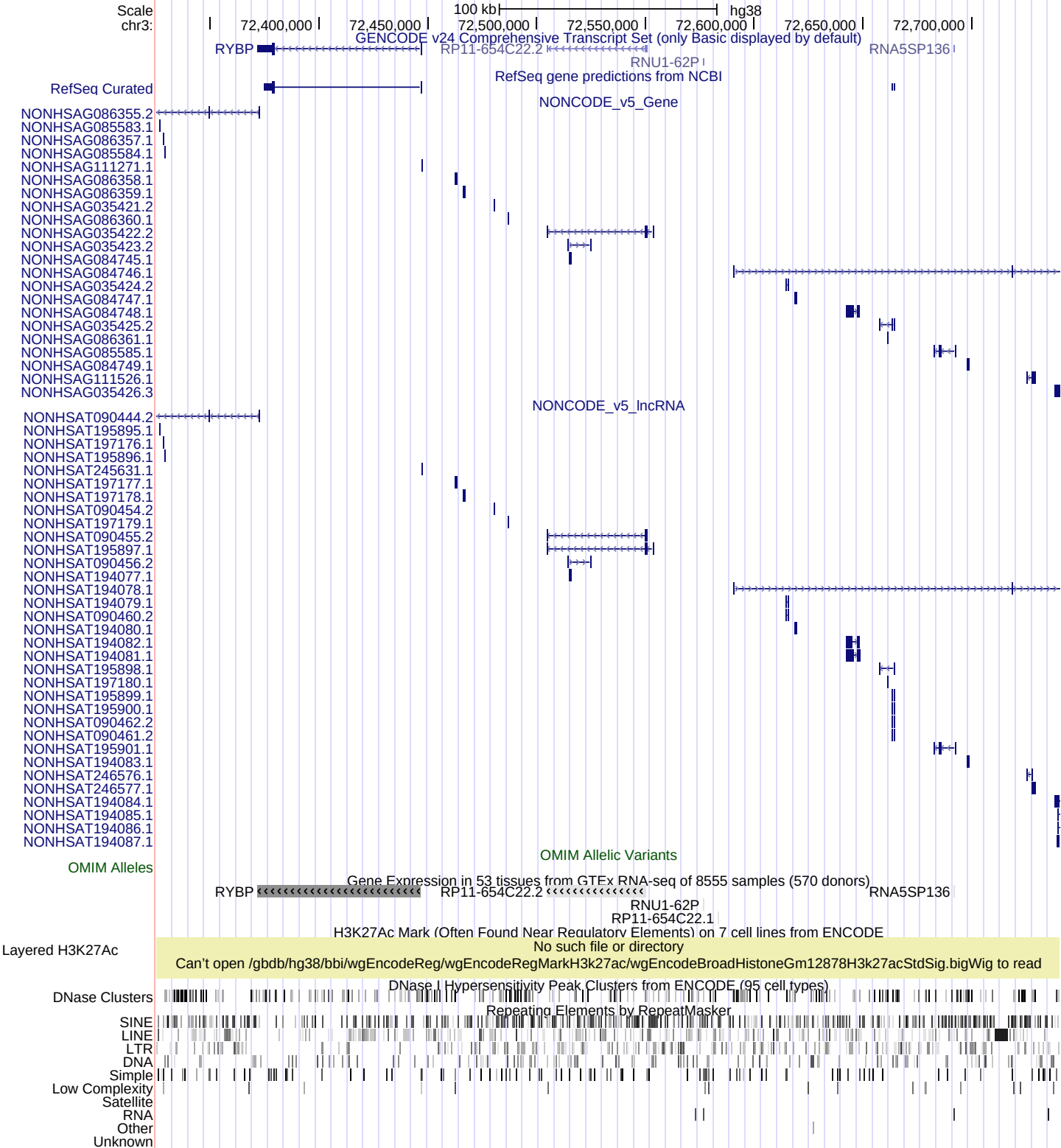
Scale
chr3:
100 kb
hg38
72,400,000
72,450,000
72,500,000
72,550,000
72,600,000
72,650,000
72,700,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
RYBP
RP11-654C22.2
RNA5SP136
RNU1-62P
RefSeq gene predictions from NCBI
RefSeq Curated
NONCODE_v5_Gene
NONHSAG086355.2
NONHSAG085583.1
NONHSAG086357.1
NONHSAG085584.1
NONHSAG111271.1
NONHSAG086358.1
NONHSAG086359.1
NONHSAG035421.2
NONHSAG086360.1
NONHSAG035422.2
NONHSAG035423.2
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NONHSAG084746.1
NONHSAG035424.2
NONHSAG084747.1
NONHSAG084748.1
NONHSAG035425.2
NONHSAG086361.1
NONHSAG085585.1
NONHSAG084749.1
NONHSAG111526.1
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NONCODE_v5_lncRNA
NONHSAT090444.2
NONHSAT195895.1
NONHSAT197176.1
NONHSAT195896.1
NONHSAT245631.1
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NONHSAT195898.1
NONHSAT197180.1
NONHSAT195899.1
NONHSAT195900.1
NONHSAT090462.2
NONHSAT090461.2
NONHSAT195901.1
NONHSAT194083.1
NONHSAT246576.1
NONHSAT246577.1
NONHSAT194084.1
NONHSAT194085.1
NONHSAT194086.1
NONHSAT194087.1
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
RYBP
RP11-654C22.2
RNA5SP136
RNU1-62P
RP11-654C22.1
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 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