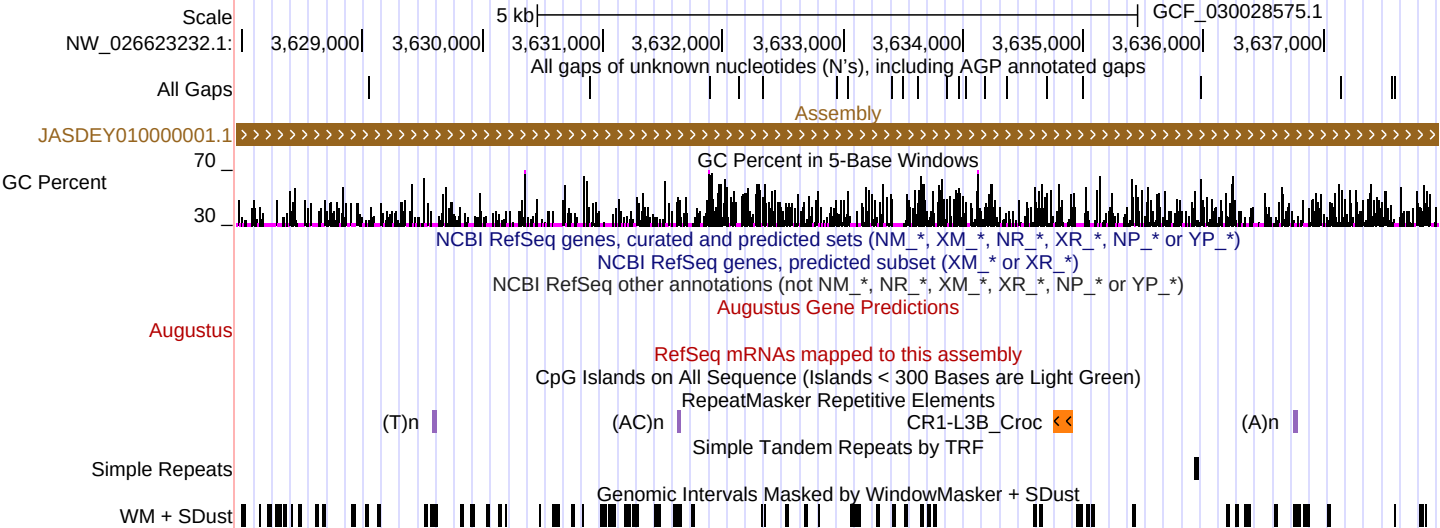
Scale
5 kb
GCF_030028575.1
NW_026623232.1:
3,629,000
3,630,000
3,631,000
3,632,000
3,633,000
3,634,000
3,635,000
3,636,000
3,637,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JASDEY010000001.1
70
GC Percent
30
GC Percent in 5-Base Windows
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(T)n
(AC)n
CR1-L3B_Croc
(A)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust