

Scale

5 kb

GCA_030936125.1

CM061645.1: 44,475,000| 44,476,000| 44,477,000| 44,478,000| 44,479,000| 44,480,000| 44,481,000| 44,482,000| 44,483,000| 44,484,000|

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

UCON26

(TTTTC)_n

A-rich

A-rich

UCON18

DR0079598

>>

DR0079568

>>

A-rich

DR0079496

<<<

DR0079568

<<

DR0079559

<<<<

Simple Tandem Repeats by TRF

Simple Repeats

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

