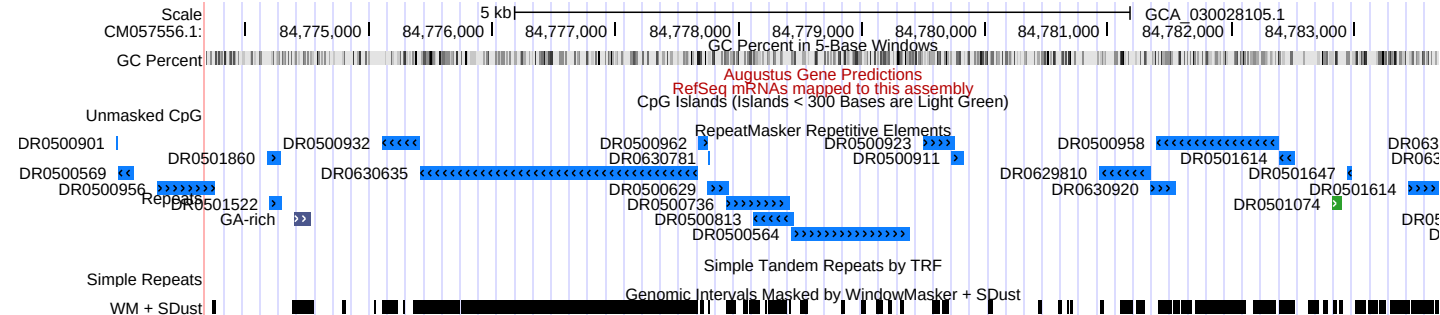
Scale
5 kb
GCA_030028105.1
CM057556.1:
84,775,000
84,776,000
84,777,000
84,778,000
84,779,000
84,780,000
84,781,000
84,782,000
84,783,000
GC Percent in 5-Base Windows
GC Percent
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
DR0500901
DR0500932
DR0500962
DR0500923
DR0500958
DR0501860
DR0630781
DR0500911
DR0501614
DR0500569
DR0630635
DR0629810
DR0501647
DR0500956
DR0630920
DR0501614
Repeats
DR0501522
DR0500629
DR0500736
DR0501074
GA-rich
DR0500813
DR0500564
Simple Repeats
Simple Tandem Repeats by TRF
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust