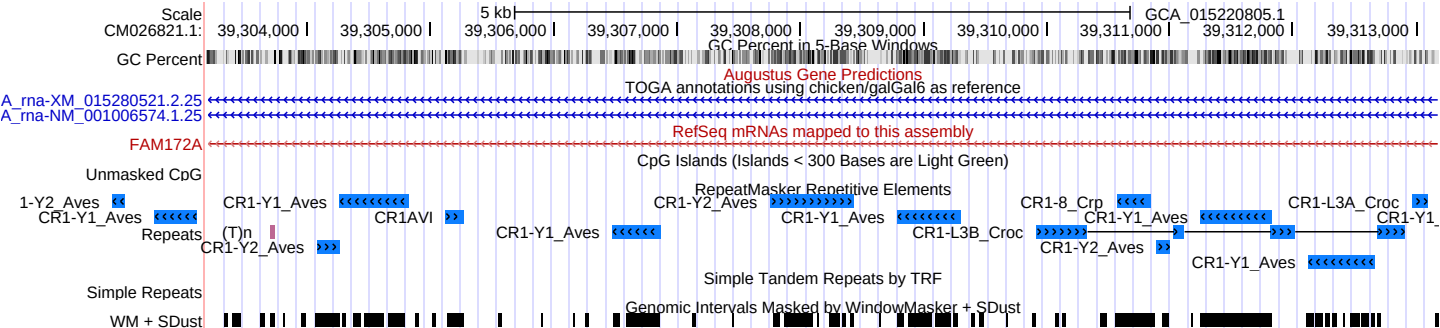
Scale
5 kb
GCA_015220805.1
CM026821.1:
39,304,000
39,305,000
39,306,000
39,307,000
39,308,000
39,309,000
39,310,000
39,311,000
39,312,000
39,313,000
GC Percent in 5-Base Windows
GC Percent
Augustus Gene Predictions
TOGA annotations using chicken/galGal6 as reference
A_rna-XM_015280521.2.25
A_rna-NM_001006574.1.25
RefSeq mRNAs mapped to this assembly
FAM172A
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
1-Y2_Aves
CR1-Y1_Aves
CR1-Y1_Aves
CR1AVI
CR1-Y2_Aves
CR1-Y1_Aves
CR1-8_Crp
CR1-Y1_Aves
CR1-L3A_Croc
CR1-Y1
Repeats
(T)n
CR1-Y1_Aves
CR1-L3B_Croc
CR1-Y2_Aves
CR1-Y2_Aves
CR1-Y1_Aves
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust