

5 kb

GCA_017310615.1

Scale

chr1:

| 3,120,500 | 3,121,000 | 3,121,500 | 3,122,000 | 3,122,500 | 3,123,000 | 3,123,500 | 3,124,000 | 3,124,500 | 3,125,000 | 3,125,500 | 3,126,000 | 3,126,500 | 3,127,000 | 3,127,500 | 3,128,000 | 3,128,500 | 3,129,000 | 3,129,500 |

GC Percent in 5-Base Windows

GC Percent



Augustus Gene Predictions

Augustus



Gene models submitted to NCBI

RML2



TRM11



I7I53_09587



RefSeq mRNAs mapped to this assembly

NM_208023.1



NM_001184216.1



CpG Islands (Islands < 300 Bases are Light Green)

CpG: 125



CpG: 39



CpG Islands (Islands < 300 Bases are Light Green)

Unmasked CpG



RepeatMasker Repetitive Elements

(CACAT)n



(GAA)n



(CTTT)n



(CT)n



rnd-3_family-493



(T)n



Simple Repeats

Simple Tandem Repeats by TRF

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

