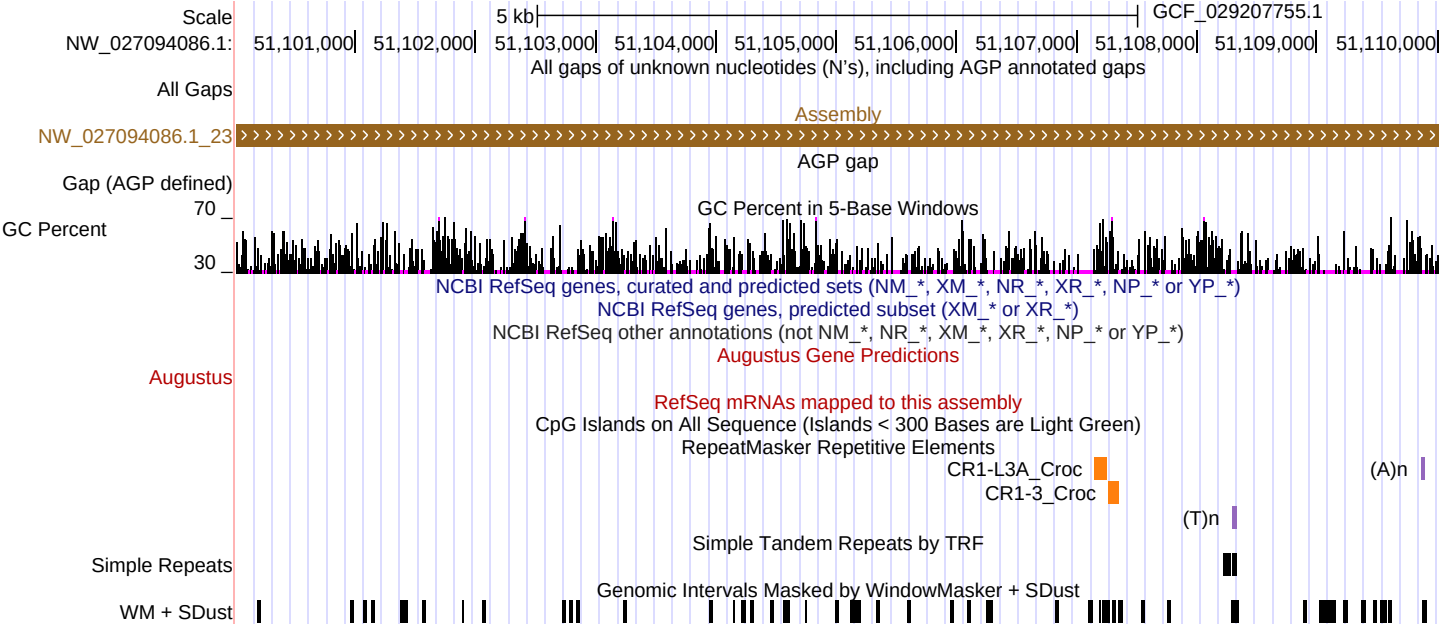
Scale
5 kb
GCF_029207755.1
NW_027094086.1:
51,101,000
51,102,000
51,103,000
51,104,000
51,105,000
51,106,000
51,107,000
51,108,000
51,109,000
51,110,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
NW_027094086.1_23
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
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
CR1-L3A_Croc
(A)n
CR1-3_Croc
(T)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust