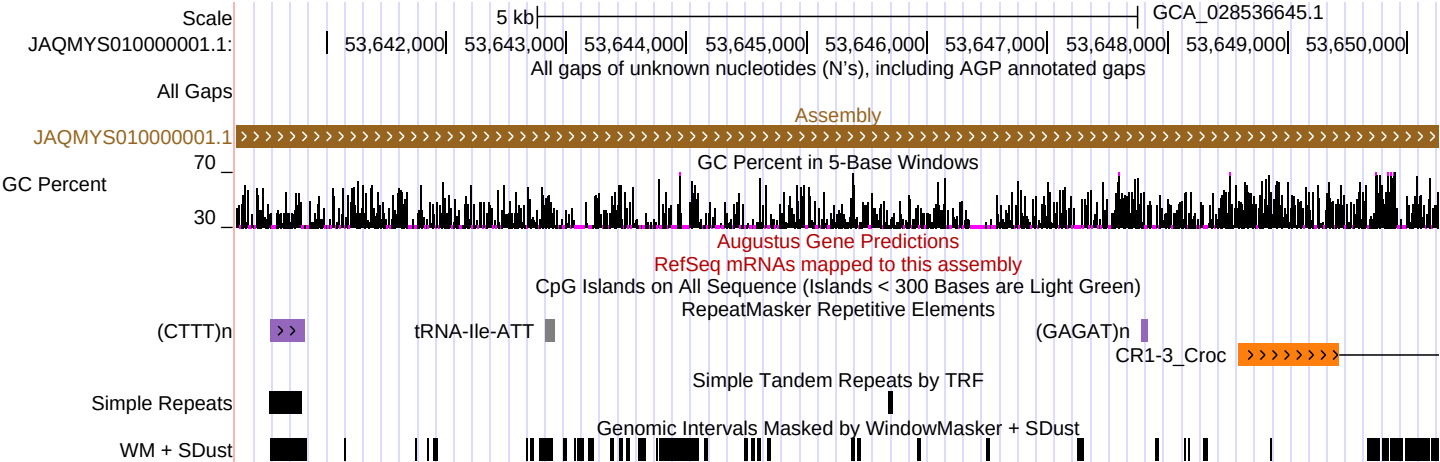
Scale
5 kb
GCA_028536645.1
JAQMYS010000001.1:
53,642,000
53,643,000
53,644,000
53,645,000
53,646,000
53,647,000
53,648,000
53,649,000
53,650,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAQMYS010000001.1
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(CTTT)n
tRNA-Ile-ATT
(GAGAT)n
CR1-3_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust