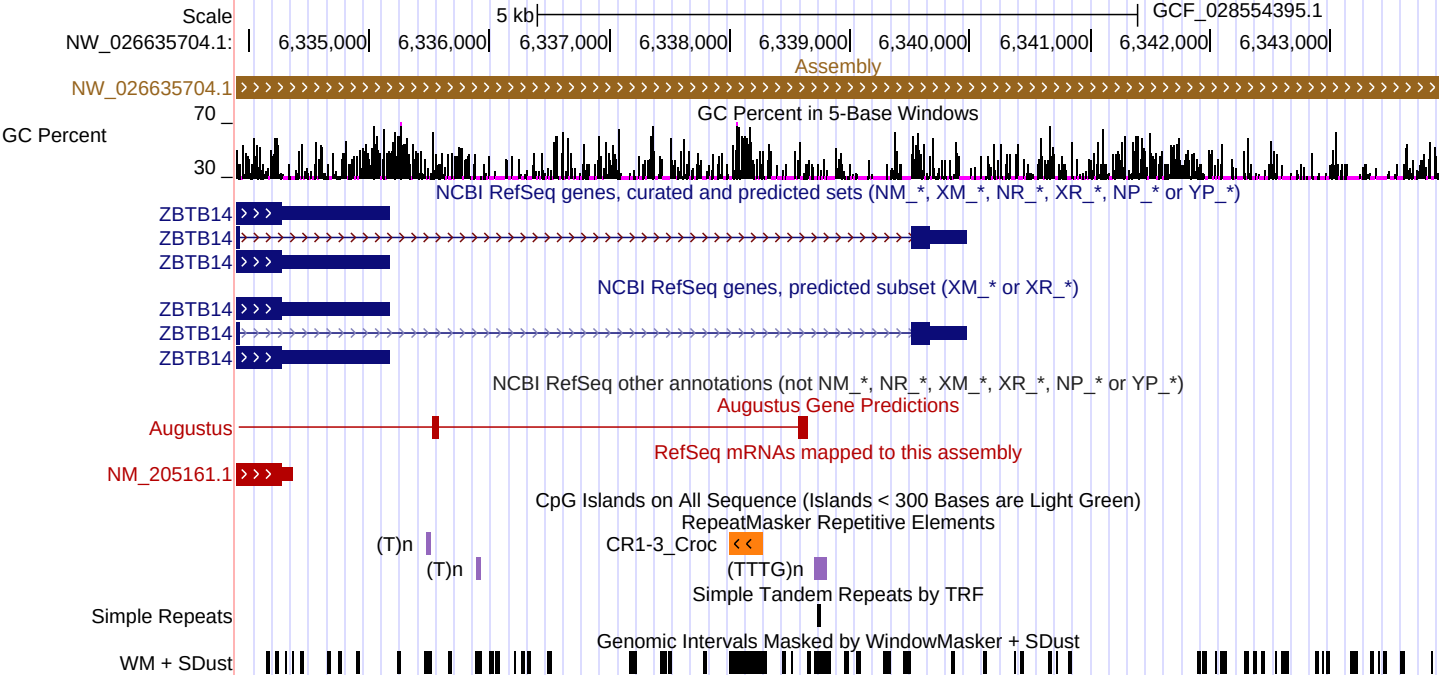
Scale
5 kb
GCF_028554395.1
NW_026635704.1:
6,335,000
6,336,000
6,337,000
6,338,000
6,339,000
6,340,000
6,341,000
6,342,000
6,343,000
Assembly
NW_026635704.1
70
GC Percent
30
GC Percent in 5-Base Windows
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
ZBTB14
ZBTB14
ZBTB14
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
ZBTB14
ZBTB14
ZBTB14
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
NM_205161.1
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(T)n
CR1-3_Croc
(T)n
(TTTG)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust