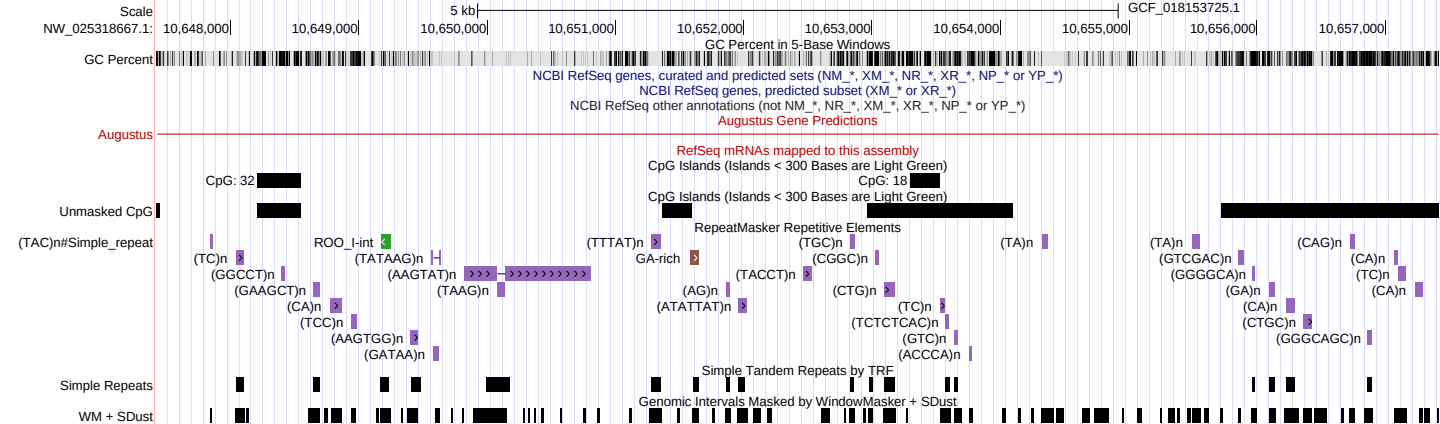
Scale
NW_025318667.1:
5 kb
GCF_018153725.1
10,648,000
10,649,000
10,650,000
10,651,000
10,652,000
10,653,000
10,654,000
10,655,000
10,656,000
10,657,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
CpG Islands (Islands < 300 Bases are Light Green)
CpG: 32
CpG: 18
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
(TAC)n#Simple_repeat
ROO_I-int
(TTTAT)n
(TGC)n
(TA)n
(TA)n
(CAG)n
(TC)n
(TATAAG)n
GA-rich
(CGGC)n
(GTCGAC)n
(CA)n
(GGCCT)n
(AAGTAT)n
(TACCT)n
(GGGGCA)n
(TC)n
(GAAGCT)n
(TAAG)n
(AG)n
(CTG)n
(GA)n
(CA)n
(CA)n
(ATATTAT)n
(TC)n
(CA)n
(TCC)n
(TCTCTCAC)n
(CTGC)n
(AAGTGG)n
(GTC)n
(GGGCAGC)n
(GATAA)n
(ACCCA)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust