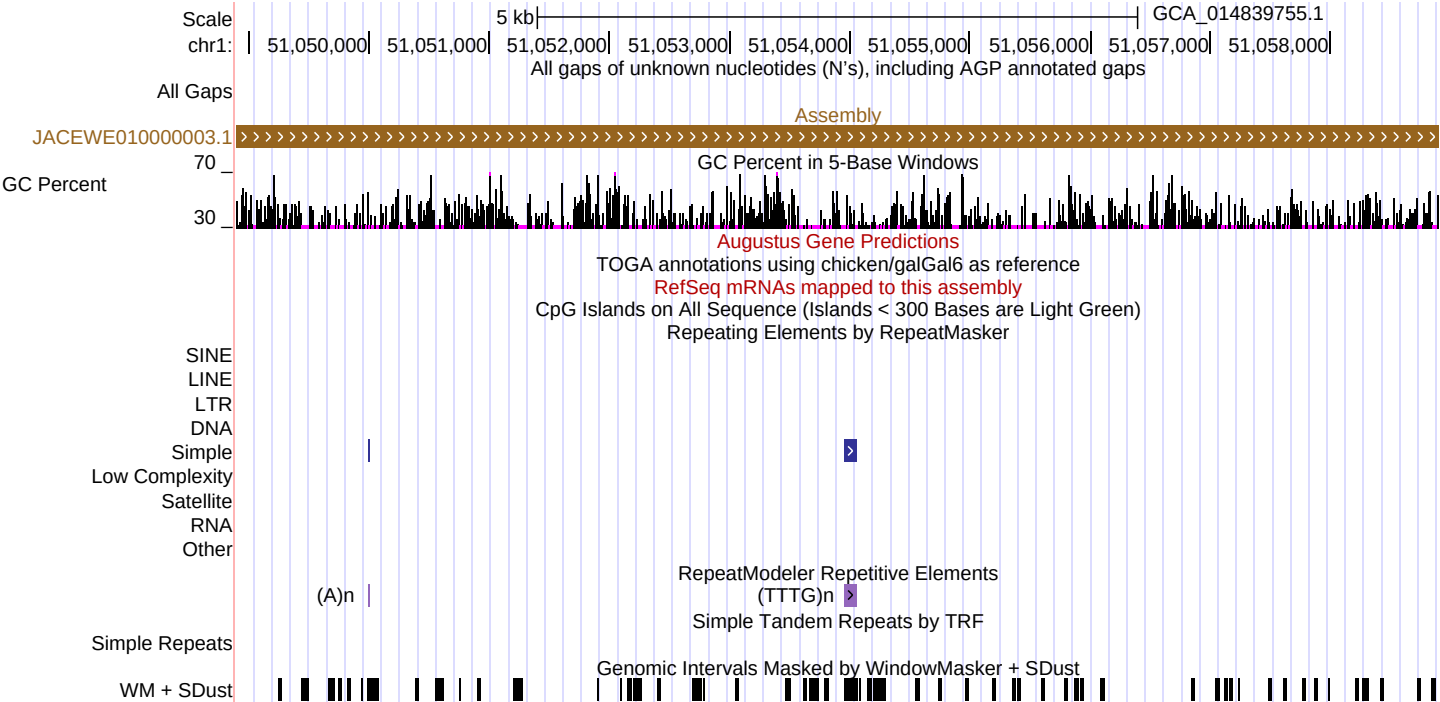
Scale
5 kb
GCA_014839755.1
chr1:
51,050,000
51,051,000
51,052,000
51,053,000
51,054,000
51,055,000
51,056,000
51,057,000
51,058,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JACEWE010000003.1
70
GC Percent in 5-Base Windows
GC Percent
30
Augustus Gene Predictions
TOGA annotations using chicken/galGal6 as reference
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
Repeating Elements by RepeatMasker
SINE
LINE
LTR
DNA
Simple
Low Complexity
Satellite
RNA
Other
RepeatModeler Repetitive Elements
(A)n
(TTTG)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust