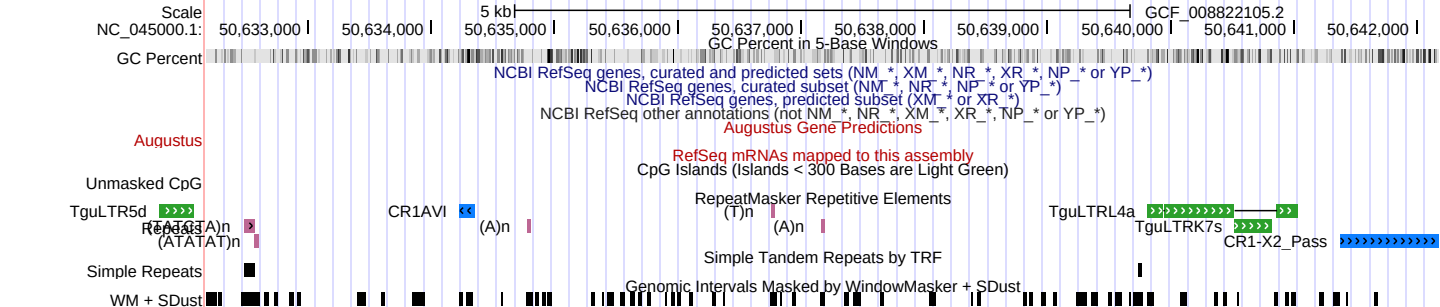
Scale
5 kb
GCF_008822105.2
NC_045000.1:
50,633,000
50,634,000
50,635,000
50,636,000
50,637,000
50,638,000
50,639,000
50,640,000
50,641,000
50,642,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMaster Repetitive Elements
TguLTR5d
CR1AVI
(T)n
TguLTRL4a
Repeats
(TTATA)n
(A)n
(A)n
TguLTRK7s
(ATATAT)n
CR1-X2_Pass
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust