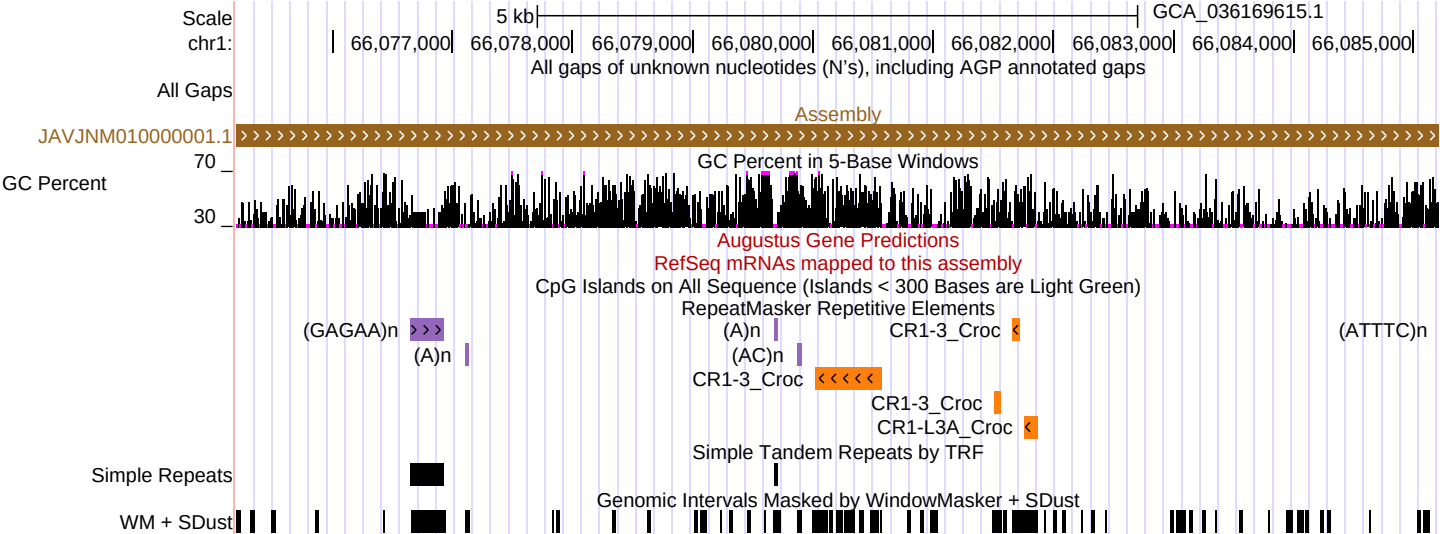
Scale
5 kb
GCA_036169615.1
chr1:
66,077,000
66,078,000
66,079,000
66,080,000
66,081,000
66,082,000
66,083,000
66,084,000
66,085,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAVJNM010000001.1
70
GC Percent in 5-Base Windows
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(GAGAA)n
(A)n
CR1-3_Croc
(ATTTC)n
(A)n
(AC)n
CR1-3_Croc
CR1-3_Croc
CR1-L3A_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust