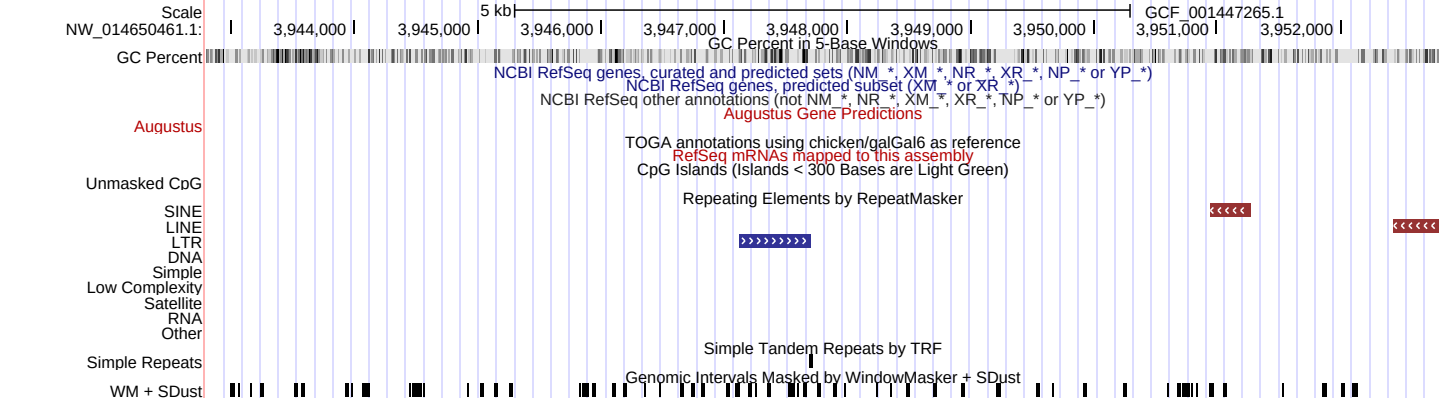
Scale
5 kb
GCF_001447265.1
NW_014650461.1:
3,944,000
3,945,000
3,946,000
3,947,000
3,948,000
3,949,000
3,950,000
3,951,000
3,952,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus
Augustus Gene Predictions
TOGA annotations using chicken/galGal6 as reference
RefSeq mRNAs mapped to this assembly
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
Repeating Elements by RepeatMasker
SINE
LINE
LTR
DNA
Simple
Low Complexity
Satellite
RNA
Other
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust