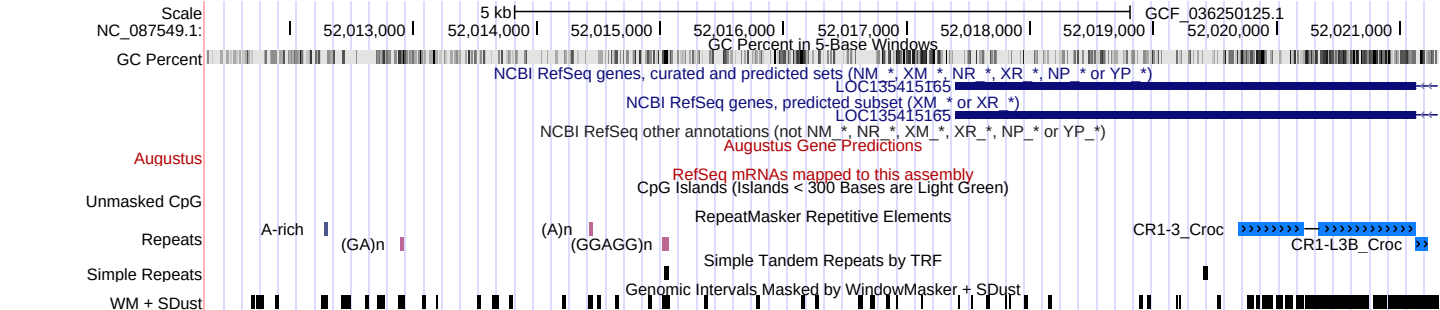
Scale
5 kb
GCF_036250125.1
NC_087549.1:
52,013,000
52,014,000
52,015,000
52,016,000
52,017,000
52,018,000
52,019,000
52,020,000
52,021,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
LOC135415165
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
LOC135415165
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
RepeatMasker Repetitive Elements
A-rich
(GA)n
(A)n
(GGAGG)n
CR1-3_Croc
CR1-L3B_Croc
Repeats
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust