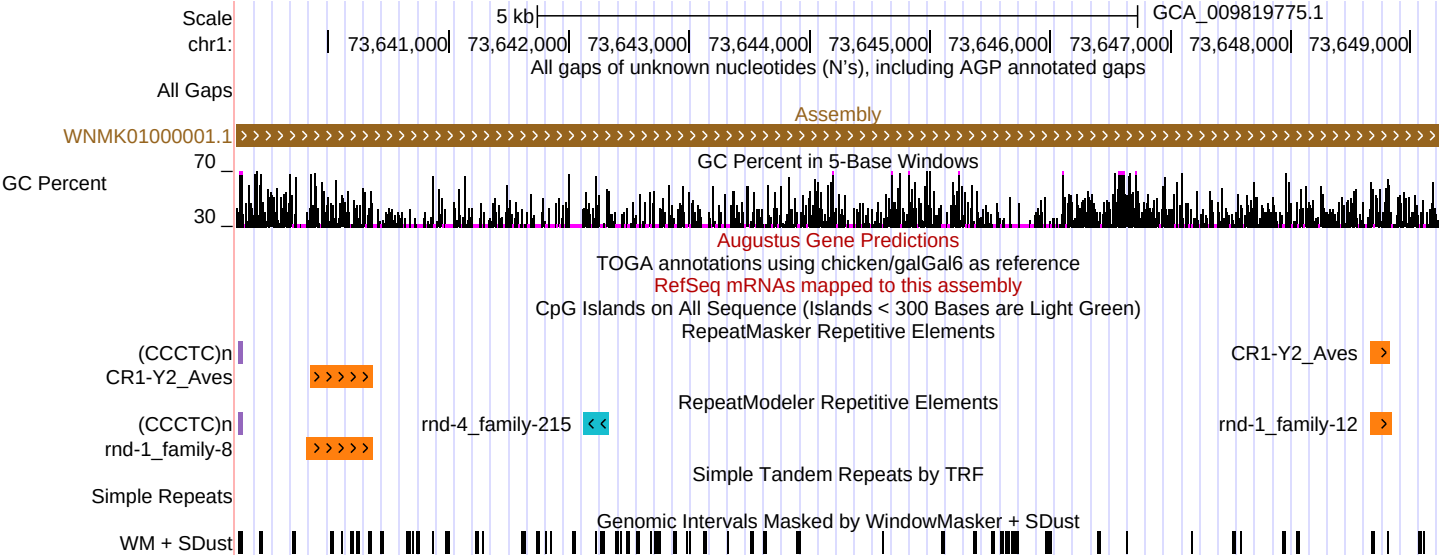
Scale
5 kb
GCA_009819775.1
chr1:
73,641,000
73,642,000
73,643,000
73,644,000
73,645,000
73,646,000
73,647,000
73,648,000
73,649,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
WNMK01000001.1
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
TOGA annotations using chicken/galGal6 as reference
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(CCCTC)n
CR1-Y2_Aves
CR1-Y2_Aves
RepeatModeler Repetitive Elements
(CCCTC)n
rnd-4_family-215
rnd-1_family-12
rnd-1_family-8
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust