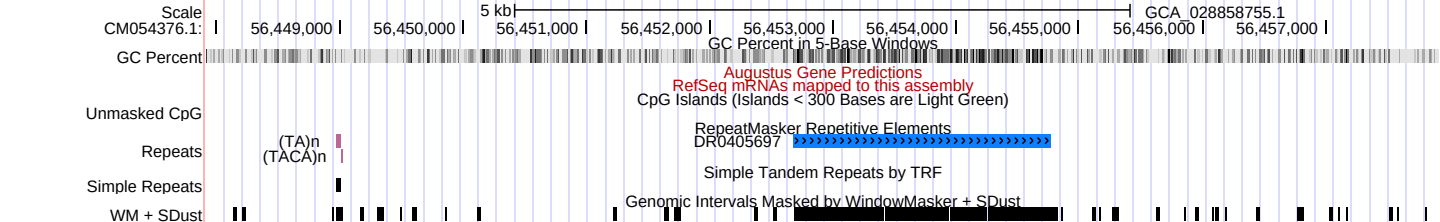
Scale
5 kb
GCA_028858755.1
CM054376.1:
56,449,000
56,450,000
56,451,000
56,452,000
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56,457,000
GC Percent in 5-Base Windows
GC Percent
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
DR0405697
(TA)n
(TACA)n
Repeats
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust