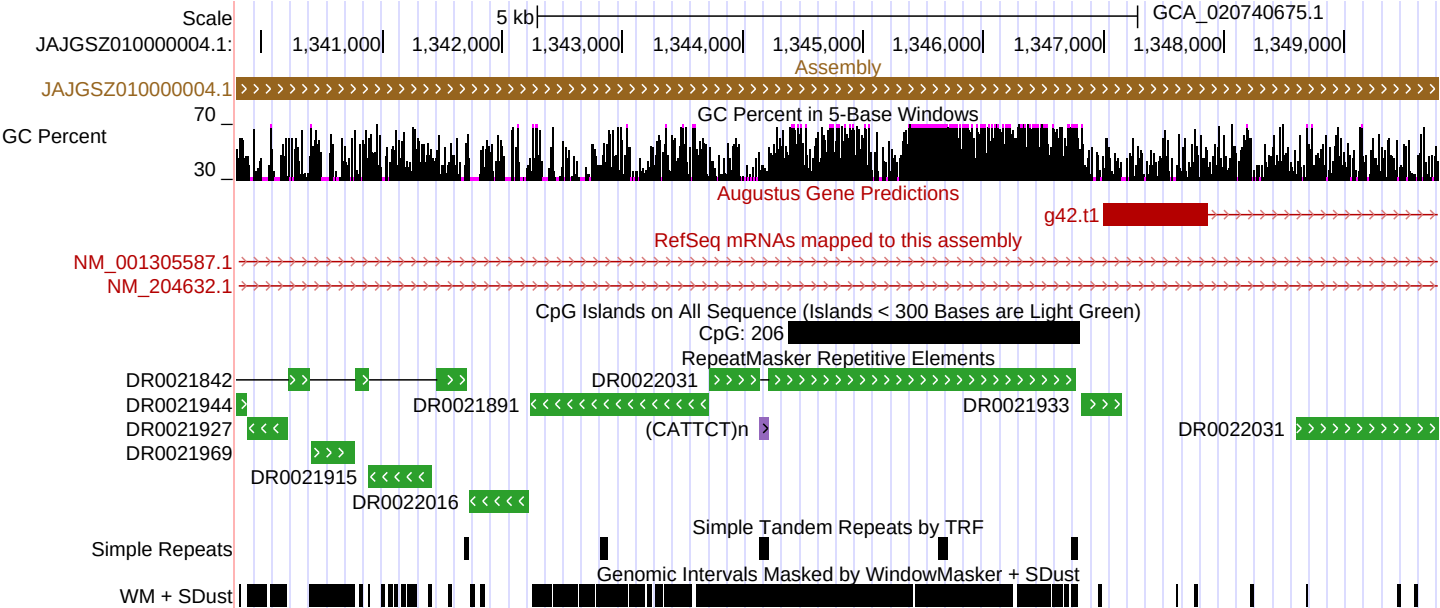
Scale
5 kb
GCA_020740675.1
JAJGSZ010000004.1:
1,341,000
1,342,000
1,343,000
1,344,000
1,345,000
1,346,000
1,347,000
1,348,000
1,349,000
Assembly
JAJGSZ010000004.1
GC Percent
70
30
GC Percent in 5-Base Windows
Augustus Gene Predictions
g42.t1
RefSeq mRNAs mapped to this assembly
NM_001305587.1
NM_204632.1
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
CpG: 206
RepeatMasker Repetitive Elements
DR0021842
DR0022031
DR0021944
DR0021891
DR0021933
DR0021927
(CATTCT)n
DR0022031
DR0021969
DR0021915
DR0022016
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust