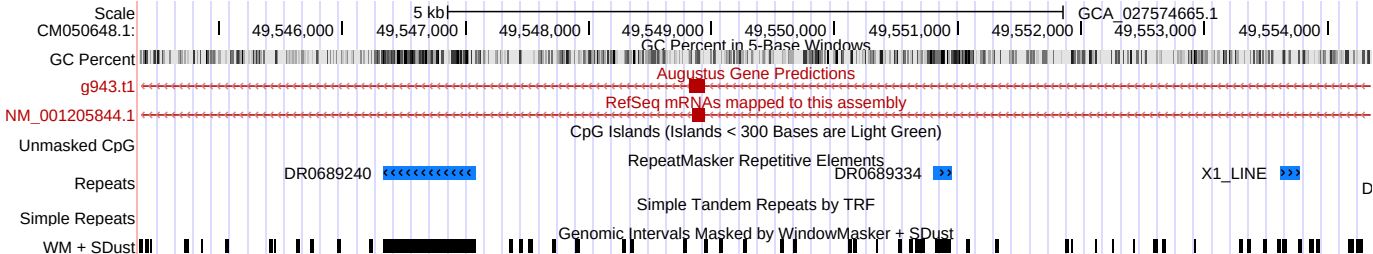
Scale
5 kb
GCA_027574665.1
CM050648.1:
49,546,000
49,547,000
49,548,000
49,549,000
49,550,000
49,551,000
49,552,000
49,553,000
49,554,000
GC Percent
GC Percent in 5-Base Windows
g943.t1
Augustus Gene Predictions
NM_001205844.1
RefSeq mRNAs mapped to this assembly
Unmasked CpG
CpG Islands (Islands < 300 Bases are Light Green)
Repeats
RepeatMasker Repetitive Elements
DR0689240
DR0689334
X1_LINE
Simple Repeats
Simple Tandem Repeats by TRF
WM + SDust
Genomic Intervals Masked by WindowMasker + SDust