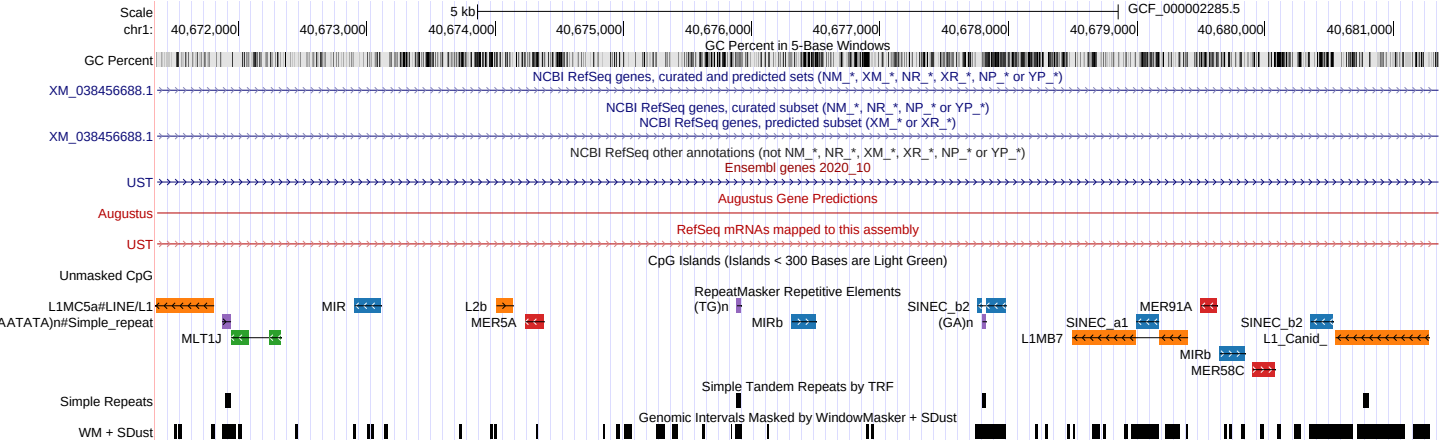
Scale
chr1:
5 kb
GCF_000002285.5
40,672,000
40,673,000
40,674,000
40,675,000
40,676,000
40,677,000
40,678,000
40,679,000
40,680,000
40,681,000
GC Percent
GC Percent in 5-Base Windows
XM_038456688.1
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
XM_038456688.1
NCBI 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_*)
UST
Ensembl genes 2020_10
Augustus
Augustus Gene Predictions
UST
RefSeq mRNAs mapped to this assembly
Unmasked CpG
CpG Islands (Islands < 300 Bases are Light Green)
L1MC5a#LINE/L1
AATATA)n#Simple_repeat
MLT1J
MIR
L2b
MER5A
RepeatMaster Repetitive Elements
(TG)n
MIRb
SINEC_b2
(GA)n
MER91A
SINEC_a1
L1MB7
MIRb
MER58C
SINEC_b2
L1_Canid_
Simple Repeats
Simple Tandem Repeats by TRF
WM + SDust
Genomic Intervals Masked by WindowMasker + SDust