

Scale

5 kb

GCA_031885435.1

chr1:

52,374,000

52,375,000

52,376,000

52,377,000

52,378,000

52,379,000

52,380,000

52,381,000

52,382,000

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

GA-rich



Simple Tandem Repeats by TRF

Simple Repeats

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

