

Scale
chr16:

75,950,000 |

50 kb |

76,000,000 |

hg38

76,050,000 |

GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
RefSeq gene predictions from NCBI

RefSeq Curated

NONHSAG071833.1

NONCODE_v5_Gene



NONHSAT173289.1

NONCODE_v5_lncRNA



OMIM Alleles

OMIM Allelic Variants

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 /gdb/hg38/bbi/wgEncodeReg/wgEncodeRegMarkH3k27ac/wgEncodeBroadHistoneGm12878H3k27acStdSig.bigWig to read

DNase Clusters

DNase I Hypersensitivity Peak Clusters from ENCODE (95 cell types)

SINE

LINE

LTR

DNA

Simple

Low Complexity

Satellite

RNA

Other

Unknown

Repeating Elements by RepeatMasker

