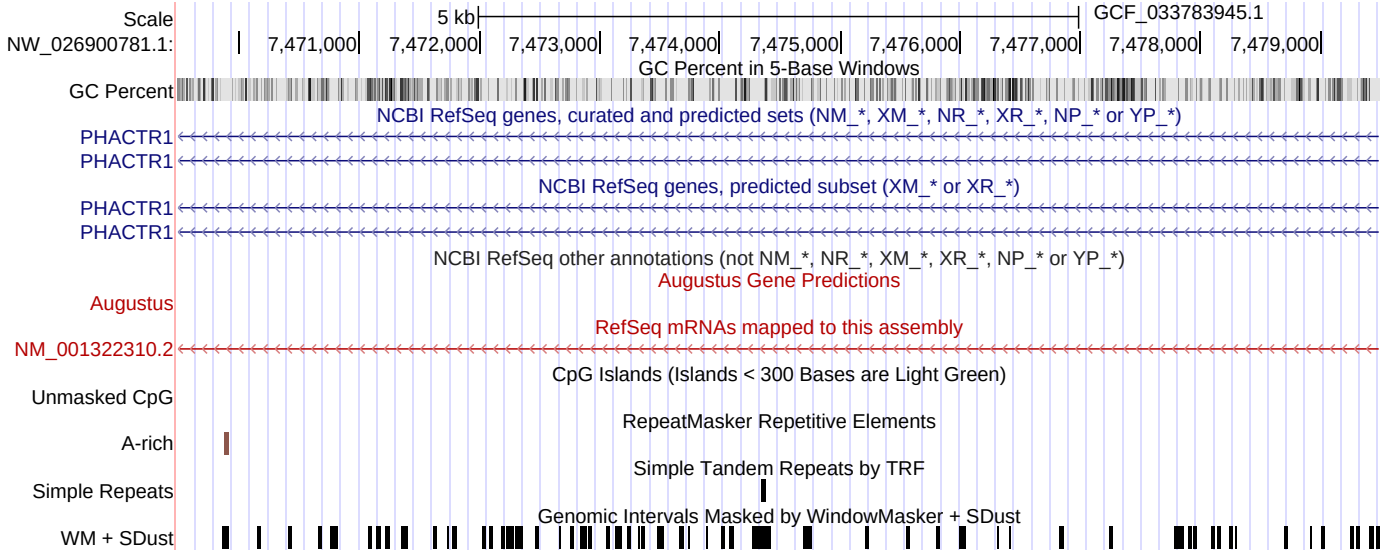
Scale
5 kb
GCF_033783945.1
NW_026900781.1:
7,471,000
7,472,000
7,473,000
7,474,000
7,475,000
7,476,000
7,477,000
7,478,000
7,479,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
PHACTR1
PHACTR1
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
PHACTR1
PHACTR1
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
NM_001322310.2
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
A-rich
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust