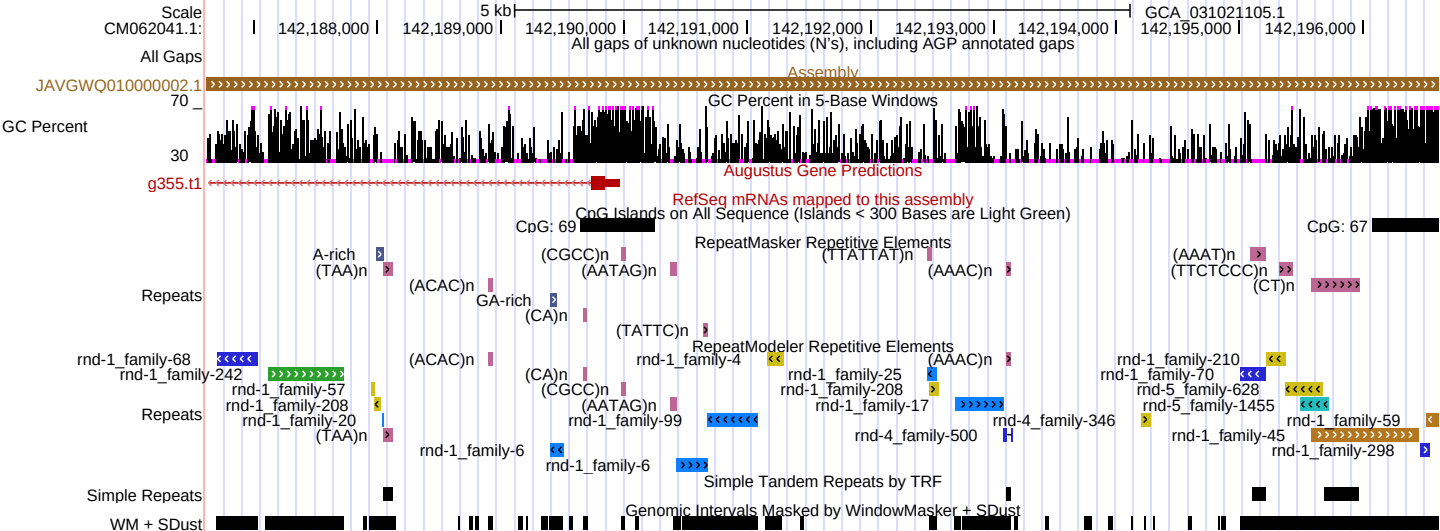
Scale
5 kb
GCA_031021105.1
CM062041.1:
142,188,000
142,189,000
142,190,000
142,191,000
142,192,000
142,193,000
142,194,000
142,195,000
142,196,000
All Gaps
All gaps of unknown nucleotides (N's), including AGP annotated gaps
Assembly
JAVGWQ010000002.1
GC Percent
70
30
GC Percent in 5-Base Windows
Augustus Gene Predictions
g355.t1
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
CpG: 69
CpG: 67
RepeatMasker Repetitive Elements
Repeats
A-rich
(TAA)n
(CGCC)n
(AATAG)n
(TTATTAT)n
(AAAC)n
(AAAT)n
(TTCTCCC)n
(CT)n
(ACAC)n
GA-rich
(CA)n
(TATTC)n
RepeatModeler Repetitive Elements
Repeats
rnd-1_family-68
rnd-1_family-242
rnd-1_family-57
rnd-1_family-208
rnd-1_family-20
(TAA)n
(ACAC)n
rnd-1_family-4
(CA)n
(CGCC)n
(AATAG)n
rnd-1_family-99
rnd-1_family-6
rnd-1_family-6
rnd-1_family-25
rnd-1_family-208
rnd-1_family-17
rnd-4_family-500
(AAAC)n
rnd-1_family-210
rnd-1_family-70
rnd-5_family-628
rnd-5_family-1455
rnd-4_family-346
rnd-1_family-45
rnd-1_family-59
rnd-1_family-298
Simple Repeats
Simple Tandem Repeats by TRF
WM + SDust
Genomic Intervals Masked by WindowMasker + SDust