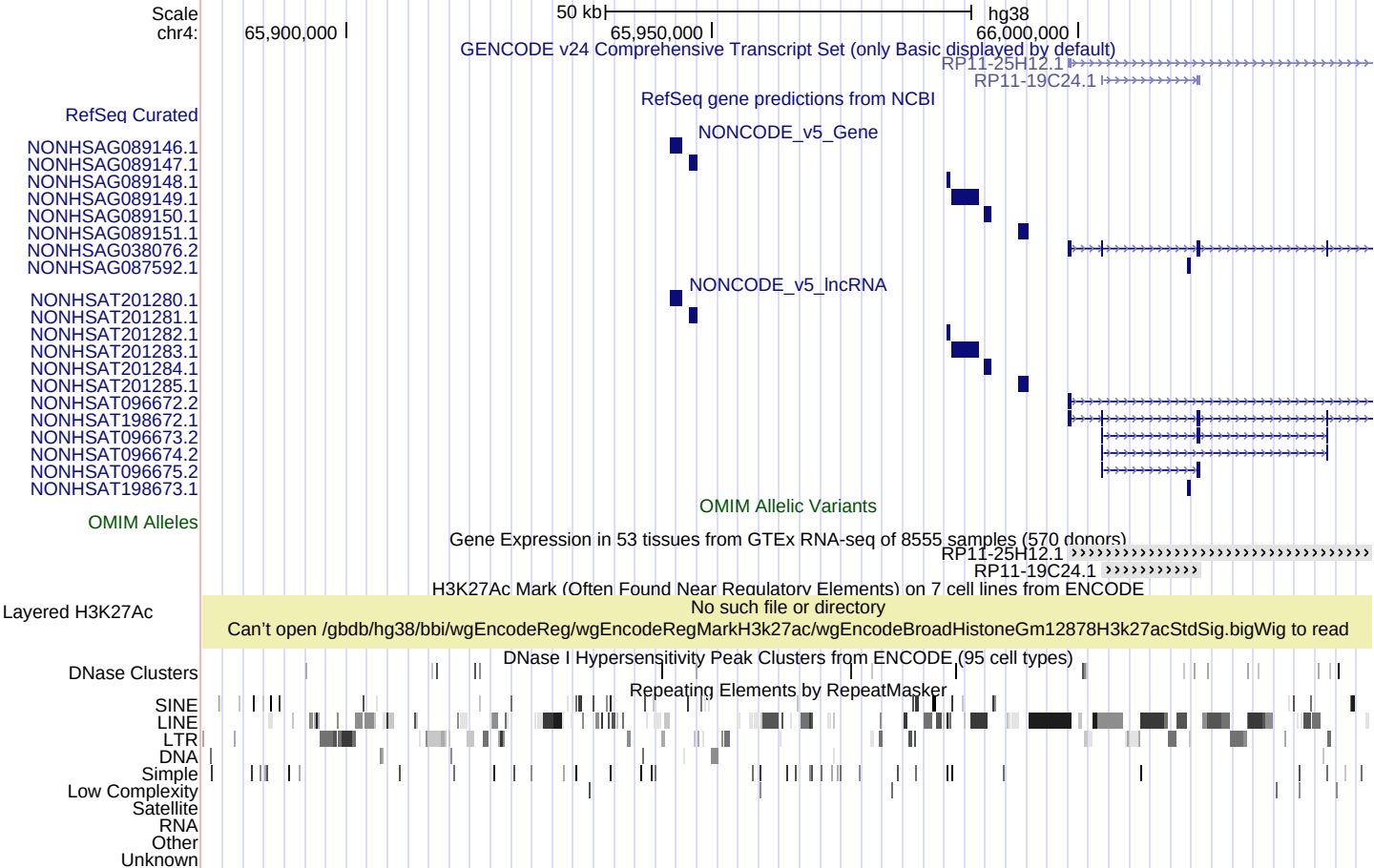
Scale
chr4:
50 kb
hg38
65,900,000
65,950,000
66,000,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
RP11-25H12.1
RP11-19C24.1
RefSeq gene predictions from NCBI
RefSeq Curated
NONCODE_v5_Gene
NONHSAG089146.1
NONHSAG089147.1
NONHSAG089148.1
NONHSAG089149.1
NONHSAG089150.1
NONHSAG089151.1
NONHSAG038076.2
NONHSAG087592.1
NONCODE_v5_lncRNA
NONHSAT201280.1
NONHSAT201281.1
NONHSAT201282.1
NONHSAT201283.1
NONHSAT201284.1
NONHSAT201285.1
NONHSAT096672.2
NONHSAT198672.1
NONHSAT096673.2
NONHSAT096674.2
NONHSAT096675.2
NONHSAT198673.1
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
RP11-25H12.1
RP11-19C24.1
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