Scale
chr9:
200 bases
hg38
133,255,100
133,255,200
133,255,300
133,255,400
133,255,500
ASM
0.5
0
Allele-Specific Methylation (ASM) averaged across all samples in hg38
CpGs
Positions of CpG sites in hg38
Sperm
Sperm methylation profiles reveal features of epigenetic inheritance and evolution in primates [Sperm]
Sperm / SRX081759 (CpG methylation)
Sperm / SRX081760 (CpG methylation)
Sperm / SRX081761 (CpG methylation)
HSPC
HSPC / SRX096518 (CpG methylation)
Cord Blood
Cord Blood / SRX096519 (CpG methylation)
B Cell
B Cell / SRX096520 (CpG methylation)
HSPC / SRX096521 (CpG methylation)
Neutrohphil
Neutrohphil / SRX096522 (CpG methylation)
Neutrohphil / SRX096523 (CpG methylation)
Cord Blood / SRX096524 (CpG methylation)
RefSeq genes from NCBI
RefSeq Curated