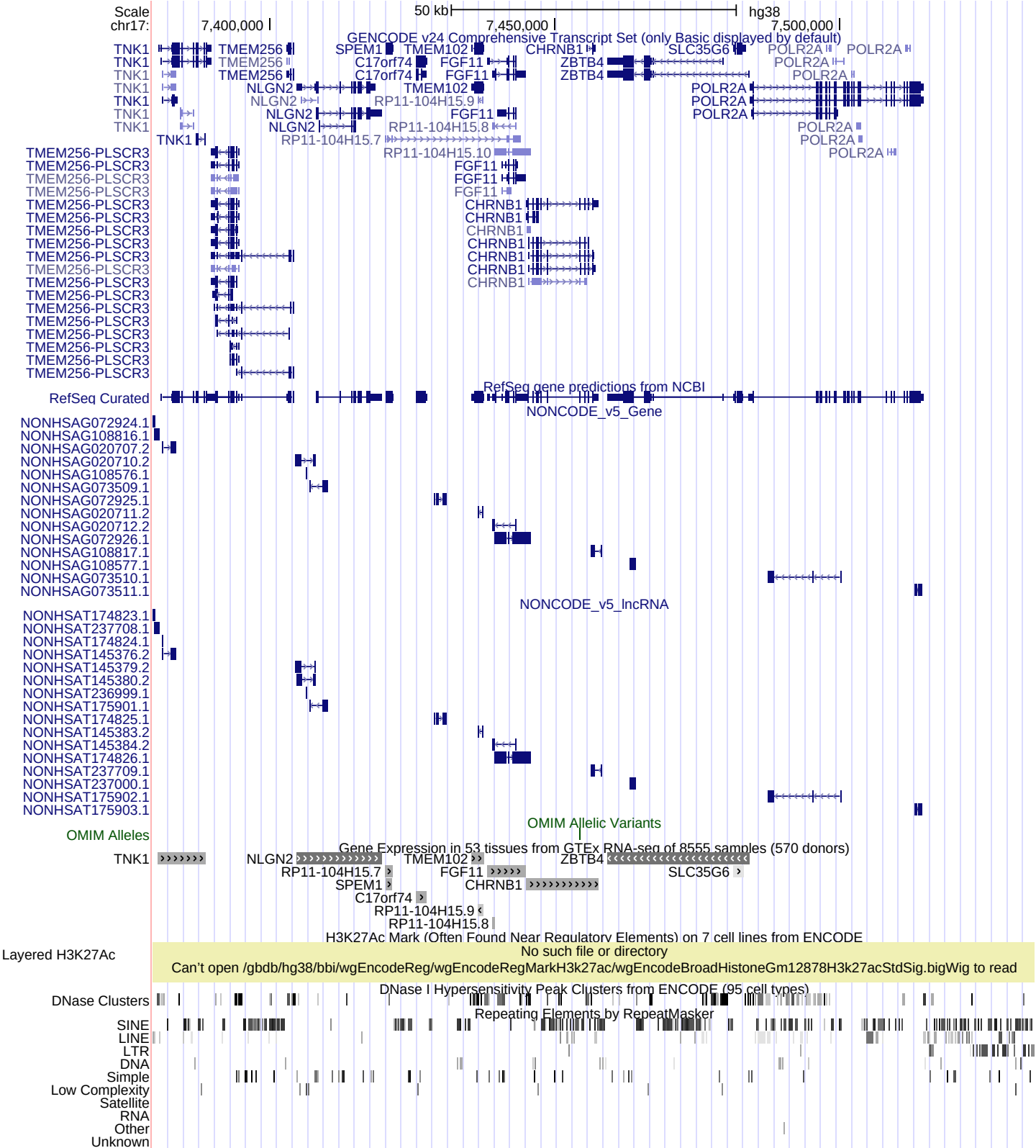
Scale
chr17:
50 kb
hg38
7,400,000
7,450,000
7,500,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
TNK1
TMEM256
SPEM1
TMEM102
CHRNB1
SLC35G6
POLR2A
C17orf74
FGF11
ZBTB4
NLGN2
RP11-104H15.9
RP11-104H15.8
RP11-104H15.7
RP11-104H15.10
TMEM256-PLSCR3
RefSeq gene predictions from NCBI
RefSeq Curated
NONCODE_v5_Gene
NONHSAG072924.1
NONHSAG108816.1
NONHSAG020707.2
NONHSAG020710.2
NONHSAG108576.1
NONHSAG073509.1
NONHSAG072925.1
NONHSAG020711.2
NONHSAG020712.2
NONHSAG072926.1
NONHSAG108817.1
NONHSAG108577.1
NONHSAG073510.1
NONHSAG073511.1
NONCODE_v5_lncRNA
NONHSAT174823.1
NONHSAT237708.1
NONHSAT174824.1
NONHSAT145376.2
NONHSAT145379.2
NONHSAT145380.2
NONHSAT236999.1
NONHSAT175901.1
NONHSAT174825.1
NONHSAT145383.2
NONHSAT145384.2
NONHSAT174826.1
NONHSAT237709.1
NONHSAT237000.1
NONHSAT175902.1
NONHSAT175903.1
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
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