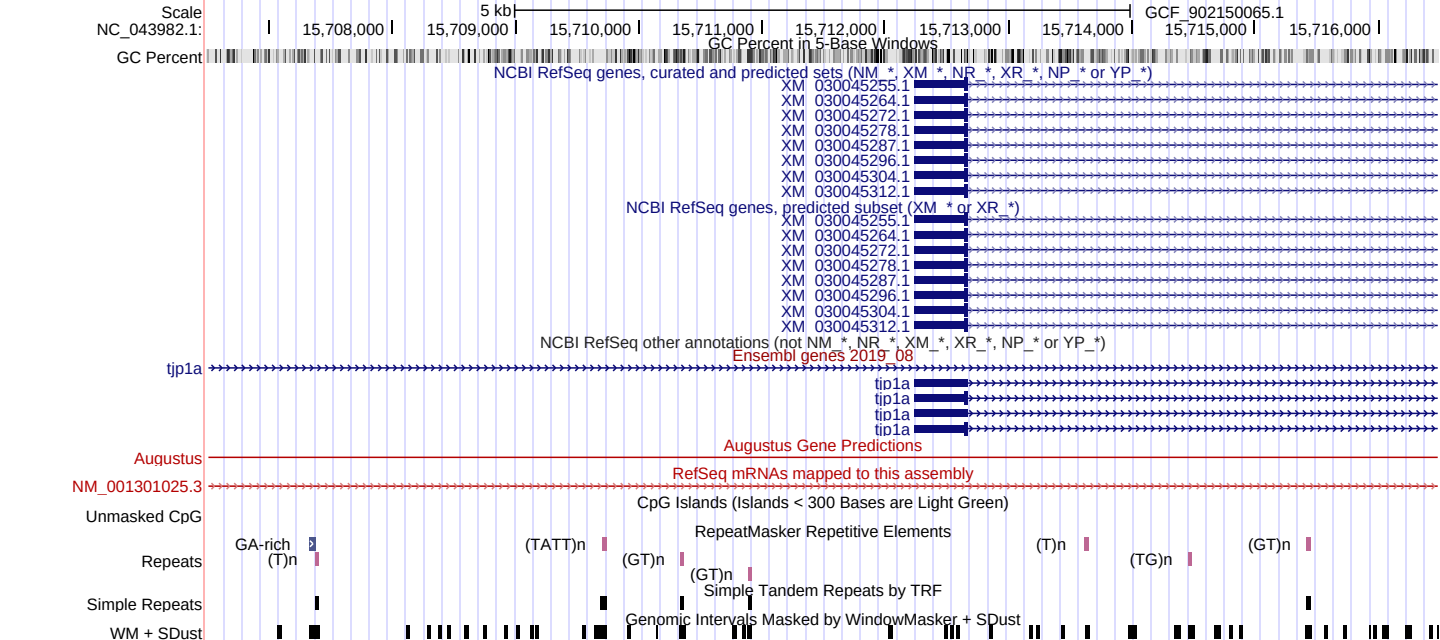
Scale
5 kb
GCF_902150065.1
NC_043982.1:
15,708,000
15,709,000
15,710,000
15,711,000
15,712,000
15,713,000
15,714,000
15,715,000
15,716,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
XM_030045255.1
XM_030045264.1
XM_030045272.1
XM_030045278.1
XM_030045287.1
XM_030045296.1
XM_030045304.1
XM_030045312.1
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
XM_030045255.1
XM_030045264.1
XM_030045272.1
XM_030045278.1
XM_030045287.1
XM_030045296.1
XM_030045304.1
XM_030045312.1
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Ensembl genes 2019_08
tjp1a
tjp1a
tjp1a
tjp1a
tjp1a
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
NM_001301025.3
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
GA-rich
(T)n
(TATT)n
(GT)n
(GT)n
(T)n
(TG)n
(GT)n
Repeats
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust