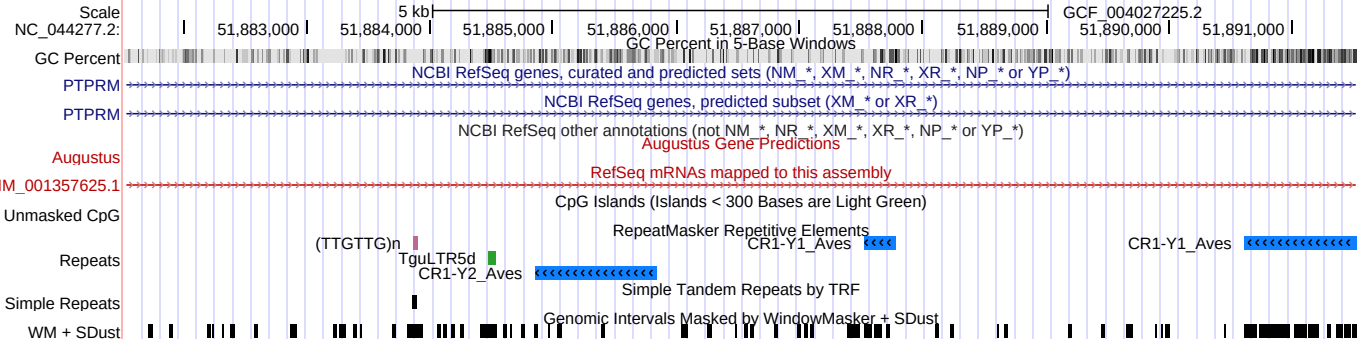
Scale
5 kb
GCF_004027225.2
NC_044277.2:
51,883,000
51,884,000
51,885,000
51,886,000
51,887,000
51,888,000
51,889,000
51,890,000
51,891,000
GC Percent
GC Percent in 5-Base Windows
PTPRM
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
PTPRM
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus
Augustus Gene Predictions
NM_001357625.1
RefSeq mRNAs mapped to this assembly
Unmasked CpG
CpG Islands (Islands < 300 Bases are Light Green)
Repeats
RepeatMasker Repetitive Elements
(TTGTTG)n
TguLTR5d
CR1-Y2_Aves
CR1-Y1_Aves
CR1-Y1_Aves
Simple Repeats
Simple Tandem Repeats by TRF
WM + SDust
Genomic Intervals Masked by WindowMasker + SDust