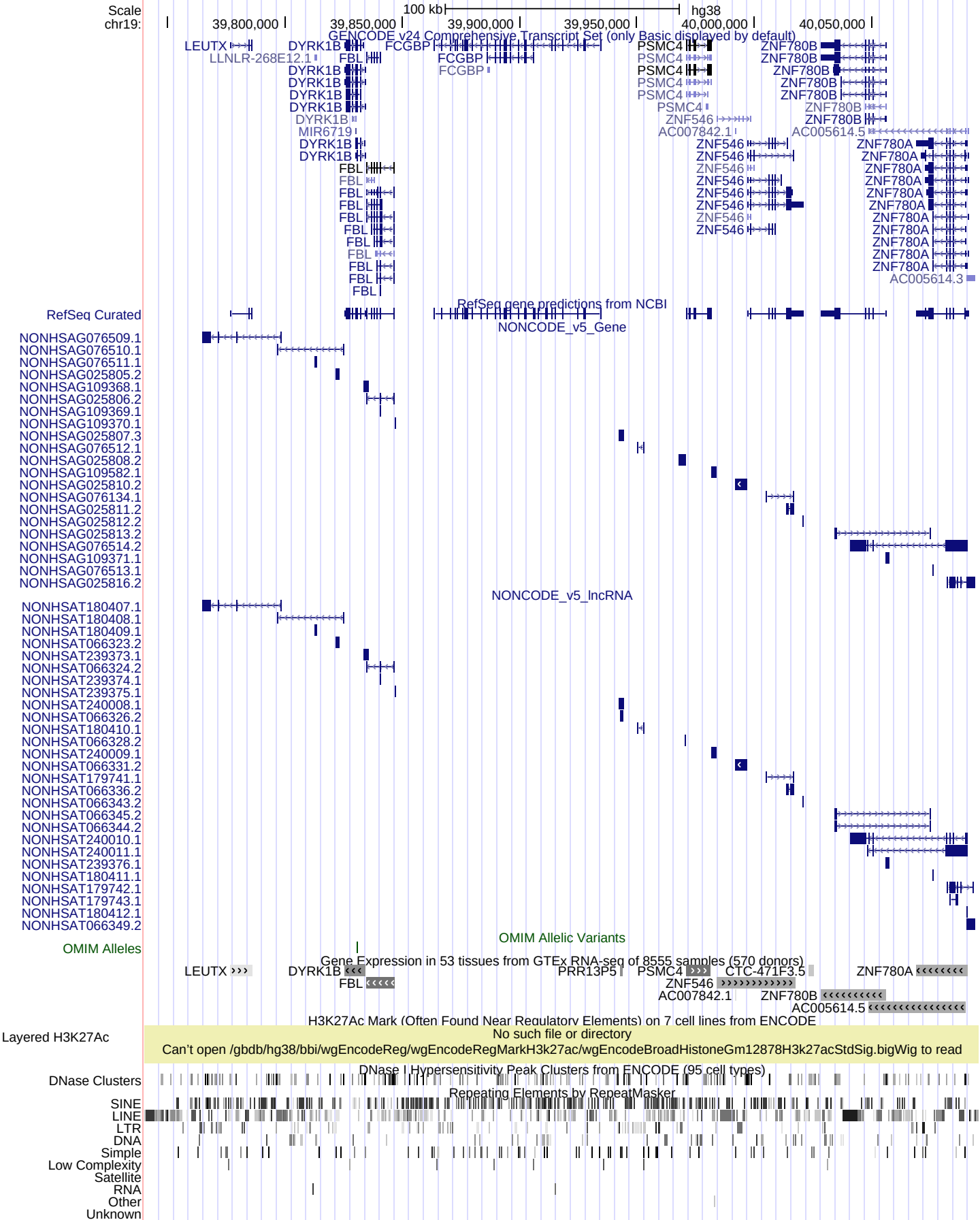
Scale
chr19:
100 kb
hg38
39,800,000
39,850,000
39,900,000
39,950,000
40,000,000
40,050,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
LEUTX
LLNLR-268E12.1
DYRK1B
FBL
FCGBP
PSMC4
ZNF780B
ZNF546
AC007842.1
AC005614.5
ZNF780A
MIR6719
AC005614.3
RefSeq gene predictions from NCBI
RefSeq Curated
NONCODE_v5_Gene
NONHSAG076509.1
NONHSAG076510.1
NONHSAG076511.1
NONHSAG025805.2
NONHSAG109368.1
NONHSAG025806.2
NONHSAG109369.1
NONHSAG109370.1
NONHSAG025807.3
NONHSAG076512.1
NONHSAG025808.2
NONHSAG109582.1
NONHSAG025810.2
NONHSAG076134.1
NONHSAG025811.2
NONHSAG025812.2
NONHSAG025813.2
NONHSAG076514.2
NONHSAG109371.1
NONHSAG076513.1
NONHSAG025816.2
NONCODE_v5_lncRNA
NONHSAT180407.1
NONHSAT180408.1
NONHSAT180409.1
NONHSAT066323.2
NONHSAT239373.1
NONHSAT066324.2
NONHSAT239374.1
NONHSAT239375.1
NONHSAT240008.1
NONHSAT066326.2
NONHSAT180410.1
NONHSAT066328.2
NONHSAT240009.1
NONHSAT066331.2
NONHSAT179741.1
NONHSAT066336.2
NONHSAT066343.2
NONHSAT066345.2
NONHSAT066344.2
NONHSAT240010.1
NONHSAT240011.1
NONHSAT239376.1
NONHSAT180411.1
NONHSAT179742.1
NONHSAT179743.1
NONHSAT180412.1
NONHSAT066349.2
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
LEUTX
DYRK1B
PRR13P5
PSMC4
CTC-471F3.5
ZNF780A
FBL
ZNF546
AC007842.1
ZNF780B
AC005614.5
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