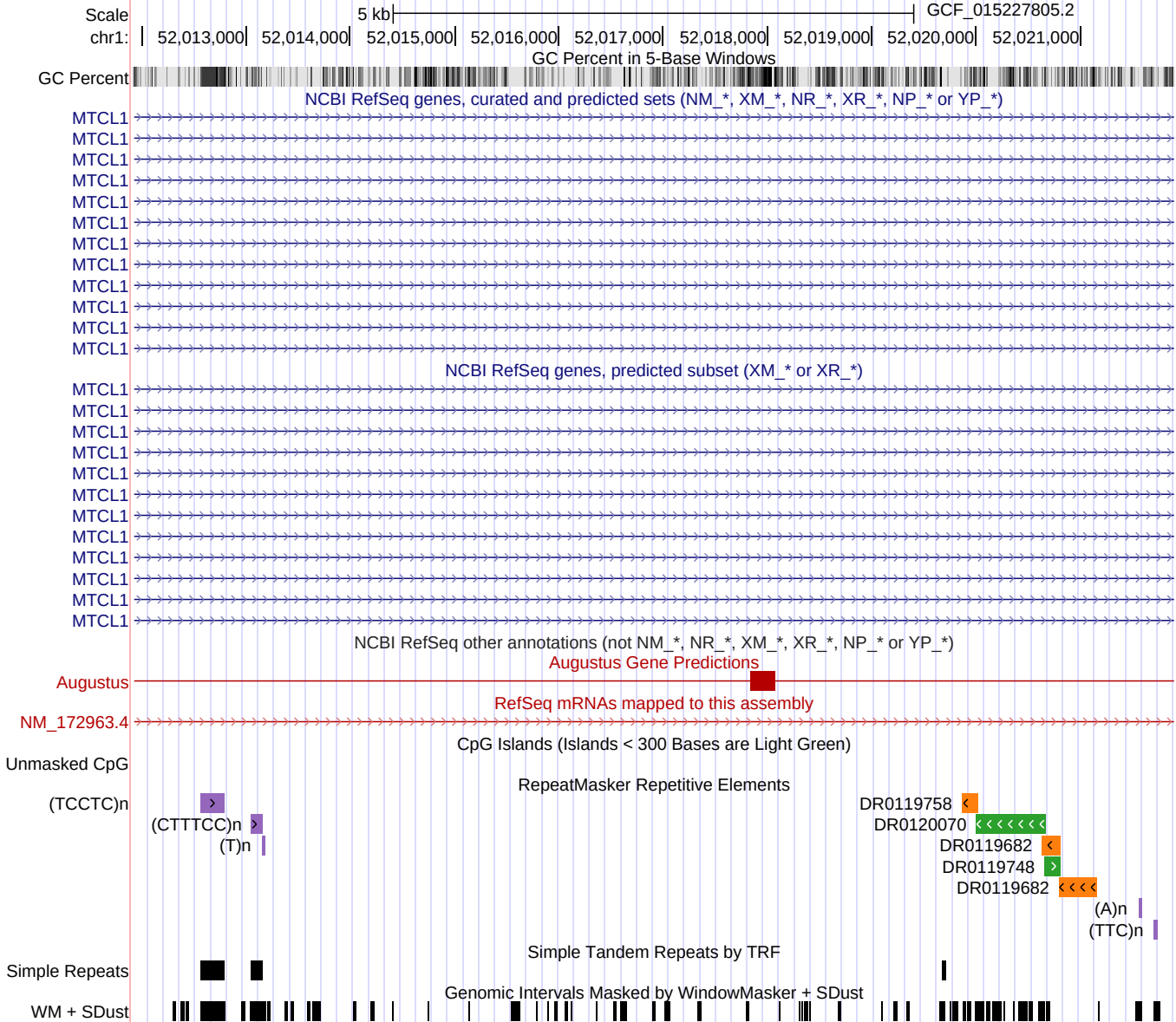
Scale
5 kb
GCF_015227805.2
chr1:
52,013,000
52,014,000
52,015,000
52,016,000
52,017,000
52,018,000
52,019,000
52,020,000
52,021,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
MTCL1
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
NM_172963.4
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
(TCCTC)n
(CTTTCC)n
(T)n
DR0119758
DR0120070
DR0119682
DR0119748
DR0119682
(A)n
(TTC)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust