

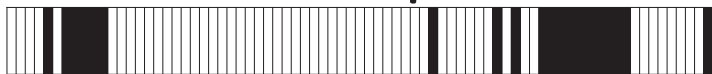
chr10:131,155,100
CpG site 1

Miseq sequencing

MSP

Exon1

chr10:131,155,761
77



Methylation percentage on average

