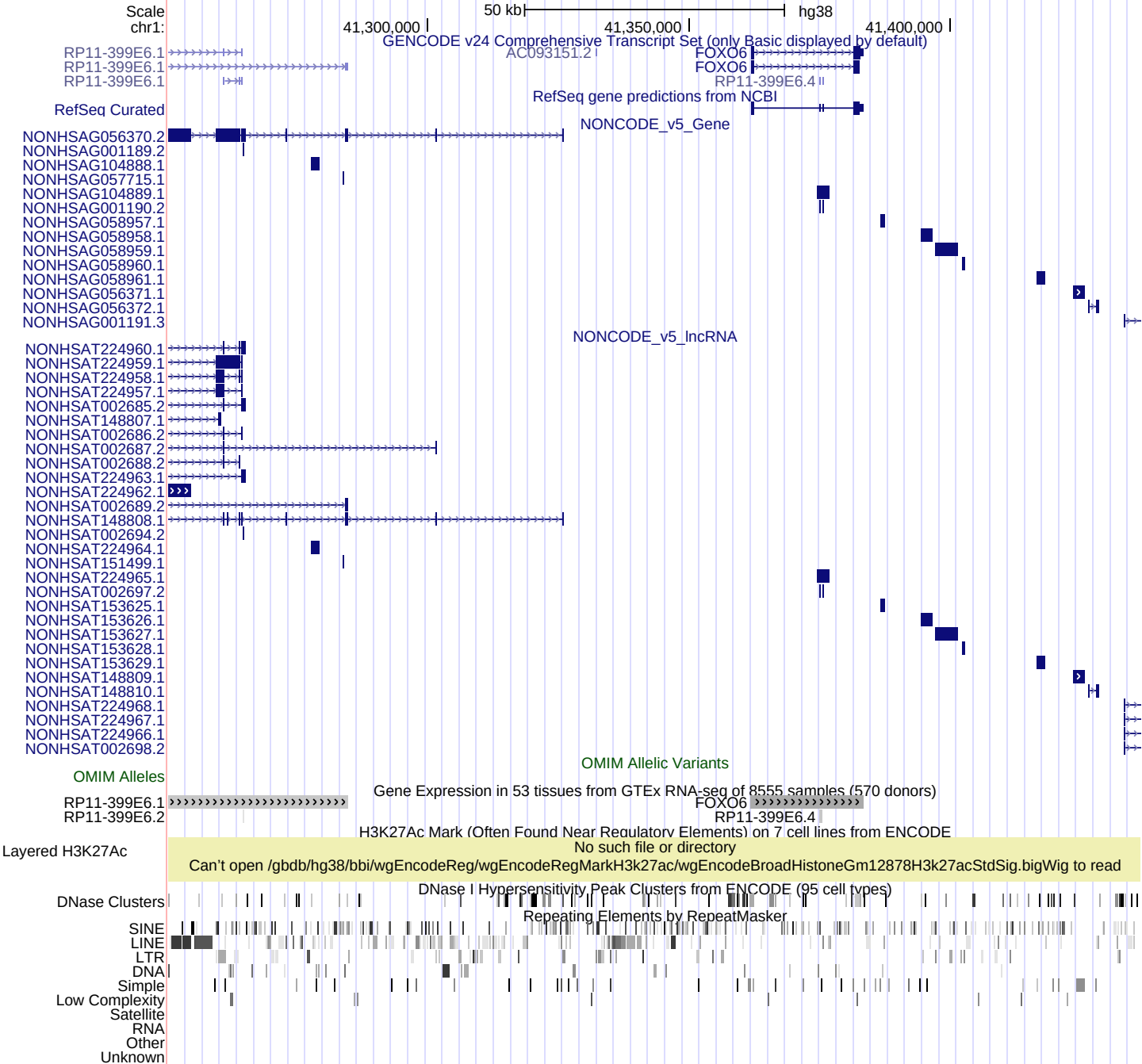
Scale
chr1:
50 kb
hg38
41,300,000
41,350,000
41,400,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
RP11-399E6.1
RP11-399E6.1
RP11-399E6.1
AC093151.2
FOXO6
FOXO6
RP11-399E6.4
RefSeq gene predictions from NCBI
RefSeq Curated
NONCODE_v5_Gene
NONHSAG056370.2
NONHSAG001189.2
NONHSAG104888.1
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NONHSAG104889.1
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NONHSAG001191.3
NONCODE_v5_lncRNA
NONHSAT224960.1
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NONHSAT224958.1
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NONHSAT148807.1
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NONHSAT002688.2
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NONHSAT224962.1
NONHSAT002689.2
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NONHSAT224968.1
NONHSAT224967.1
NONHSAT224966.1
NONHSAT002698.2
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
RP11-399E6.1
RP11-399E6.2
FOXO6
RP11-399E6.4
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