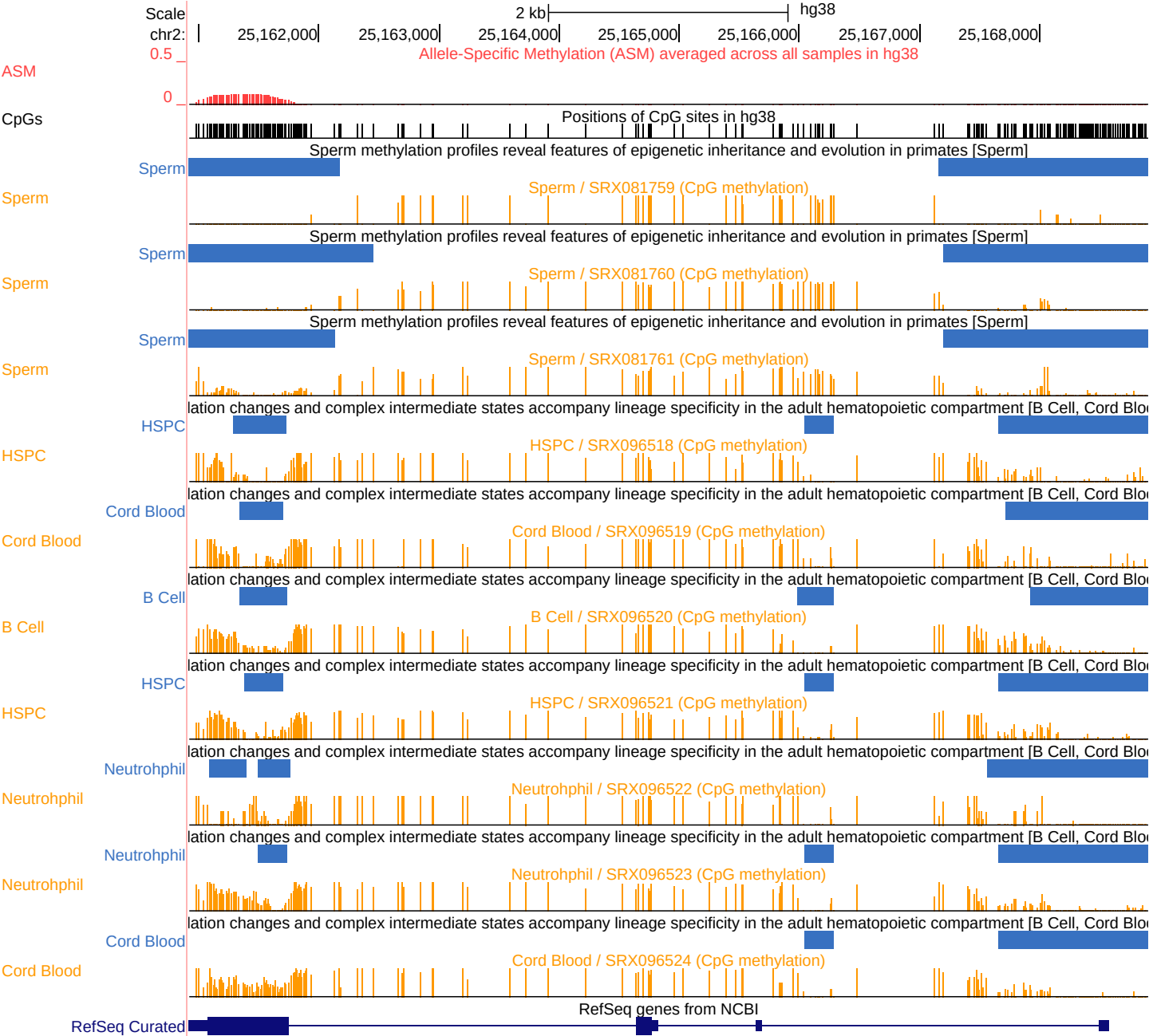
Scale
chr2:
2 kb
hg38
25,162,000
25,163,000
25,164,000
25,165,000
25,166,000
25,167,000
25,168,000
0.5
0
ASM
Allele-Specific Methylation (ASM) averaged across all samples in hg38
CpGs
Positions of CpG sites in hg38
Sperm methylation profiles reveal features of epigenetic inheritance and evolution in primates [Sperm]
Sperm
Sperm / SRX081759 (CpG methylation)
Sperm methylation profiles reveal features of epigenetic inheritance and evolution in primates [Sperm]
Sperm
Sperm / SRX081760 (CpG methylation)
Sperm methylation profiles reveal features of epigenetic inheritance and evolution in primates [Sperm]
Sperm
Sperm / SRX081761 (CpG methylation)
lation changes and complex intermediate states accompany lineage specificity in the adult hematopoietic compartment [B Cell, Cord Blo
HSPC
HSPC / SRX096518 (CpG methylation)
lation changes and complex intermediate states accompany lineage specificity in the adult hematopoietic compartment [B Cell, Cord Blo
Cord Blood
Cord Blood / SRX096519 (CpG methylation)
lation changes and complex intermediate states accompany lineage specificity in the adult hematopoietic compartment [B Cell, Cord Blo
B Cell
B Cell / SRX096520 (CpG methylation)
lation changes and complex intermediate states accompany lineage specificity in the adult hematopoietic compartment [B Cell, Cord Blo
HSPC
HSPC / SRX096521 (CpG methylation)
lation changes and complex intermediate states accompany lineage specificity in the adult hematopoietic compartment [B Cell, Cord Blo
Neutrohphil
Neutrohphil / SRX096522 (CpG methylation)
lation changes and complex intermediate states accompany lineage specificity in the adult hematopoietic compartment [B Cell, Cord Blo
Neutrohphil
Neutrohphil / SRX096523 (CpG methylation)
lation changes and complex intermediate states accompany lineage specificity in the adult hematopoietic compartment [B Cell, Cord Blo
Cord Blood
Cord Blood / SRX096524 (CpG methylation)
RefSeq genes from NCBI
RefSeq Curated