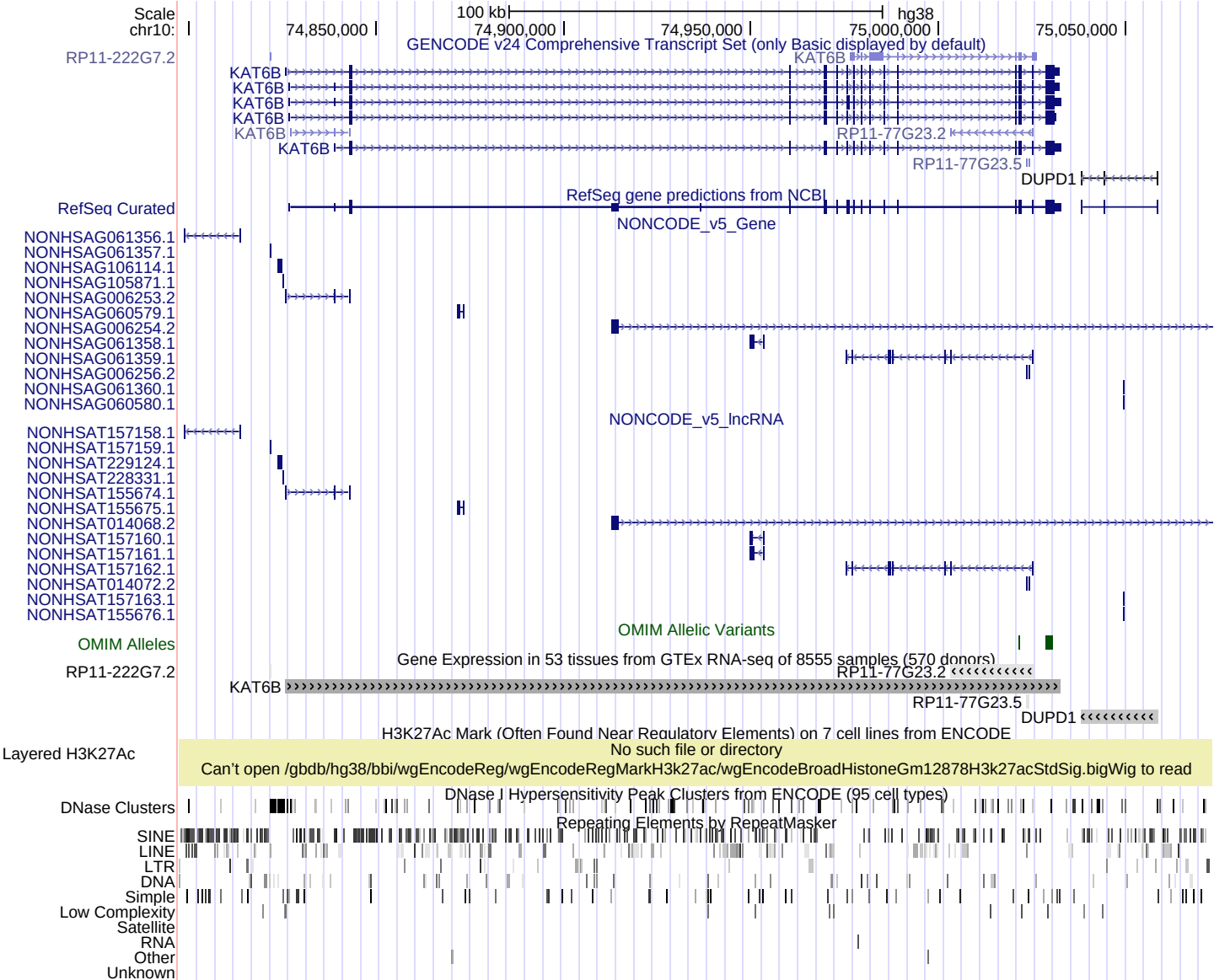
Scale
chr10:
100 kb
hg38
74,850,000
74,900,000
74,950,000
75,000,000
75,050,000
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
RP11-222G7.2
KAT6B
RP11-77G23.2
RP11-77G23.5
DUPD1
RefSeq Curated
RefSeq gene predictions from NCBI
NONCODE_v5_Gene
NONHSAG061356.1
NONHSAG061357.1
NONHSAG106114.1
NONHSAG105871.1
NONHSAG006253.2
NONHSAG060579.1
NONHSAG006254.2
NONHSAG061358.1
NONHSAG061359.1
NONHSAG006256.2
NONHSAG061360.1
NONHSAG060580.1
NONCODE_v5_lncRNA
NONHSAT157158.1
NONHSAT157159.1
NONHSAT229124.1
NONHSAT228331.1
NONHSAT155674.1
NONHSAT155675.1
NONHSAT014068.2
NONHSAT157160.1
NONHSAT157161.1
NONHSAT157162.1
NONHSAT014072.2
NONHSAT157163.1
NONHSAT155676.1
OMIM Allelic Variants
OMIM Alleles
Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
RP11-222G7.2
RP11-77G23.2
KAT6B
RP11-77G23.5
DUPD1
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