

Scale
chrY:

56,830,000 | 56,840,000 | 56,850,000 | 56,860,000 | 56,870,000 | 56,880,000 |

20 kb | hg38
GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)
RefSeq gene predictions from NCBI

RefSeq Curated

NONHSAG056015.2
NONHSAG056016.2
NONHSAG056017.2
NONHSAG104753.1
NONHSAG104754.1
NONHSAG104755.1
NONHSAG104756.1

NONHSAT139816.2
NONHSAT139817.2
NONHSAT139818.2
NONHSAT224526.1
NONHSAT224527.1
NONHSAT224528.1
NONHSAT224529.1

OMIM Alleles

OMIM Allelic Variants

Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
CTBP2P1 >

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

Repeating Elements by RepeatMasker

SINE

LINE

LTR

DNA

Simple

Low Complexity

Satellite

RNA

Other

Unknown