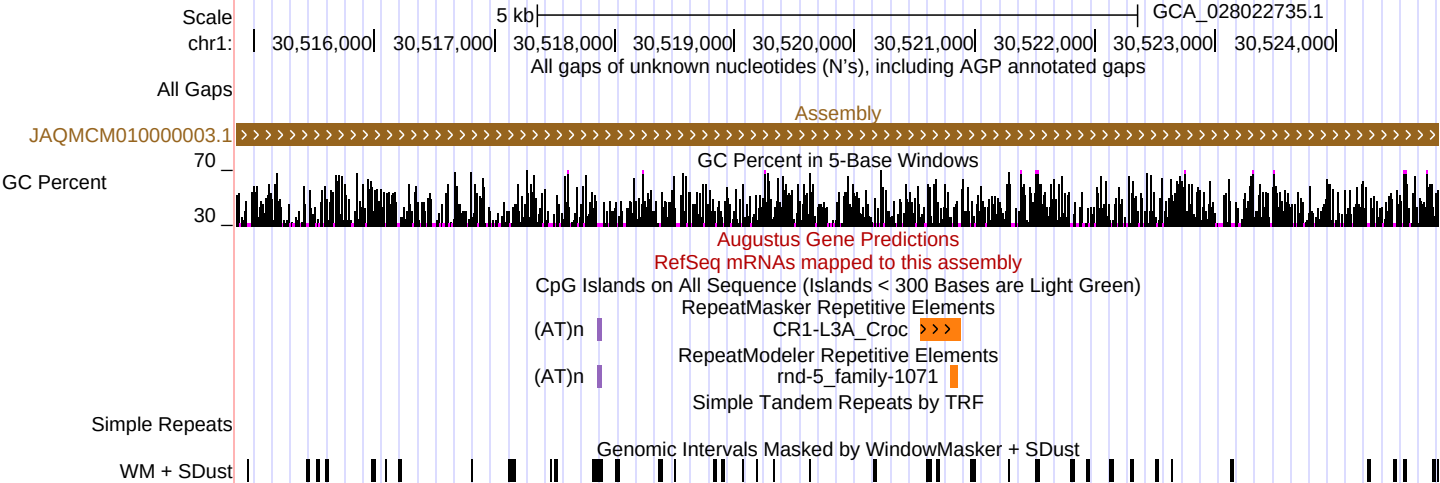
Scale
5 kb
GCA_028022735.1
chr1:
30,516,000
30,517,000
30,518,000
30,519,000
30,520,000
30,521,000
30,522,000
30,523,000
30,524,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAQMCM010000003.1
70
GC Percent
30
GC Percent in 5-Base Windows
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(AT)n
CR1-L3A_Croc
RepeatModeler Repetitive Elements
(AT)n
rnd-5_family-1071
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust