

Scale

5 kb

GCA\_964195595.1

chr1:

73,704,000

73,705,000

73,706,000

73,707,000

73,708,000

73,709,000

73,710,000

73,711,000

73,712,000

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

(TG)n

(TATGTA)n

(T)n

(T)n

Simple Tandem Repeats by TRF

Simple Repeats

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

