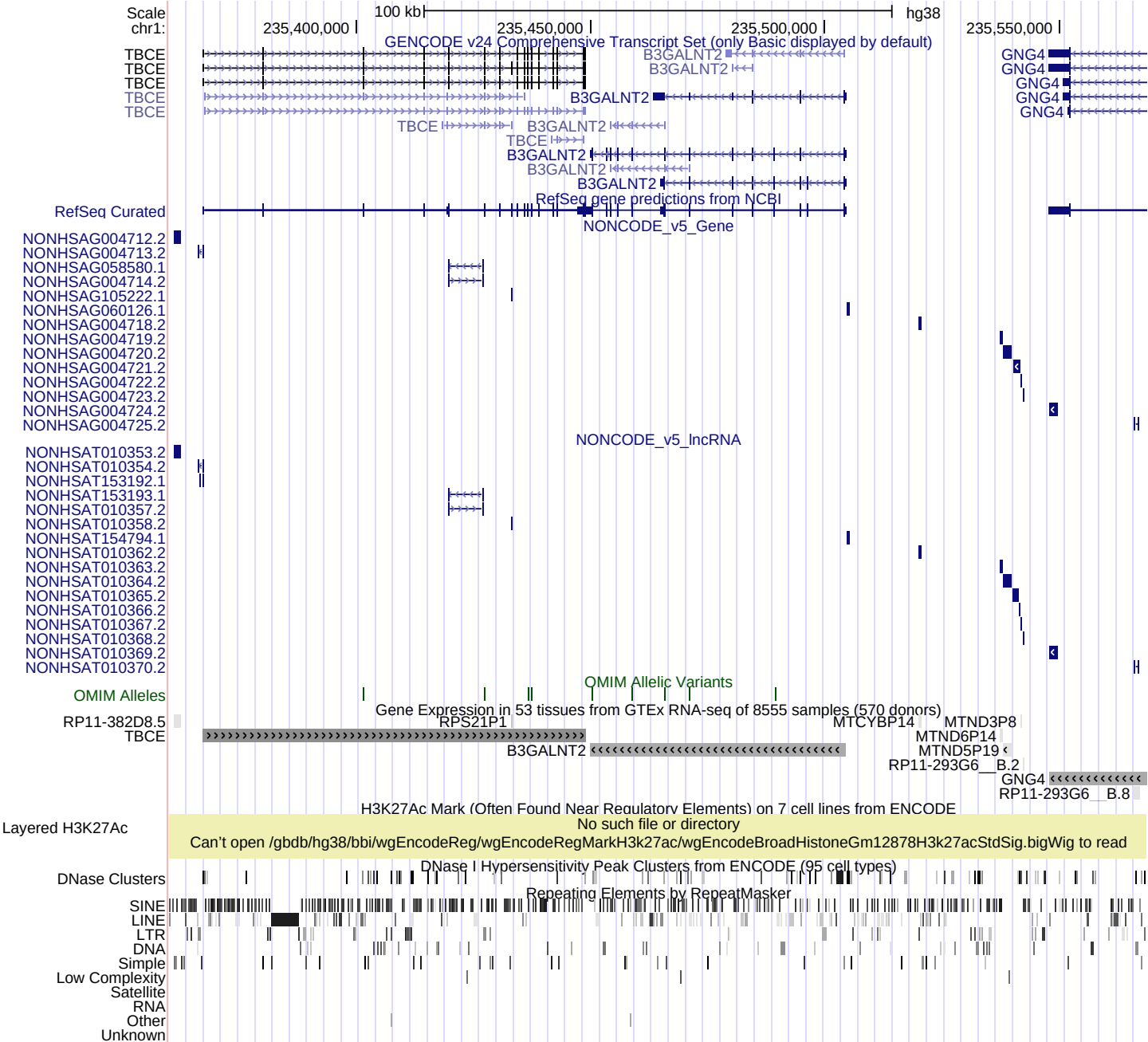
Scale
chr1:
100 kb
hg38
235,400,000
235,450,000
235,500,000
235,550,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
TBCE
B3GALNT2
GNG4
RefSeq Curated
RefSeq gene predictions from NCBI
NONCODE_v5_Gene
NONHSAG004712.2
NONHSAG004713.2
NONHSAG058580.1
NONHSAG004714.2
NONHSAG105222.1
NONHSAG060126.1
NONHSAG004718.2
NONHSAG004719.2
NONHSAG004720.2
NONHSAG004721.2
NONHSAG004722.2
NONHSAG004723.2
NONHSAG004724.2
NONHSAG004725.2
NONCODE_v5_lncRNA
NONHSAT010353.2
NONHSAT010354.2
NONHSAT153192.1
NONHSAT153193.1
NONHSAT010357.2
NONHSAT010358.2
NONHSAT154794.1
NONHSAT010362.2
NONHSAT010363.2
NONHSAT010364.2
NONHSAT010365.2
NONHSAT010366.2
NONHSAT010367.2
NONHSAT010368.2
NONHSAT010369.2
NONHSAT010370.2
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
RP11-382D8.5
RPS21P1
MTCYBP14
MTND3P8
MTND6P14
MTND5P19
RP11-293G6__B.2
RP11-293G6__B.8
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 I Hypersensitivity Peak Clusters from ENCODE (95 cell types)
DNase Clusters
Repeating Elements by RepeatMasker
SINE
LINE
LTR
DNA
Simple
Low Complexity
Satellite
RNA
Other
Unknown