

The Cancer Genome Atlas Pilot Project
Johns Hopkins University and University of Southern California
Segmented Data Description

Platform: Illumina GoldenGate

Segmented Data

The majority of loci on the Illumina platform will be CpG islands that are normally unmethylated, and for which hypermethylation in the tumor sample will be of interest. Therefore, we will rank loci by the beta value obtained for each locus within each sample, with highly methylated loci of highest interest.