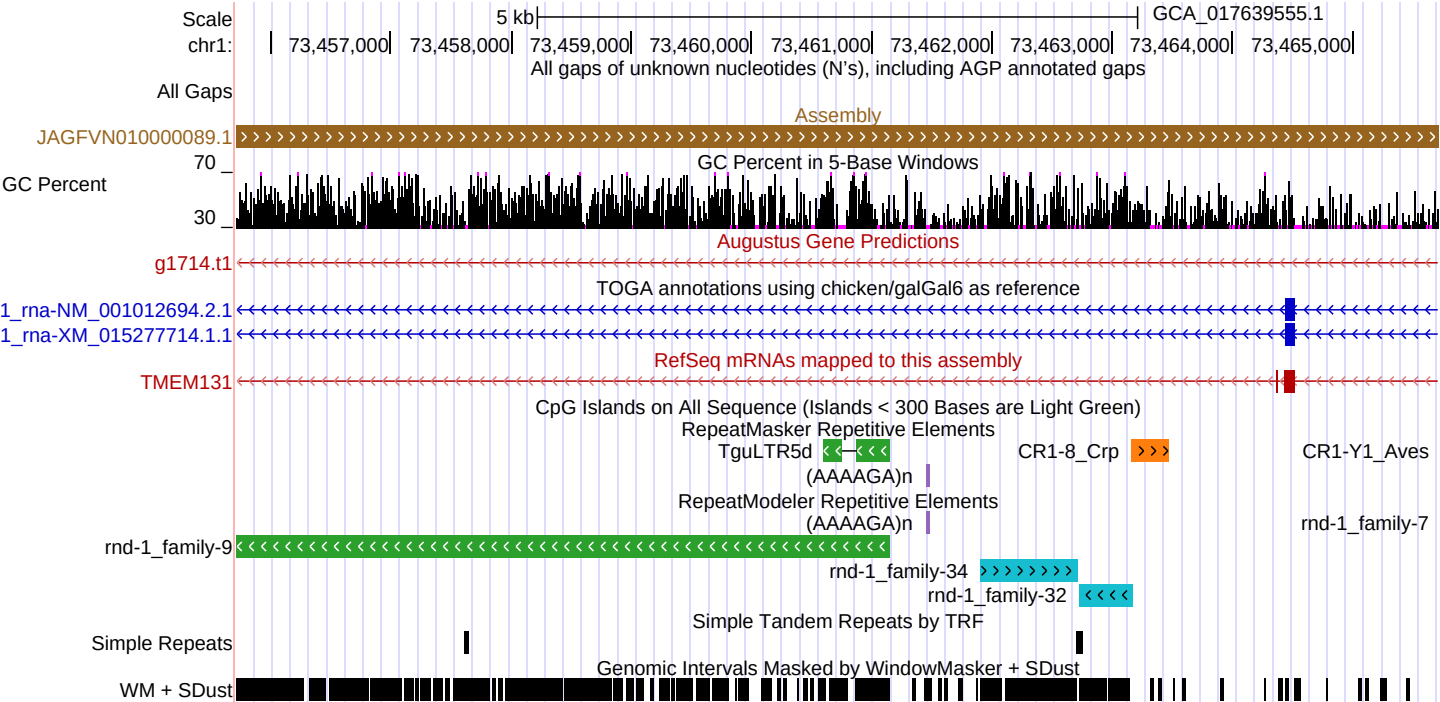
Scale
5 kb
GCA_017639555.1
chr1:
73,457,000
73,458,000
73,459,000
73,460,000
73,461,000
73,462,000
73,463,000
73,464,000
73,465,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAGFVN010000089.1
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
g1714.t1
TOGA annotations using chicken/galGal6 as reference
1_rna-NM_001012694.2.1
1_rna-XM_015277714.1.1
RefSeq mRNAs mapped to this assembly
TMEM131
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
TguLTR5d
CR1-8_Crp
CR1-Y1_Aves
(AAAAGA)n
RepeatModeler Repetitive Elements
(AAAAGA)n
rnd-1_family-7
rnd-1_family-9
rnd-1_family-34
rnd-1_family-32
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust