

Scale chr18:

22,450,000 | 50 kb | hg38 | 22,500,000 | 22,550,000 |
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)

CTAGE1
CTAGE1

RefSeq gene predictions from NCBI

NONCODE_v5_Gene

NONCODE_v5_lncRNA

OMIM Allelic Variants

Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)

RPS4XP18

CTAGE1 <<<<
RP11-863N1.4

H3K27Ac Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE

No such file or directory

Can't open /gdb/hg38/bbi/wgEncodeReg/wgEncodeRegMarkH3k27ac/wgEncodeBroadHistoneGm12878H3k27acStdSig.bigWig to read

Layered H3K27Ac

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