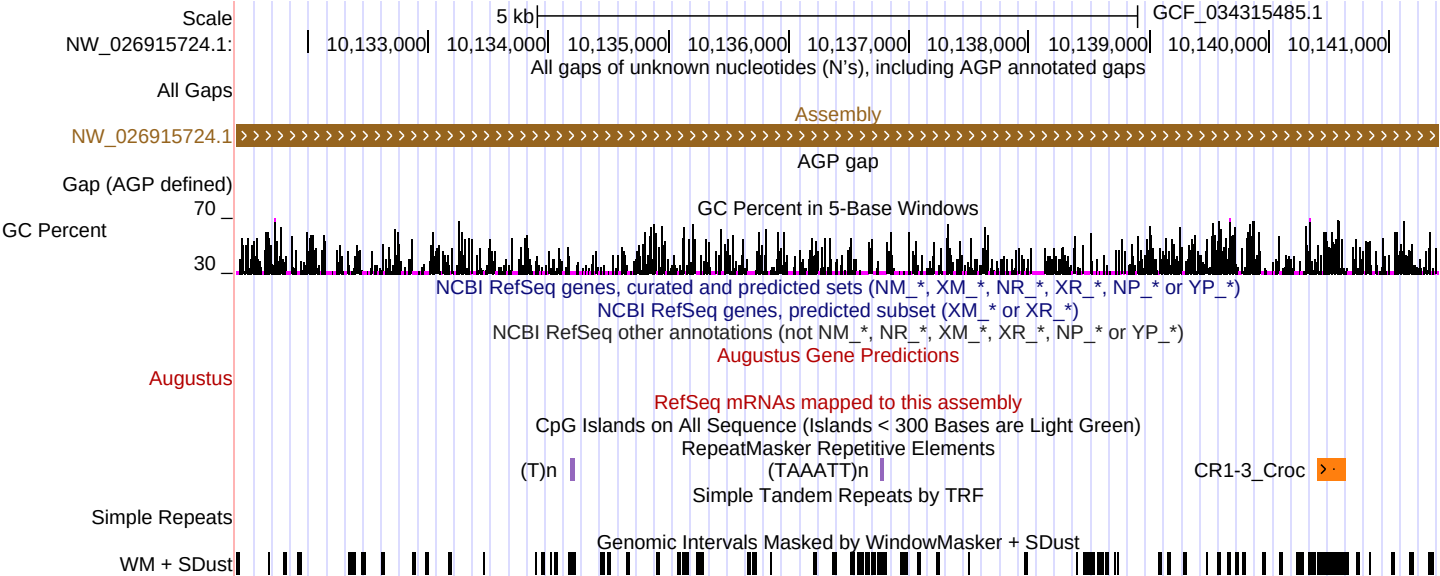
Scale
5 kb
GCF_034315485.1
NW_026915724.1:
10,133,000
10,134,000
10,135,000
10,136,000
10,137,000
10,138,000
10,139,000
10,140,000
10,141,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
NW_026915724.1
AGP gap
Gap (AGP defined)
70
GC Percent
GC Percent in 5-Base Windows
30
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
(TAAATT)n
CR1-3_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust