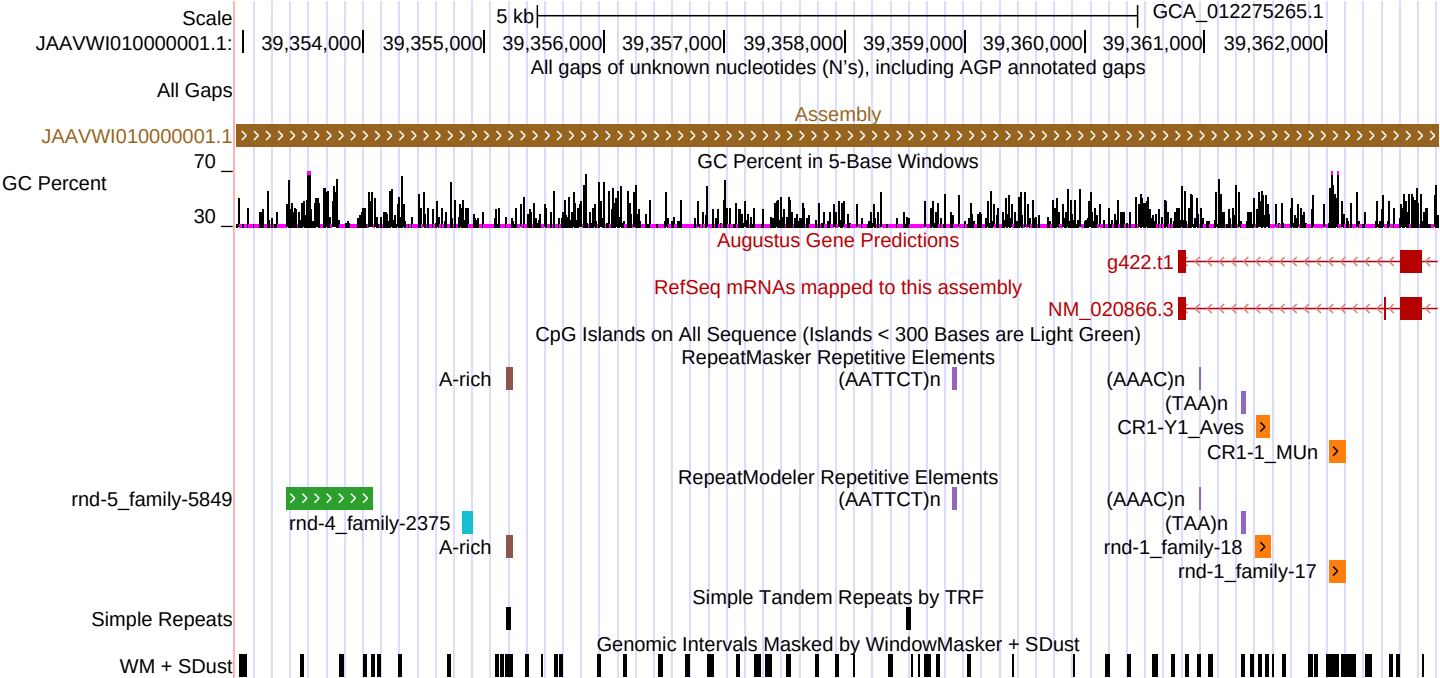
Scale
5 kb
GCA_012275265.1
JAAVWI010000001.1:
39,354,000
39,355,000
39,356,000
39,357,000
39,358,000
39,359,000
39,360,000
39,361,000
39,362,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAAVWI010000001.1
70
GC Percent in 5-Base Windows
GC Percent
30
Augustus Gene Predictions
g422.t1
RefSeq mRNAs mapped to this assembly
NM_020866.3
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
A-rich
(AATTCT)n
(AAAC)n
(TAA)n
CR1-Y1_Aves
CR1-1_MUn
RepeatModeler Repetitive Elements
rnd-5_family-5849
(AATTCT)n
(AAAC)n
rnd-4_family-2375
(TAA)n
A-rich
rnd-1_family-18
rnd-1_family-17
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust