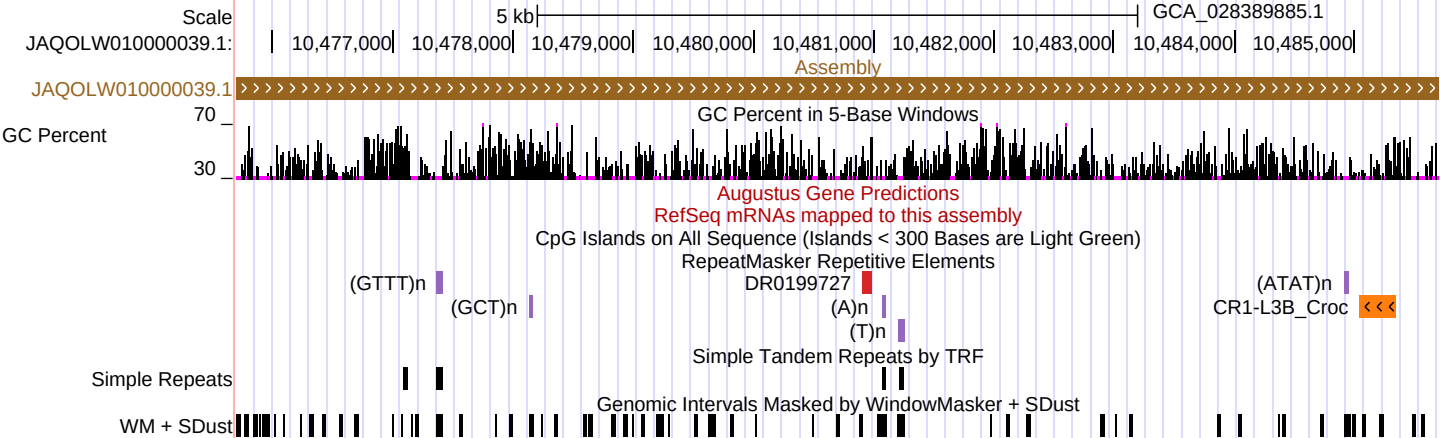
Scale
5 kb
GCA_028389885.1
JAQOLW010000039.1:
10,477,000
10,478,000
10,479,000
10,480,000
10,481,000
10,482,000
10,483,000
10,484,000
10,485,000
Assembly
JAQOLW010000039.1
70
GC Percent
30
GC Percent in 5-Base Windows
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(GTTT)n
DR0199727
(ATAT)n
(GCT)n
(A)n
CR1-L3B_Croc
(T)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust