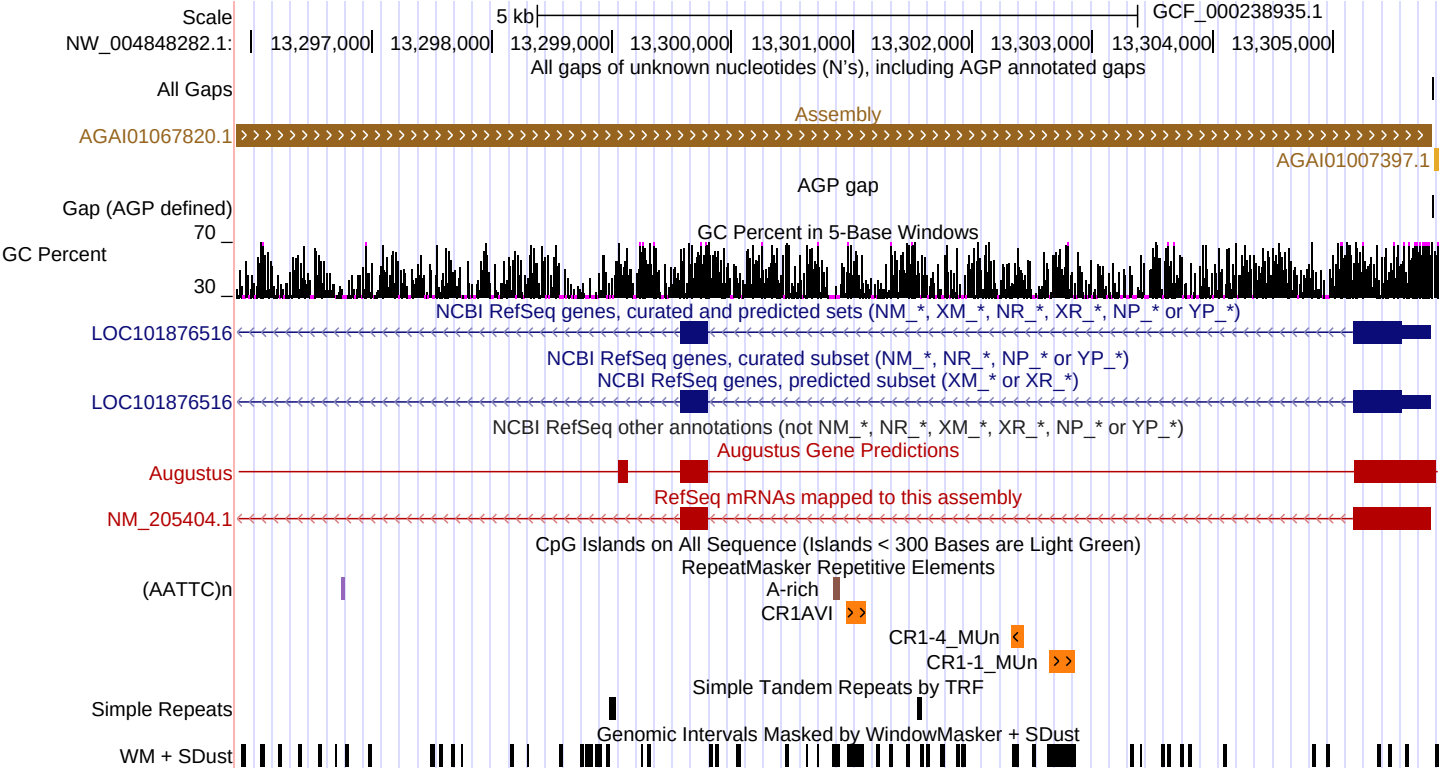
Scale
5 kb
GCF_000238935.1
NW_004848282.1:
13,297,000
13,298,000
13,299,000
13,300,000
13,301,000
13,302,000
13,303,000
13,304,000
13,305,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
AGAI01067820.1
AGAI01007397.1
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
GC Percent
70
30
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
LOC101876516
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
LOC101876516
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
NM_205404.1
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(AATTC)n
A-rich
CR1AVI
CR1-4_MUn
CR1-1_MUn
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust