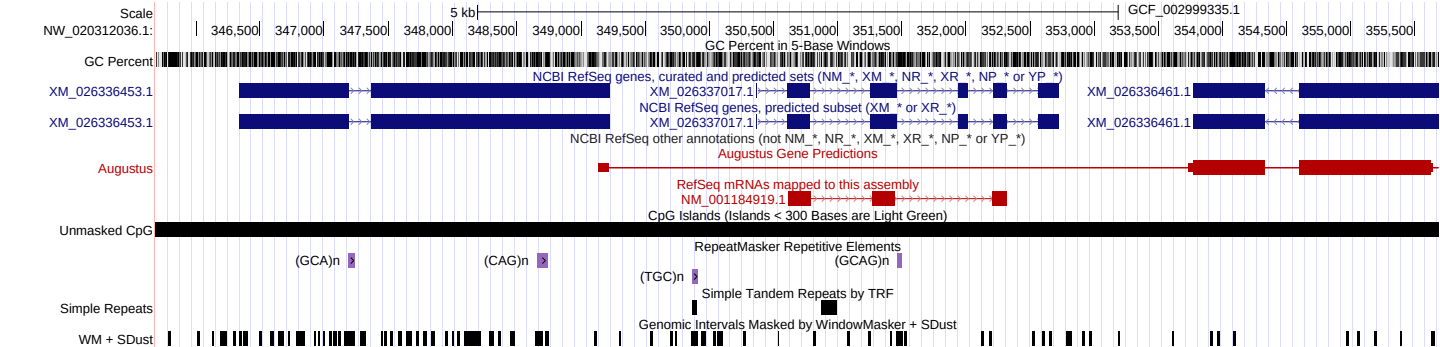
Scale
5 kb
GCF_002999335.1
NW_020312036.1:
346,500
347,000
347,500
348,000
348,500
349,000
349,500
350,000
350,500
351,000
351,500
352,000
352,500
353,000
353,500
354,000
354,500
355,000
355,500
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
XM_026336453.1
XM_026337017.1
XM_026336461.1
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
XM_026336453.1
XM_026337017.1
XM_026336461.1
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
NM_001184919.1
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
(GCA)n
(CAG)n
(GCAG)n
(TGC)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust