Scale
chr11:
50 kb
hg38
119,000,000
119,050,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
FOXR1
Metazoa_SRP
RP11-110I1.12
CCDC84
TRAPPC4
SLC37A4
RP11-110I1.6
HYOU1
VPS11
RP11-110I1.11
RPS25
RP11-110I1.14
HMBS
MIR3656
RP11-110I1.13
RefSeq Curated
RefSeq gene predictions from NCBI
NONCODE_v5_Gene
NONHSAG063137.1
NONHSAG009789.3
NONHSAG009788.2
NONHSAG009791.2
NONHSAG009792.2
NONHSAG009793.2
NONHSAG009790.2
NONHSAG009794.2
NONHSAG009796.2
NONHSAG009797.2
NONHSAG009799.2
NONHSAG009800.3
NONCODE_v5_lncRNA
NONHSAT159746.1
NONHSAT230847.1
NONHSAT230846.1
NONHSAT024534.2
NONHSAT024531.2
NONHSAT230848.1
NONHSAT024532.2
NONHSAT024536.2
NONHSAT024538.2
NONHSAT024539.2
NONHSAT159749.1
NONHSAT024545.2
NONHSAT024544.2
NONHSAT024543.2
NONHSAT024559.2
NONHSAT024560.2
NONHSAT024563.2
NONHSAT159750.1
NONHSAT024567.2
NONHSAT230849.1
NONHSAT230850.1
NONHSAT024569.2
NONHSAT024570.2
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
RPL23AP64
H3K27Ac Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE
Layered H3K27Ac
No such file or directory
Can't open /gbdb/hg38/bbi/wgEncodeReg/wgEncodeRegMarkH3k27ac/wgEncodeBroadHistoneGm12878H3k27acStdSig.bigWig to read
DNase Clusters
DNase I Hypersensitivity Peak Clusters from ENCODE (95 cell types)
Repeating Elements by RepeatMasker
SINE
LINE
LTR
DNA
Simple
Low Complexity
Satellite
RNA
Other
Unknown