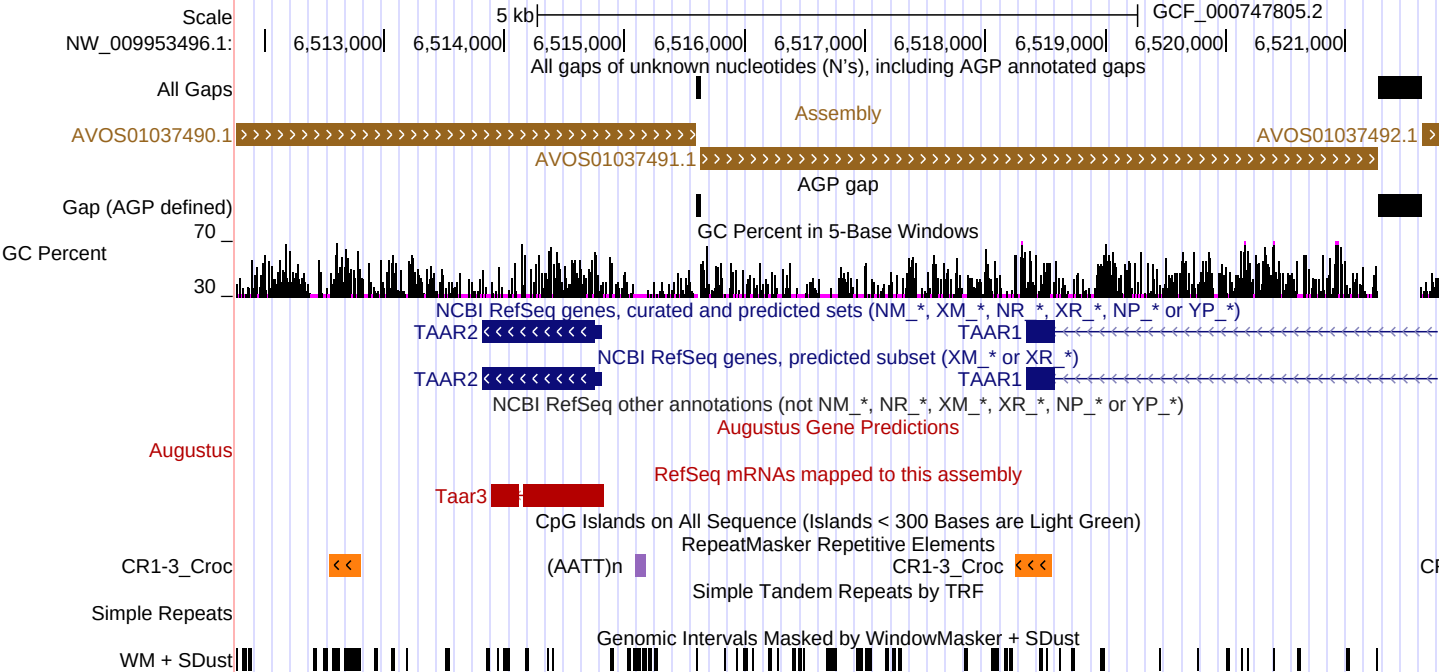
Scale
5 kb
GCF_000747805.2
NW_009953496.1:
6,513,000
6,514,000
6,515,000
6,516,000
6,517,000
6,518,000
6,519,000
6,520,000
6,521,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
AVOS01037490.1
AVOS01037492.1
AVOS01037491.1
AGP gap
Gap (AGP defined)
70
GC Percent in 5-Base Windows
GC Percent
30
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
TAAR2
TAAR1
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
TAAR2
TAAR1
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
Taar3
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
CR1-3_Croc
(AATT)n
CR1-3_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust