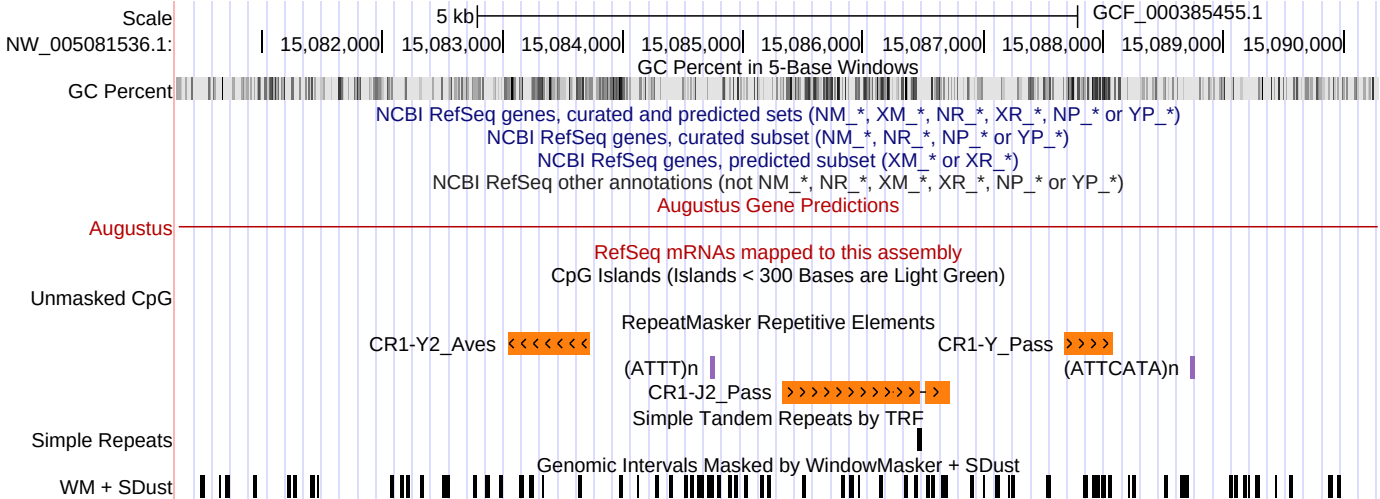
Scale
5 kb
GCF_000385455.1
NW_005081536.1:
15,082,000
15,083,000
15,084,000
15,085,000
15,086,000
15,087,000
15,088,000
15,089,000
15,090,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
NCBI RefSeq genes, curated subset (NM_*, NR_*, 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 (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
CR1-Y2_Aves
CR1-Y_Pass
(ATTT)n
(ATTCATA)n
CR1-J2_Pass
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust