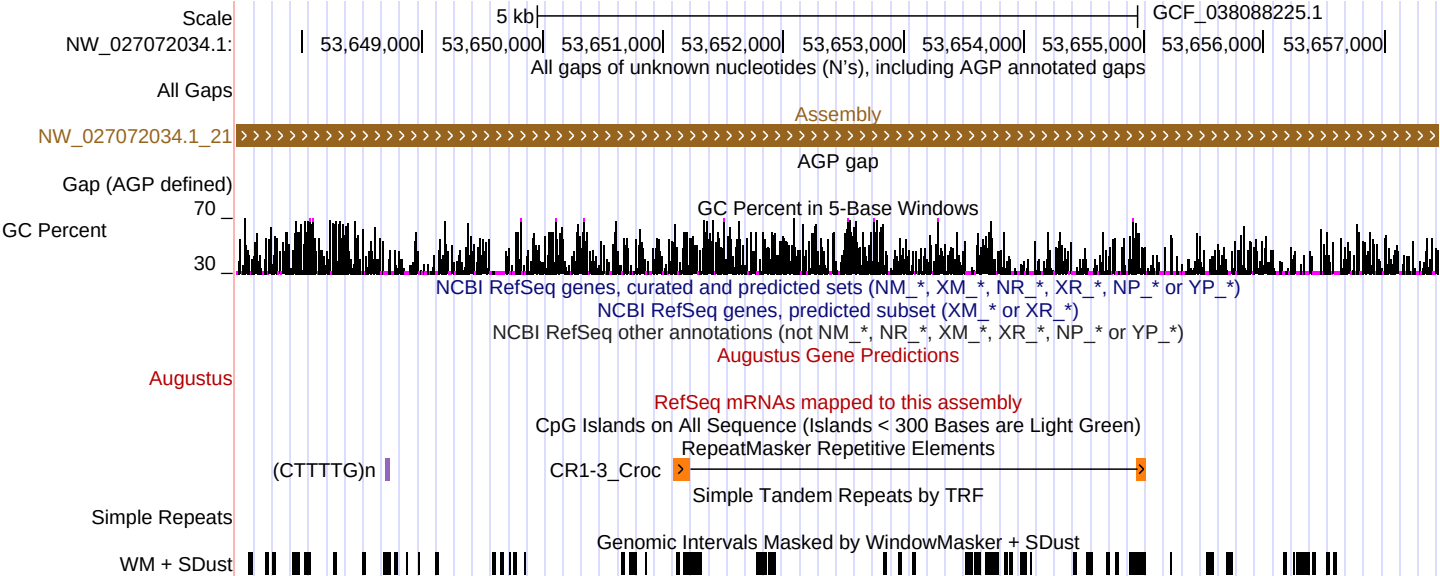
Scale
5 kb
GCF_038088225.1
NW_027072034.1:
53,649,000
53,650,000
53,651,000
53,652,000
53,653,000
53,654,000
53,655,000
53,656,000
53,657,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
NW_027072034.1_21
AGP gap
Gap (AGP defined)
70
GC Percent
30
GC Percent in 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
(CTTTTG)n
CR1-3_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust