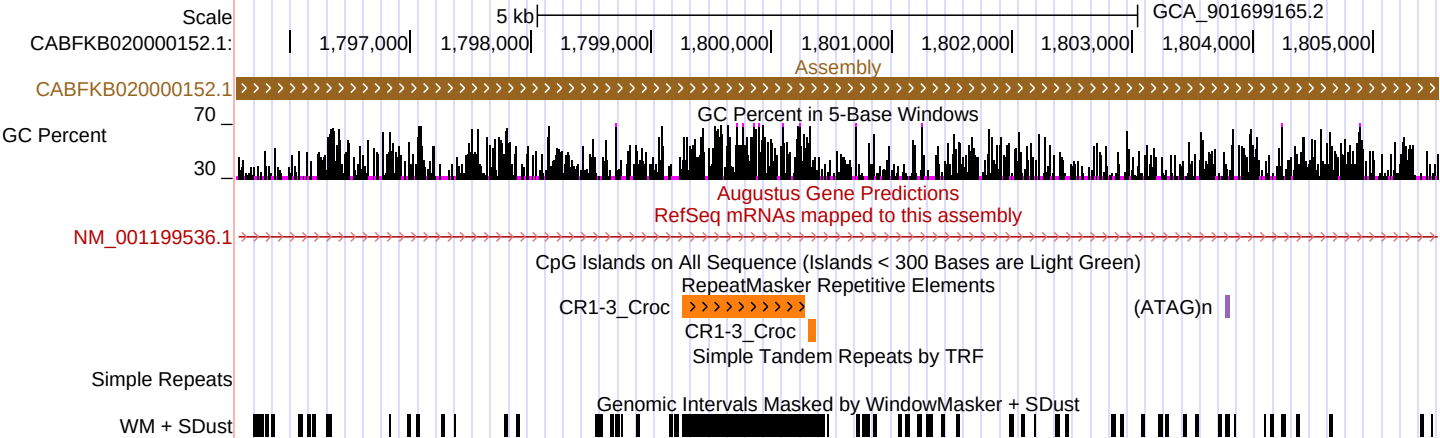
Scale
5 kb
GCA_901699165.2
CABFKB020000152.1:
1,797,000
1,798,000
1,799,000
1,800,000
1,801,000
1,802,000
1,803,000
1,804,000
1,805,000
Assembly
CABFKB020000152.1
GC Percent
70
30
GC Percent in 5-Base Windows
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
NM_001199536.1
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
CR1-3_Croc
(ATAG)n
CR1-3_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust