

Scale 5 kb | GCA_014706295.1

JACSVZ010000002.1: | 50,738,000 | 50,739,000 | 50,740,000 | 50,741,000 | 50,742,000 | 50,743,000 | 50,744,000 | 50,745,000 | 50,746,000 |

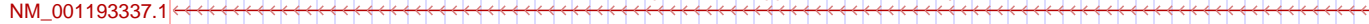
GC Percent in 5-Base Windows



Augustus Gene Predictions



RefSeq mRNAs mapped to this assembly



CpG Islands (Islands < 300 Bases are Light Green)

Unmasked CpG

RepeatMasker Repetitive Elements

TguLTR5d



CR1-Y2_Aves >>



UCON12A <



X7C_LINE



(TAA)n



CR1-Y2_Aves <



(TTAAG)n



Simple Tandem Repeats by TRF

Simple Repeats

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

