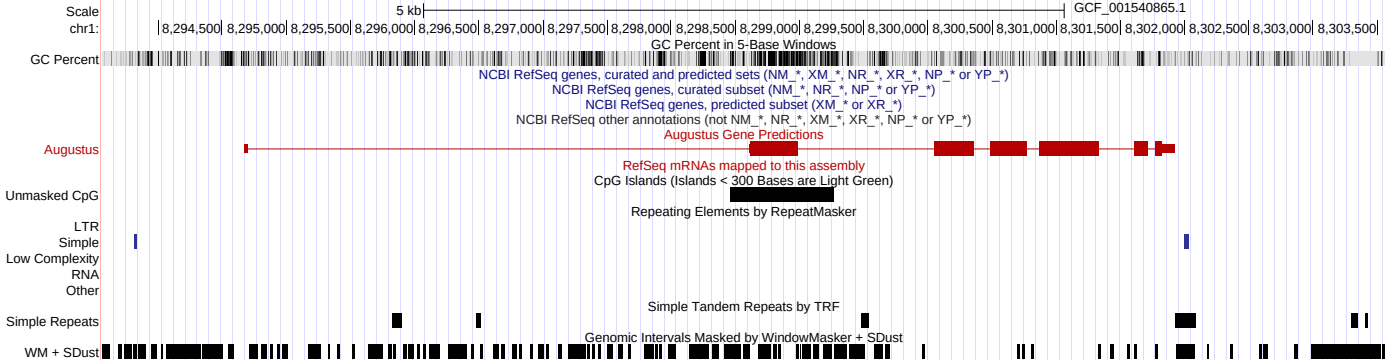
Scale
chr1:
5 kb
GCF_001540865.1
8,294,500
8,295,000
8,295,500
8,296,000
8,296,500
8,297,000
8,297,500
8,298,000
8,298,500
8,299,000
8,299,500
8,300,000
8,300,500
8,301,000
8,301,500
8,302,000
8,302,500
8,303,000
8,303,500
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted 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
Repeating Elements by RepeatMasker
LTR
Simple
Low Complexity
RNA
Other
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust