

Scale 200 bases hg38  
chr19: 13,618,800 13,618,900 13,619,000 13,619,100 13,619,200

ChOP-seq universal TTSs

ChOP-seq

Background universal TTSs

Background

Capture-seq universal TTSs

Capture-seq

ENCODE v29 Comprehensive Transcript Set (only Basic displayed by default)

CACNA1A

Repeating Elements by RepeatMasker

SINE

LINE

LTR

DNA

Simple

Low Complexity

Satellite

RNA

Other

Unknown