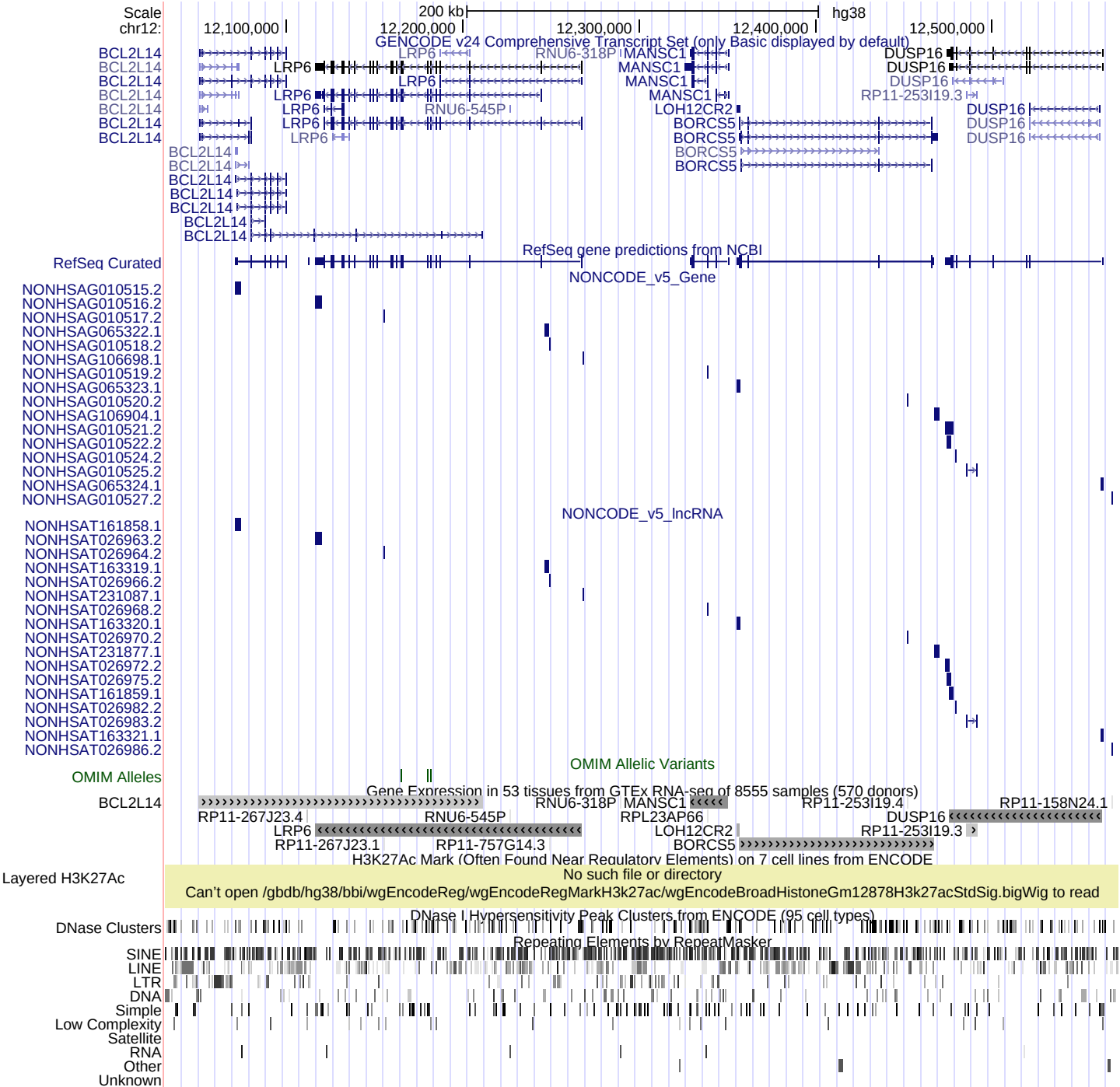
Scale
chr12:
200 kb
hg38
12,100,000
12,200,000
12,300,000
12,400,000
12,500,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
BCL2L14
LRP6
RNU6-318P
MANSC1
DUSP16
RP11-253I19.3
RNU6-545P
LOH12CR2
BORCS5
RefSeq Curated
RefSeq gene predictions from NCBI
NONCODE_v5_Gene
NONHSAG010515.2
NONHSAG010516.2
NONHSAG010517.2
NONHSAG065322.1
NONHSAG010518.2
NONHSAG106698.1
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NONHSAG010524.2
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NONHSAG065324.1
NONHSAG010527.2
NONCODE_v5_lncRNA
NONHSAT161858.1
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NONHSAT026982.2
NONHSAT026983.2
NONHSAT163321.1
NONHSAT026986.2
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
RP11-267J23.4
RP11-158N24.1
RPL23AP66
RP11-267J23.1
RP11-757G14.3
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