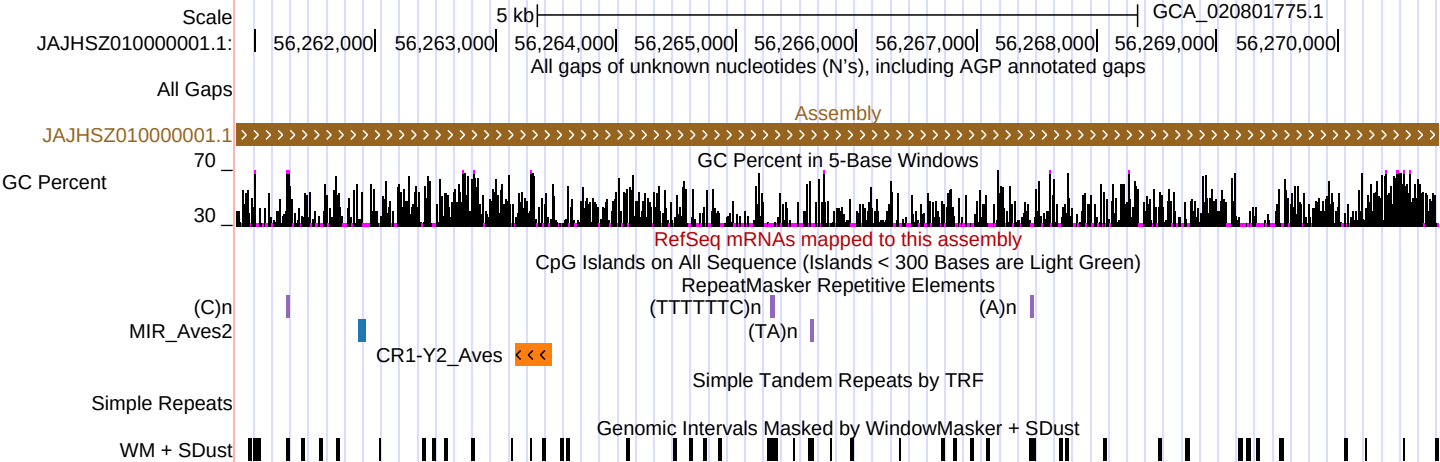
Scale
5 kb
GCA_020801775.1
JAJHSZ010000001.1:
56,262,000
56,263,000
56,264,000
56,265,000
56,266,000
56,267,000
56,268,000
56,269,000
56,270,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAJHSZ010000001.1
GC Percent in 5-Base Windows
70
GC Percent
30
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(C)n
(TTTTTTC)n
(A)n
MIR_Aves2
(TA)n
CR1-Y2_Aves
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust