

Scale

5 kb

GCA\_036417535.1

chr1:

71,899,000

71,900,000

71,901,000

71,902,000

71,903,000

71,904,000

71,905,000

71,906,000

71,907,000

GC Percent in 5-Base Windows

GC Percent

Augustus Gene Predictions

RefSeq mRNAs mapped to this assembly

CpG Islands (Islands < 300 Bases are Light Green)

Unmasked CpG

RepeatMasker Repetitive Elements

(T)n

DR0008641

>>

(T)n

(T)n >

Simple Tandem Repeats by TRF

Simple Repeats

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust