Scale
chr2:
200 kb
hg38
206,000,000
206,100,000
206,200,000
206,300,000
206,400,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
RN7SKP178
AC007679.3
INO80D
NDUFS1
GPR1
ZDBF2
GPR1-AS
AC017081.1
Vault
AC007383.3
RN7SKP200
Y_RNA
EEF1B2
RN7SKP260
AC007383.4
uc_338
snoZ196
SNORD51
SNORA41
RefSeq Curated
RefSeq gene predictions from NCBI
NONCODE_v5_Gene
NONHSAG030329.2
NONHSAG078110.1
NONHSAG030330.2
NONHSAG078111.1
NONHSAG080947.1
NONHSAG080948.1
NONHSAG030331.2
NONHSAG030332.2
NONHSAG079355.1
NONHSAG030333.3
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NONHSAG030334.2
NONHSAG079357.1
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NONHSAG030336.3
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NONHSAG030338.2
NONHSAG078112.1
NONHSAG079358.1
NONHSAG078113.1
NONHSAG030342.3
NONHSAG030340.2
NONHSAG030341.2
NONHSAG030343.2
NONHSAG080949.1
NONHSAG030344.2
NONHSAG080950.1
NONCODE_v5_lncRNA
NONHSAT076529.2
NONHSAT183439.1
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NONHSAT183440.1
NONHSAT187981.1
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NONHSAT186241.1
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NONHSAT241359.1
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NONHSAT241362.1
NONHSAT187983.1
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NONHSAT187984.1
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
AC007679.1
AC007383.6
AC017081.2
HNRNPA1P51
HMGN1P6
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