

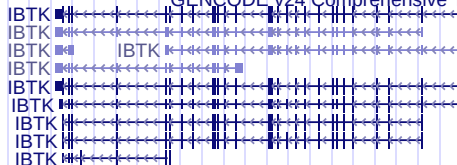
Scale
chr6:

100 kb|

hg38

82,200,000 | 82,250,000 | 82,300,000 | 82,350,000 |

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



TPBG
TPBG
TPBG

RefSeq Curated

NONHSAG044231.2
NONHSAG094306.1
NONHSAG112973.1
NONHSAG044232.2
NONHSAG044233.2
NONHSAG113298.1
NONHSAG044234.2

NONHSAT209364.1
NONHSAT209365.1
NONHSAT251271.1
NONHSAT113761.2
NONHSAT113762.2
NONHSAT207550.1
NONHSAT252466.1
NONHSAT252467.1
NONHSAT113763.2

OMIM Alleles

RefSeq gene predictions from NCBI

NONCODE_v5_Gene

NONCODE_v5_lncRNA

OMIM Allelic Variants

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

IBTK

RNU6-130P RP3-492P14.2

TPBG

H3K27Ac Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE

No such file or directory

Layered H3K27Ac

Can't open /gdb/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