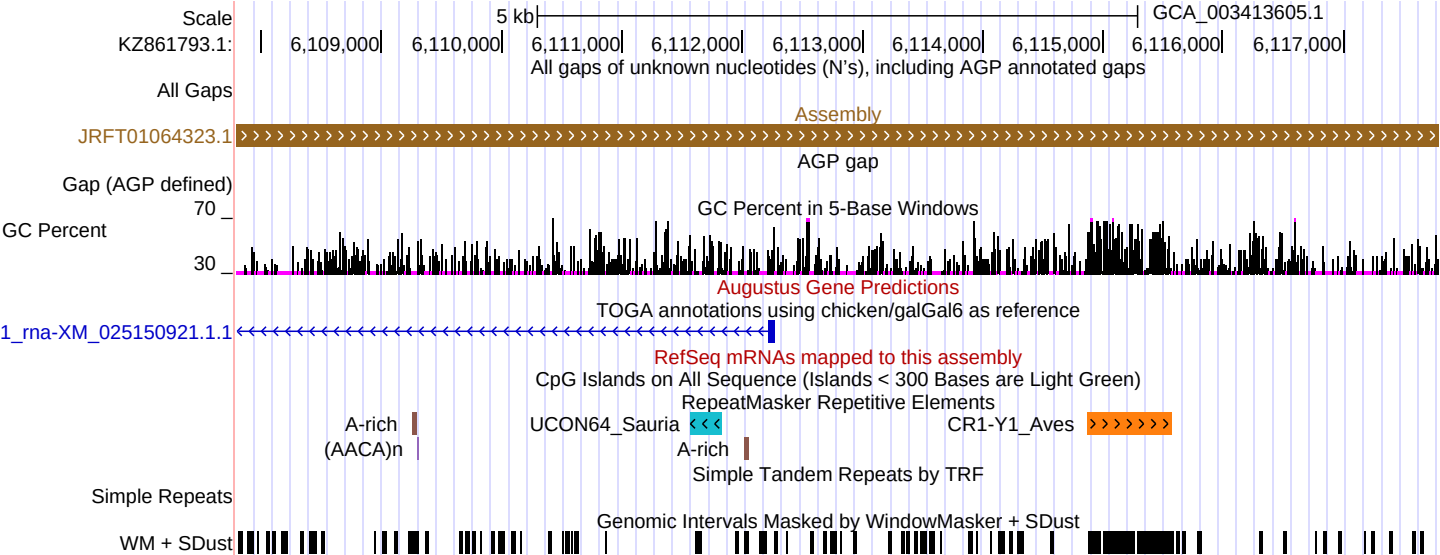
Scale
5 kb
GCA_003413605.1
KZ861793.1:
6,109,000
6,110,000
6,111,000
6,112,000
6,113,000
6,114,000
6,115,000
6,116,000
6,117,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JRFT01064323.1
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
TOGA annotations using chicken/galGal6 as reference
1_rna-XM_025150921.1.1
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
A-rich
UCON64_Sauria
CR1-Y1_Aves
(AACA)n
A-rich
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust