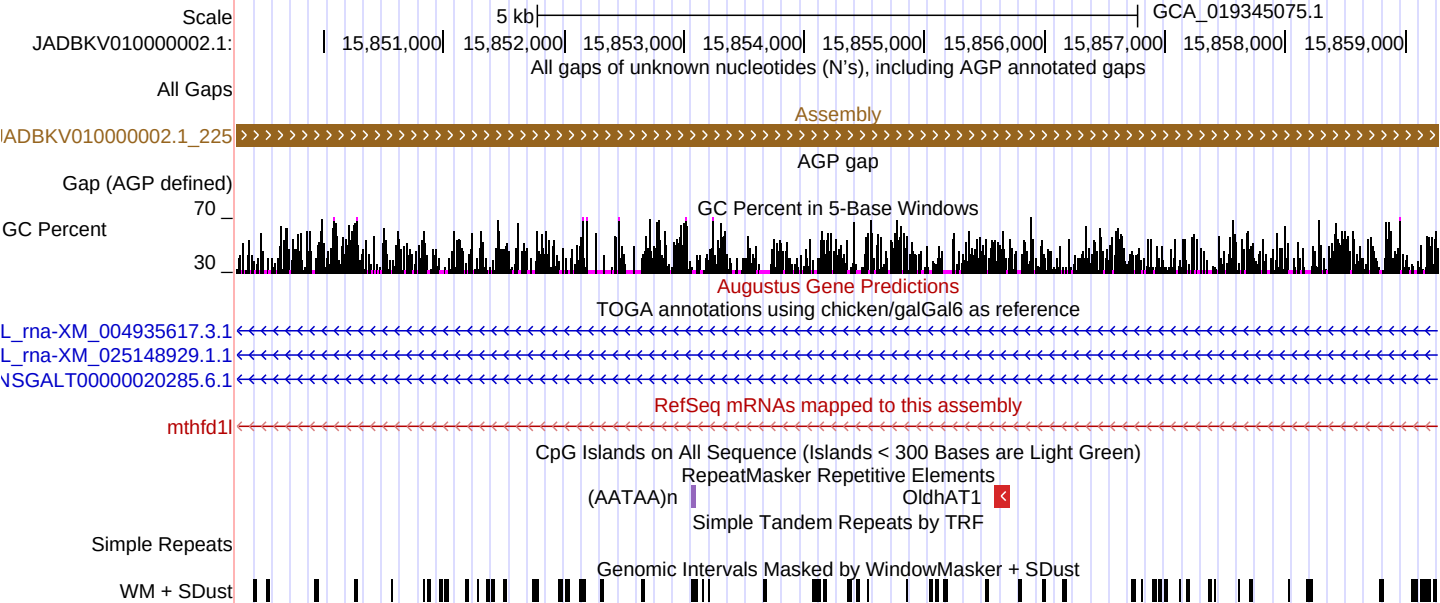
Scale
5 kb
GCA_019345075.1
JADBKV010000002.1:
15,851,000
15,852,000
15,853,000
15,854,000
15,855,000
15,856,000
15,857,000
15,858,000
15,859,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
ADBKV010000002.1_225
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
TOGA annotations using chicken/galGal6 as reference
L_rna-XM_004935617.3.1
L_rna-XM_025148929.1.1
NSGALT00000020285.6.1
RefSeq mRNAs mapped to this assembly
mthfd1l
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(AATAA)n
OldhAT1
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust