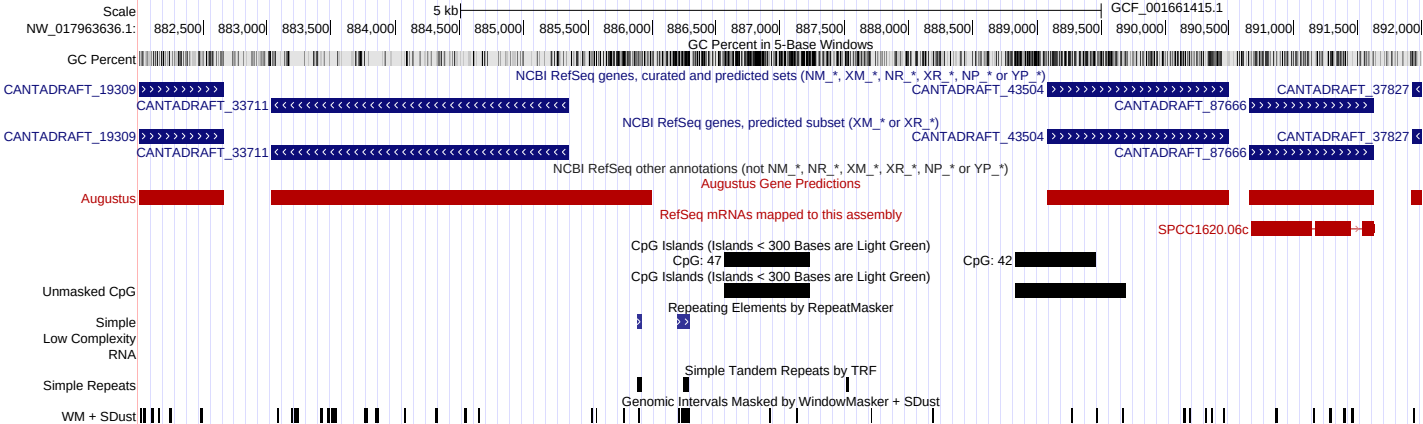
Scale
5 kb
GCF_001661415.1
NW_017963636.1:
882,500
883,000
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892,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
CANTADRAFT_19309
CANTADRAFT_43504
CANTADRAFT_37827
CANTADRAFT_33711
CANTADRAFT_87666
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
CANTADRAFT_19309
CANTADRAFT_43504
CANTADRAFT_37827
CANTADRAFT_33711
CANTADRAFT_87666
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
SPCC1620.06c
CpG Islands (Islands < 300 Bases are Light Green)
CpG: 47
CpG: 42
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
Repeating Elements by RepeatMasker
Simple
Low Complexity
RNA
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust