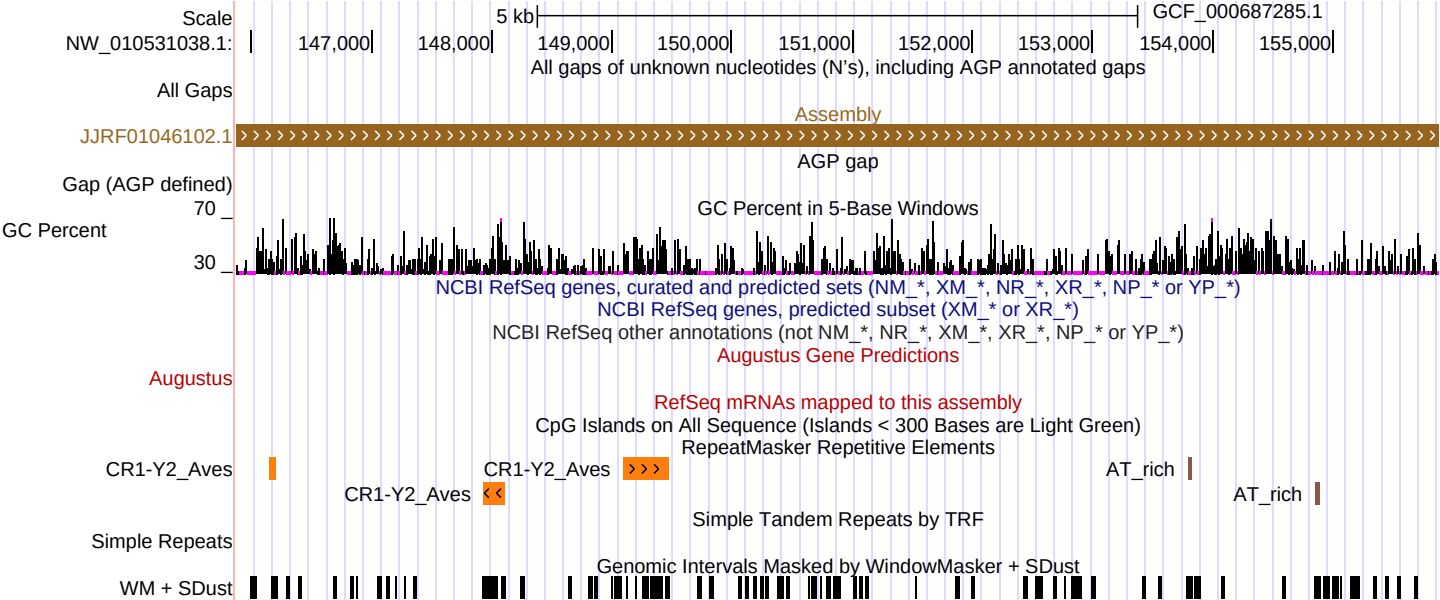
Scale
5 kb
GCF_000687285.1
NW_010531038.1:
147,000
148,000
149,000
150,000
151,000
152,000
153,000
154,000
155,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JJRF01046102.1
AGP gap
Gap (AGP defined)
70
GC Percent
30
GC Percent in 5-Base Windows
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
CR1-Y2_Aves
CR1-Y2_Aves
CR1-Y2_Aves
AT_rich
AT_rich
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust