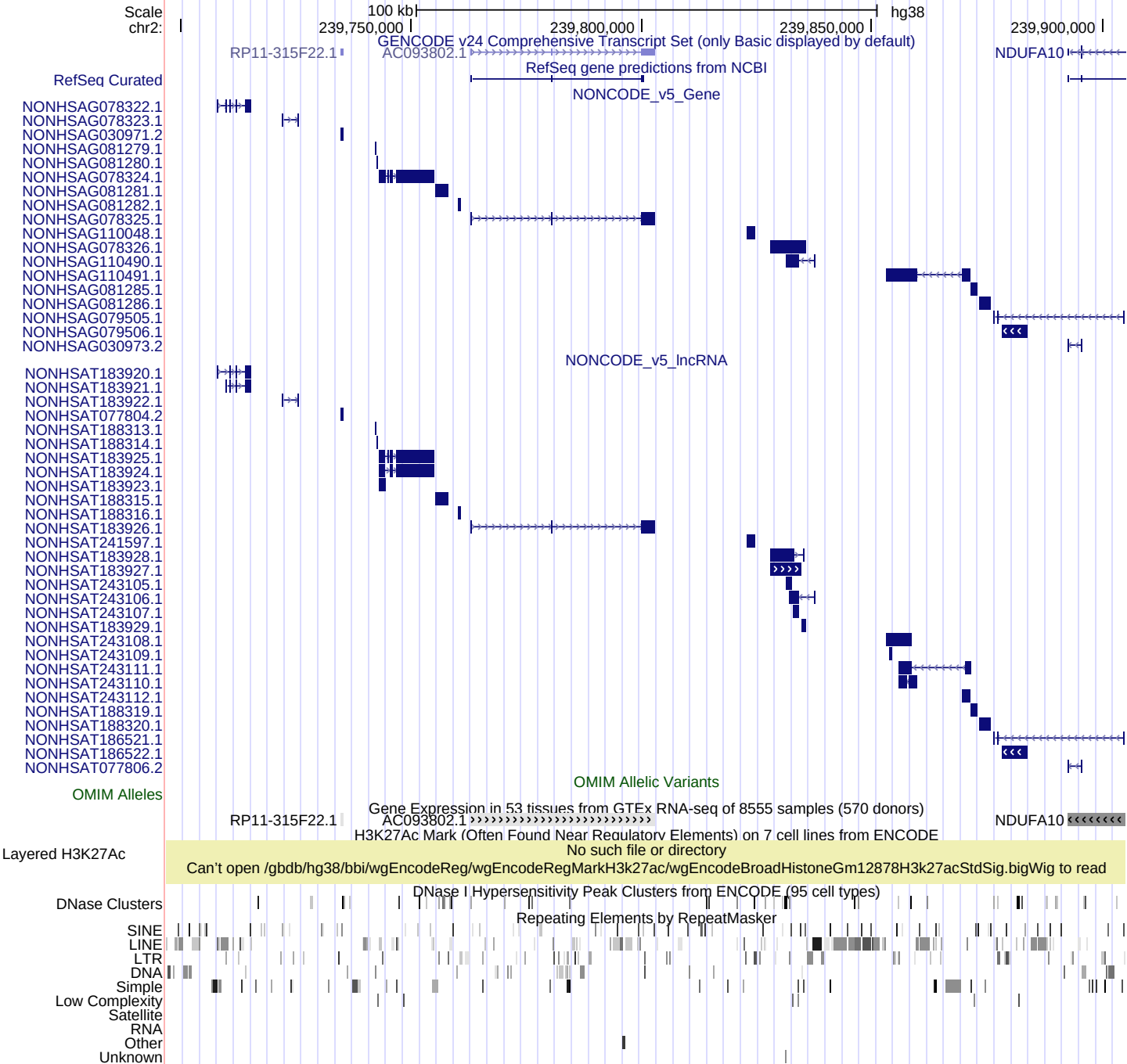
Scale
chr2:
100 kb
hg38
239,750,000
239,800,000
239,850,000
239,900,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
RP11-315F22.1
AC093802.1
NDUFA10
RefSeq gene predictions from NCBI
RefSeq Curated
NONCODE_v5_Gene
NONHSAG078322.1
NONHSAG078323.1
NONHSAG030971.2
NONHSAG081279.1
NONHSAG081280.1
NONHSAG078324.1
NONHSAG081281.1
NONHSAG081282.1
NONHSAG078325.1
NONHSAG110048.1
NONHSAG078326.1
NONHSAG110490.1
NONHSAG110491.1
NONHSAG081285.1
NONHSAG081286.1
NONHSAG079505.1
NONHSAG079506.1
NONHSAG030973.2
NONCODE_v5_lncRNA
NONHSAT183920.1
NONHSAT183921.1
NONHSAT183922.1
NONHSAT077804.2
NONHSAT188313.1
NONHSAT188314.1
NONHSAT183925.1
NONHSAT183924.1
NONHSAT183923.1
NONHSAT188315.1
NONHSAT188316.1
NONHSAT183926.1
NONHSAT241597.1
NONHSAT183928.1
NONHSAT183927.1
NONHSAT243105.1
NONHSAT243106.1
NONHSAT243107.1
NONHSAT183929.1
NONHSAT243108.1
NONHSAT243109.1
NONHSAT243111.1
NONHSAT243110.1
NONHSAT243112.1
NONHSAT188319.1
NONHSAT188320.1
NONHSAT186521.1
NONHSAT186522.1
NONHSAT077806.2
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
RP11-315F22.1
AC093802.1
NDUFA10
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