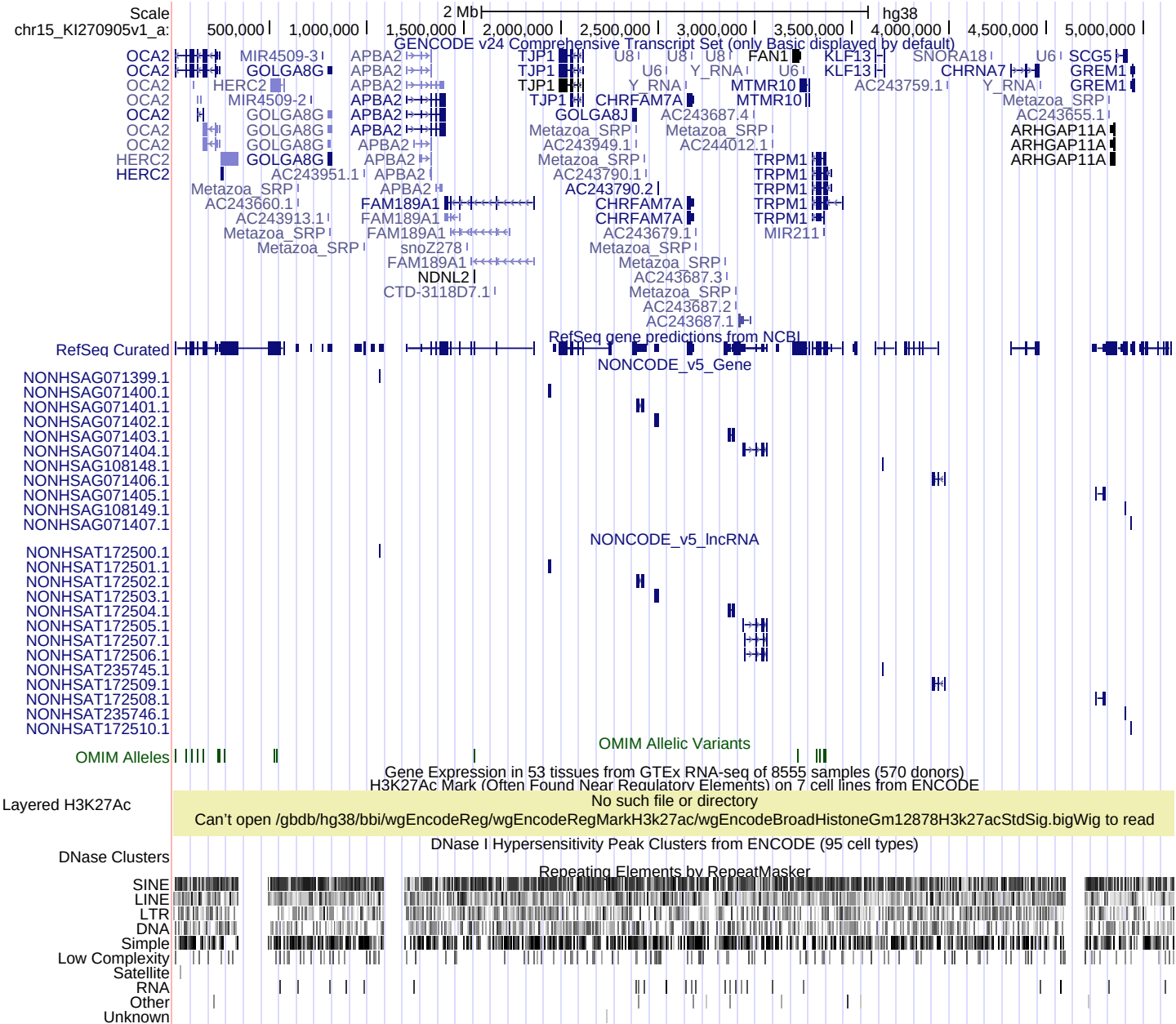
Scale
chr15_KI270905v1_a:
2 Mb
hg38
500,000
1,000,000
1,500,000
2,000,000
2,500,000
3,000,000
3,500,000
4,000,000
4,500,000
5,000,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
OCA2
HERC2
MIR4509-3
MIR4509-2
GOLGA8G
AC243951.1
Metazoa_SRP
AC243660.1
AC243913.1
APBA2
FAM189A1
snoZ278
NDNL2
CTD-3118D7.1
TJP1
CHRFAM7A
GOLGA8J
AC243687.4
AC243949.1
AC244012.1
AC243790.1
AC243790.2
AC243679.1
AC243687.3
AC243687.2
AC243687.1
U8
U6
Y_RNA
FAN1
KLF13
SNORA18
CHRNA7
SCG5
GREM1
MTMR10
AC243759.1
AC243655.1
ARHGAP11A
TRPM1
MIR211
RefSeq Curated
RefSeq gene predictions from NCBI
NONCODE_v5_Gene
NONHSAG071399.1
NONHSAG071400.1
NONHSAG071401.1
NONHSAG071402.1
NONHSAG071403.1
NONHSAG071404.1
NONHSAG108148.1
NONHSAG071406.1
NONHSAG071405.1
NONHSAG108149.1
NONHSAG071407.1
NONCODE_v5_lncRNA
NONHSAT172500.1
NONHSAT172501.1
NONHSAT172502.1
NONHSAT172503.1
NONHSAT172504.1
NONHSAT172505.1
NONHSAT172507.1
NONHSAT172506.1
NONHSAT235745.1
NONHSAT172509.1
NONHSAT172508.1
NONHSAT235746.1
NONHSAT172510.1
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