



Figure S1: Functional genomic location and neighborhood distribution of hypermethylated CpG Sites in TMX2-28 compared to MCF-7 that have decreased methylation after 5-Aza-dC treatment. Pie charts indicate the functional genomic distribution (A) and neighborhood locations (B) defined by Illumina for the 707 CpG sites identified as having increased methylation in TMX2-28 compared to MCF-7 and also decreased methylation in TMX2-28 after 5-Aza-dC treatment.