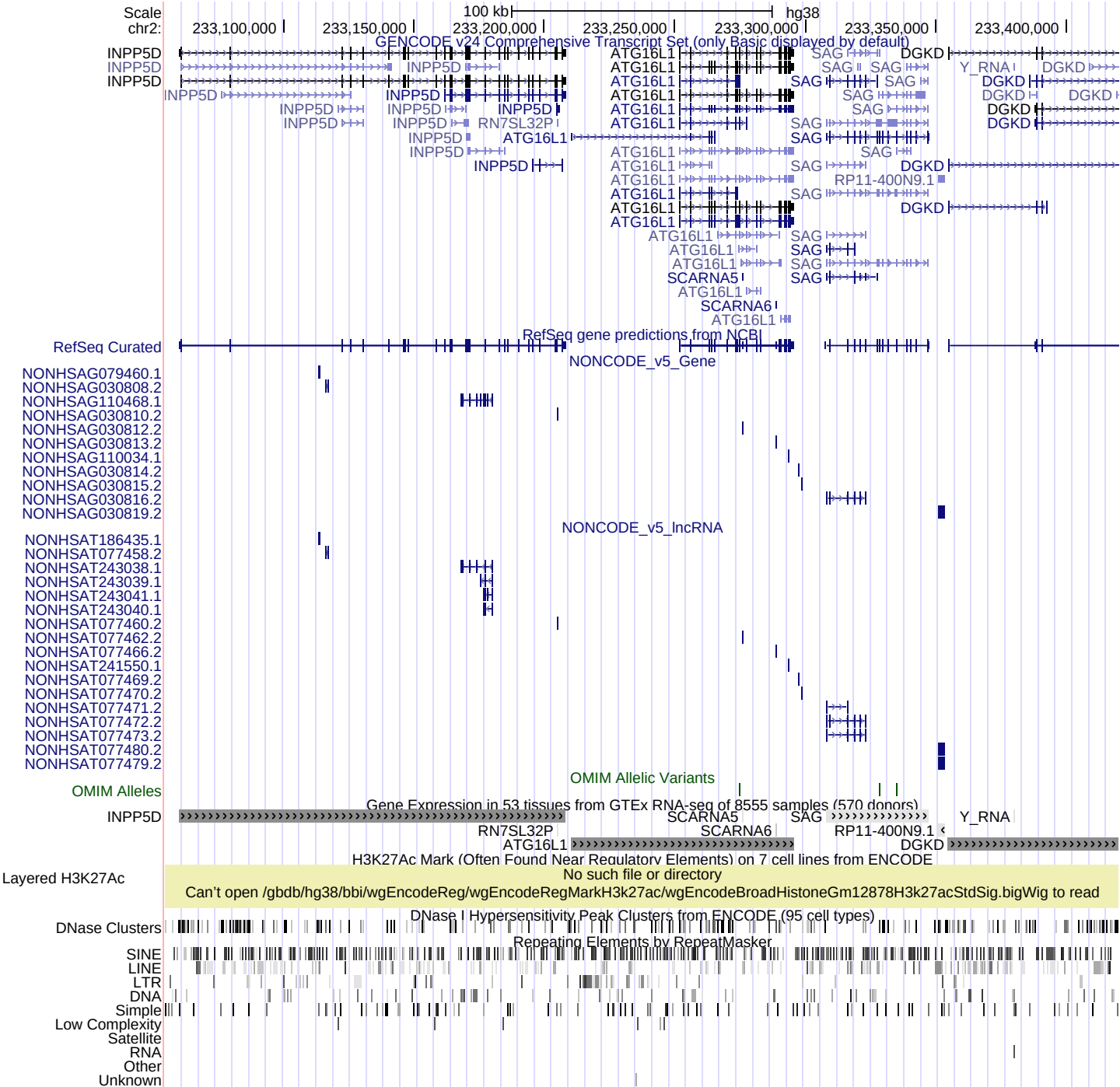
Scale
chr2:
100 kb
hg38
233,100,000
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233,400,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
INPP5D
ATG16L1
SAG
DGKD
Y_RNA
RN7SL32P
SCARNA5
SCARNA6
RP11-400N9.1
RefSeq Curated
RefSeq gene predictions from NCBI
NONCODE_v5_Gene
NONHSAG079460.1
NONHSAG030808.2
NONHSAG110468.1
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NONHSAG030819.2
NONCODE_v5_lncRNA
NONHSAT186435.1
NONHSAT077458.2
NONHSAT243038.1
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NONHSAT241550.1
NONHSAT077469.2
NONHSAT077470.2
NONHSAT077471.2
NONHSAT077472.2
NONHSAT077473.2
NONHSAT077480.2
NONHSAT077479.2
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