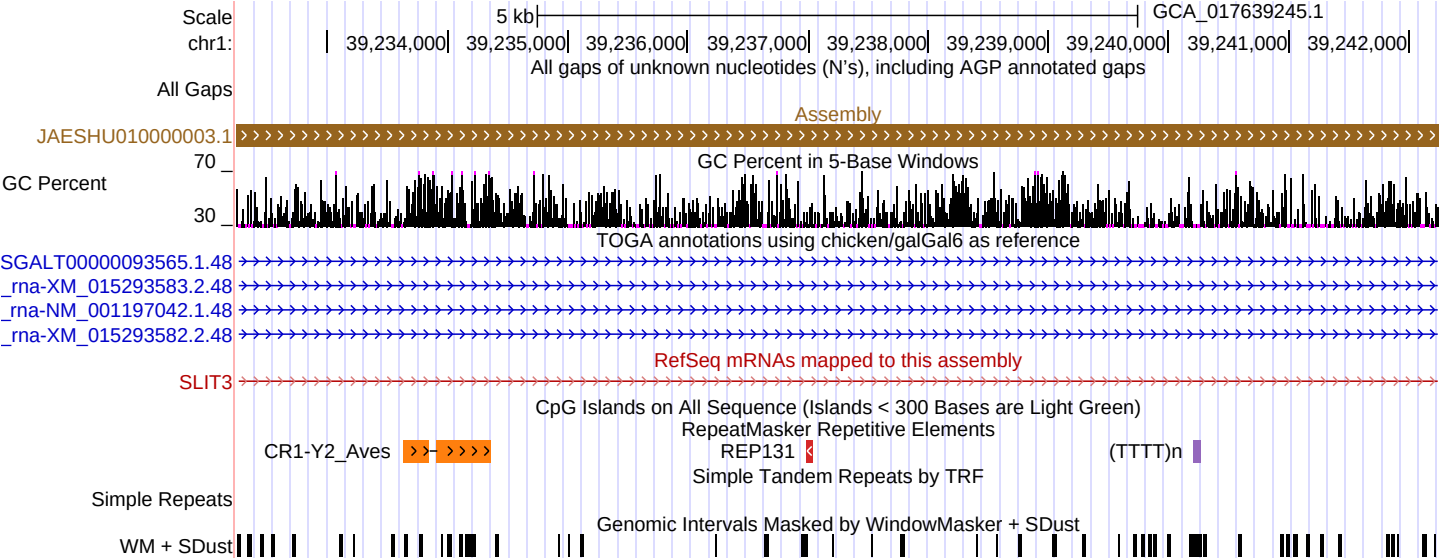
Scale
5 kb
GCA_017639245.1
chr1:
39,234,000
39,235,000
39,236,000
39,237,000
39,238,000
39,239,000
39,240,000
39,241,000
39,242,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAESHU010000003.1
GC Percent in 5-Base Windows
70
GC Percent
30
TOGA annotations using chicken/galGal6 as reference
SGALT00000093565.1.48
_rna-XM_015293583.2.48
_rna-NM_001197042.1.48
_rna-XM_015293582.2.48
RefSeq mRNAs mapped to this assembly
SLIT3
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
CR1-Y2_Aves
REP131
(TTTT)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust