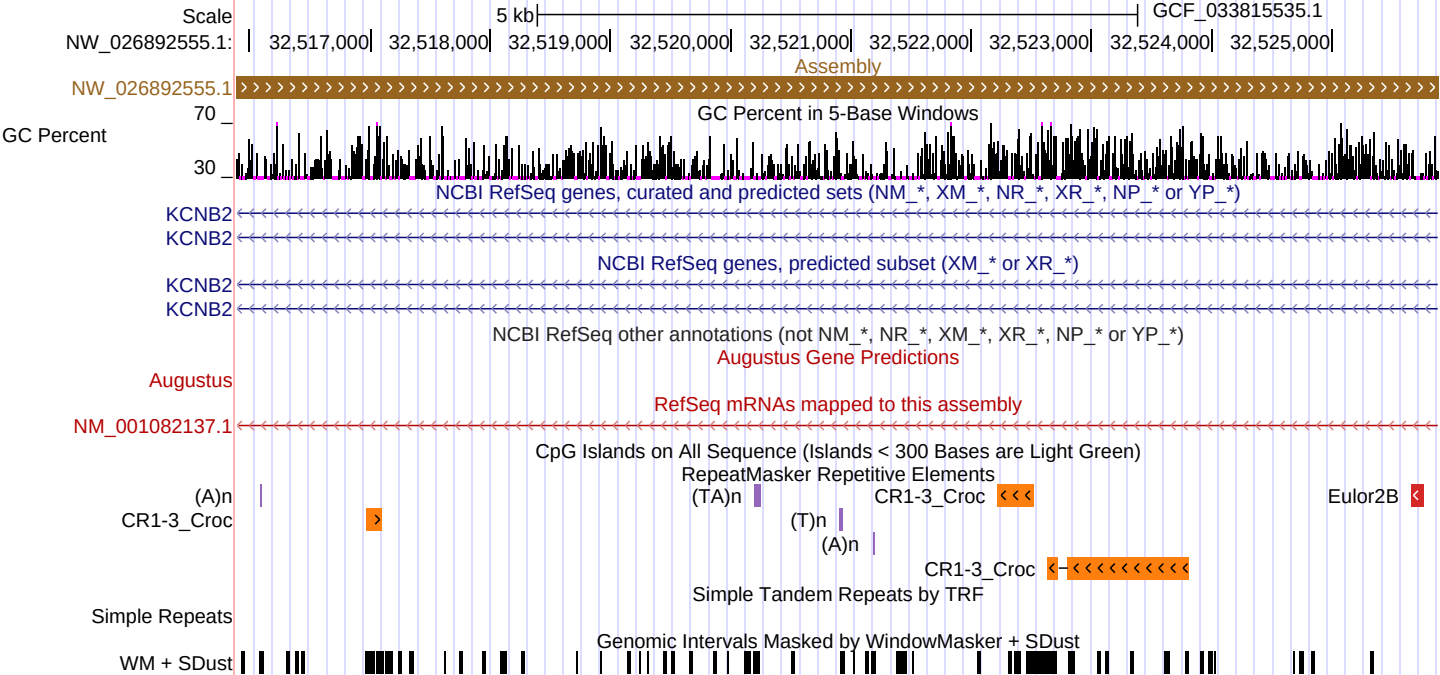
Scale
5 kb
GCF_033815535.1
NW_026892555.1:
32,517,000
32,518,000
32,519,000
32,520,000
32,521,000
32,522,000
32,523,000
32,524,000
32,525,000
Assembly
NW_026892555.1
GC Percent
70
30
GC Percent in 5-Base Windows
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
KCNB2
KCNB2
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
KCNB2
KCNB2
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
NM_001082137.1
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(A)n
(TA)n
CR1-3_Croc
Eulor2B
CR1-3_Croc
(T)n
(A)n
CR1-3_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust