Scale
chr18:

100 kb   hg38

48,600,000 | 48,650,000 | 48,700,000 | 48,750,000 | 48,800,000 | 48,850,000 | 48,900,000 |

GENCODE v24 Comprehensive Transcript Set (only Basic displayed by default)

CTIF
CTIF
CTIF   CTIF   RP11-484L8.1   CTIF
CTIF   RP11-46D1.2   CTIF
CTIF
CTIF   CTIF
CTIF
RP11-426J5.3   CTIF   RP11-426J5.1
                                    AC048380.1
                                    RP11-426J5.2

RefSeq gene predictions from NCBI
RefSeq Curated

NONCODE_v5_Gene
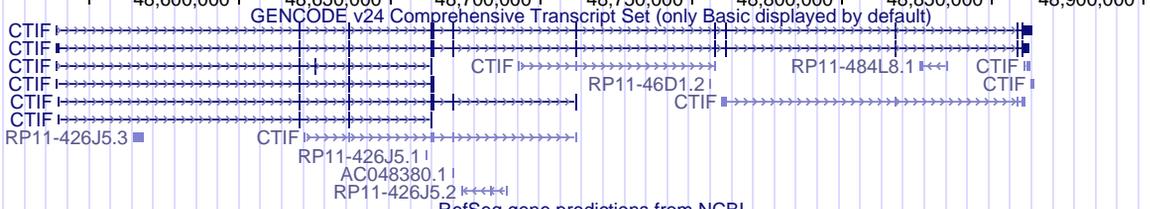NONHSAG109190.1
NONHSAG074790.1
NONHSAG075181.1
NONHSAG023816.2
NONHSAG109083.1
NONHSAG023818.2
NONHSAG023819.2
NONHSAG023820.2
NONHSAG075182.1
NONHSAG023821.3
NONHSAG074791.2
NONHSAG075678.1
NONHSAG075679.1
NONHSAG075680.1
NONHSAG075681.1

NONCODE_v5_lncRNA
NONHSAT238885.1
NONHSAT238884.1
NONHSAT177784.1
NONHSAT178468.1
NONHSAT059191.2
NONHSAT238509.1
NONHSAT059193.2
NONHSAT059196.2
NONHSAT059198.2
NONHSAT059197.2
NONHSAT178469.1
NONHSAT059200.2
NONHSAT178471.1
NONHSAT178470.1
NONHSAT238886.1
NONHSAT059201.2
NONHSAT238887.1
NONHSAT238510.1
NONHSAT177785.1
NONHSAT238511.1
NONHSAT179085.1
NONHSAT179086.1
NONHSAT179087.1
NONHSAT179088.1

OMIM Allelic Variants
OMIM Alleles

Gene Expression in 53 tissues from GTEx RNA-seq of 8555 samples (570 donors)
CTIF
                              AC048380.1
                    RP11-426J5.2   RP11-484L8.1
H3K27Ac Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE
No such file or directory
Layered H3K27Ac
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