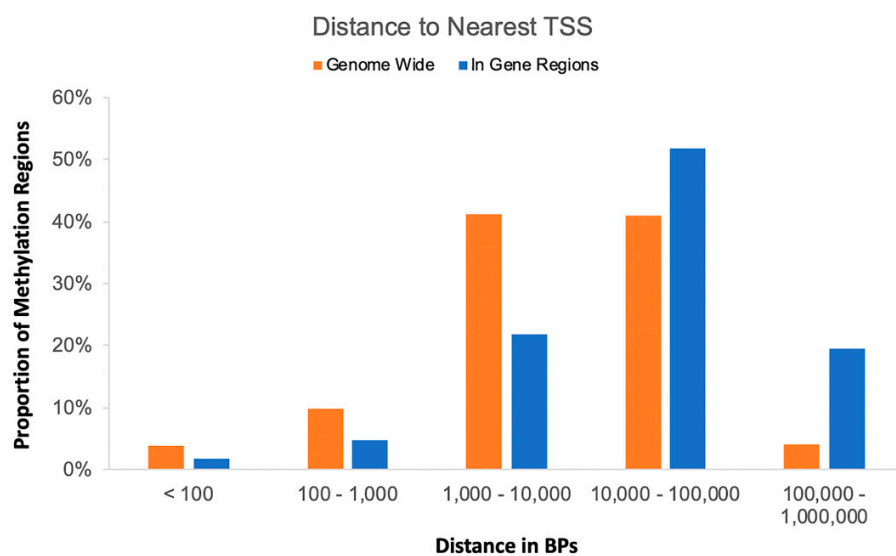


Figure S1. Distribution of methylation regions. (a) Distribution of DNA methylation regions based on size. (b) Distribution of DNA methylation regions based on distance to nearest TSS both pre- and post-filtering to remove regions located outside annotated gene regions.