

Scale
chr5: |

50 kb | hg38
21,150,000 | 21,200,000 |

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

RefSeq Curated

- NONHSAG040035.2
- NONHSAG040036.2
- NONHSAG091749.1
- NONHSAG091750.1
- NONHSAG091751.1
- NONHSAG091752.1

- NONHSAT202345.1
- NONHSAT100708.2
- NONHSAT203946.1
- NONHSAT100709.2
- NONHSAT205430.1
- NONHSAT205431.1
- NONHSAT205432.1
- NONHSAT205433.1

NONCODE_v5_Gene

NONCODE_v5_lncRNA

OMIM Alleles

OMIM Allelic Variants

Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
RP11-811J10.1

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 I Hypersensitivity Peak Clusters from ENCODE (95 cell types)

DNase Clusters

- SINE
- LINE
- LTR
- DNA
- Simple
- Low Complexity
- Satellite
- RNA
- Other
- Unknown

Repeating Elements by RepeatMasker

