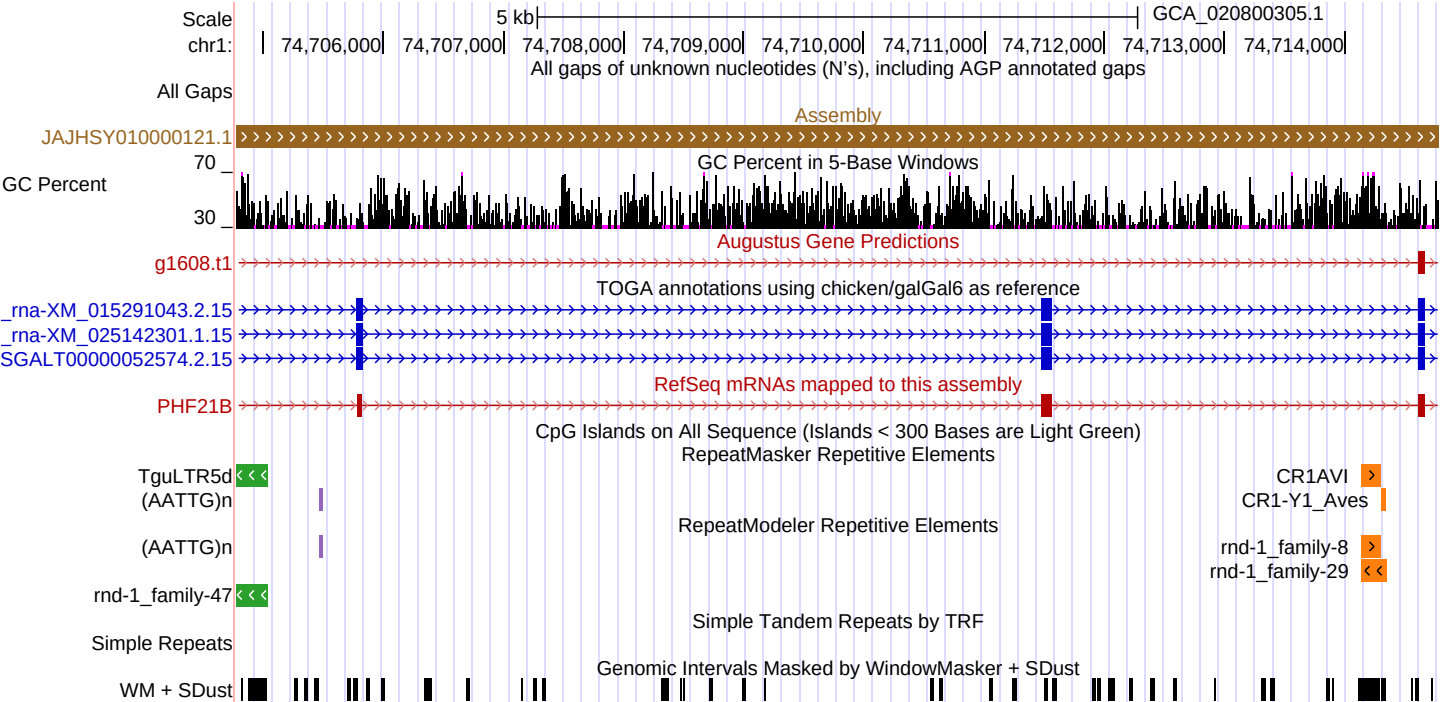
Scale
5 kb
GCA_020800305.1
chr1:
74,706,000
74,707,000
74,708,000
74,709,000
74,710,000
74,711,000
74,712,000
74,713,000
74,714,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAJHSY010000121.1
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
g1608.t1
TOGA annotations using chicken/galGal6 as reference
_rna-XM_015291043.2.15
_rna-XM_025142301.1.15
SGALT00000052574.2.15
RefSeq mRNAs mapped to this assembly
PHF21B
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
TguLTR5d
(AATTG)n
CR1AVI
CR1-Y1_Aves
RepeatModeler Repetitive Elements
(AATTG)n
rnd-1_family-8
rnd-1_family-29
rnd-1_family-47
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust