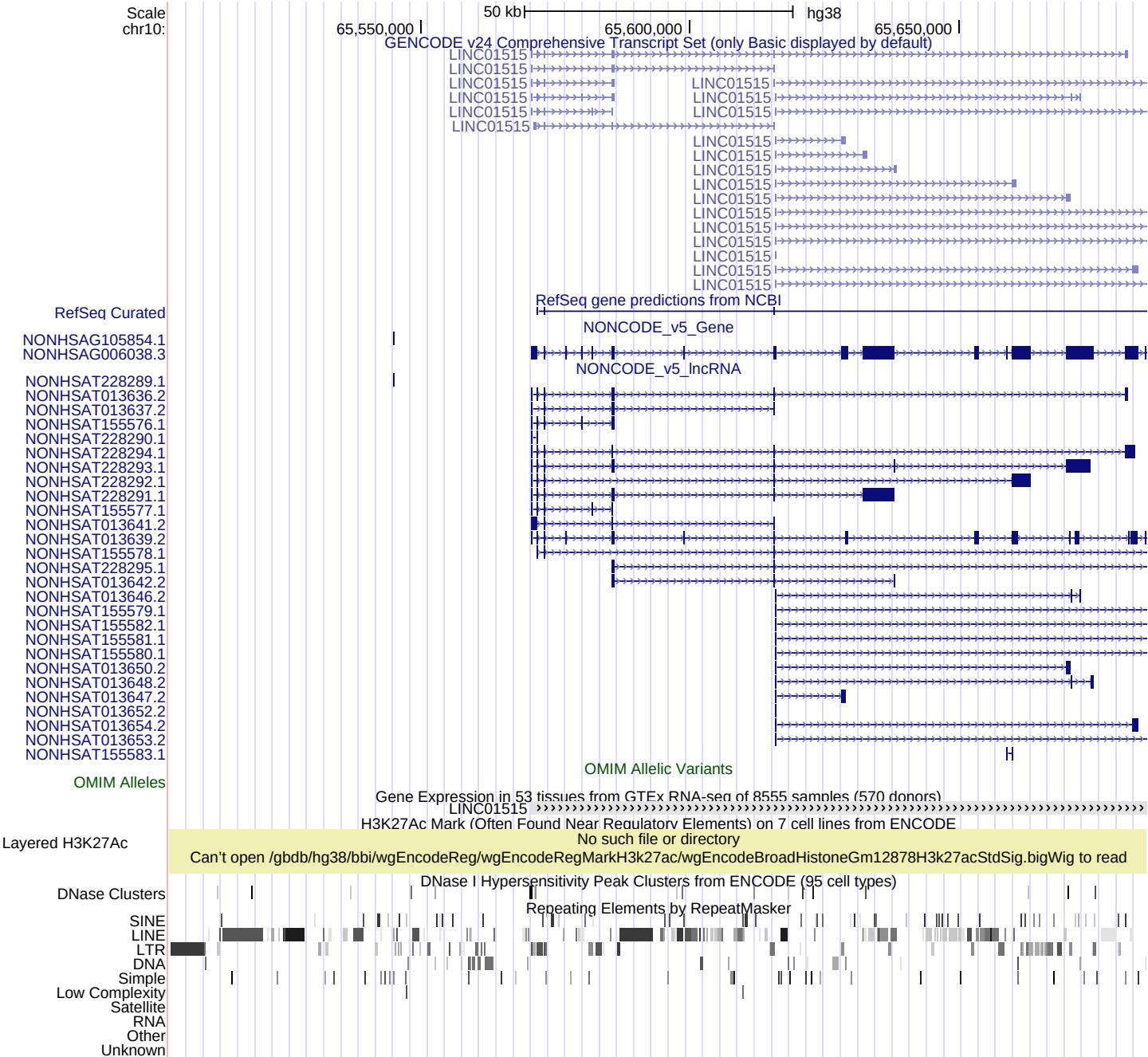
Scale
chr10:
50 kb
hg38
65,550,000
65,600,000
65,650,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
LINC01515
RefSeq Curated
RefSeq gene predictions from NCBI
NONCODE_v5_Gene
NONHSAG105854.1
NONHSAG006038.3
NONCODE_v5_lncRNA
NONHSAT228289.1
NONHSAT013636.2
NONHSAT013637.2
NONHSAT155576.1
NONHSAT228290.1
NONHSAT228294.1
NONHSAT228293.1
NONHSAT228292.1
NONHSAT228291.1
NONHSAT155577.1
NONHSAT013641.2
NONHSAT013639.2
NONHSAT155578.1
NONHSAT228295.1
NONHSAT013642.2
NONHSAT013646.2
NONHSAT155579.1
NONHSAT155582.1
NONHSAT155581.1
NONHSAT155580.1
NONHSAT013650.2
NONHSAT013648.2
NONHSAT013647.2
NONHSAT013652.2
NONHSAT013654.2
NONHSAT013653.2
NONHSAT155583.1
OMIM Alleles
OMIM Allelic Variants
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
LINC01515
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