

Scale

5 kb

GCA_964106895.1

chr1:

47,328,000

47,329,000

47,330,000

47,331,000

47,332,000

47,333,000

47,334,000

47,335,000

47,336,000

47,337,000

GC Percent in 5-Base Windows

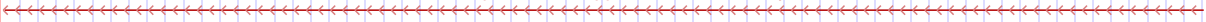
GC Percent



Augustus Gene Predictions

RefSeq mRNAs mapped to this assembly

NM_001130249.2



CpG Islands (Islands < 300 Bases are Light Green)

Unmasked CpG



RepeatMasker Repetitive Elements

(A)n

Simple Tandem Repeats by TRF

Simple Repeats



Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

