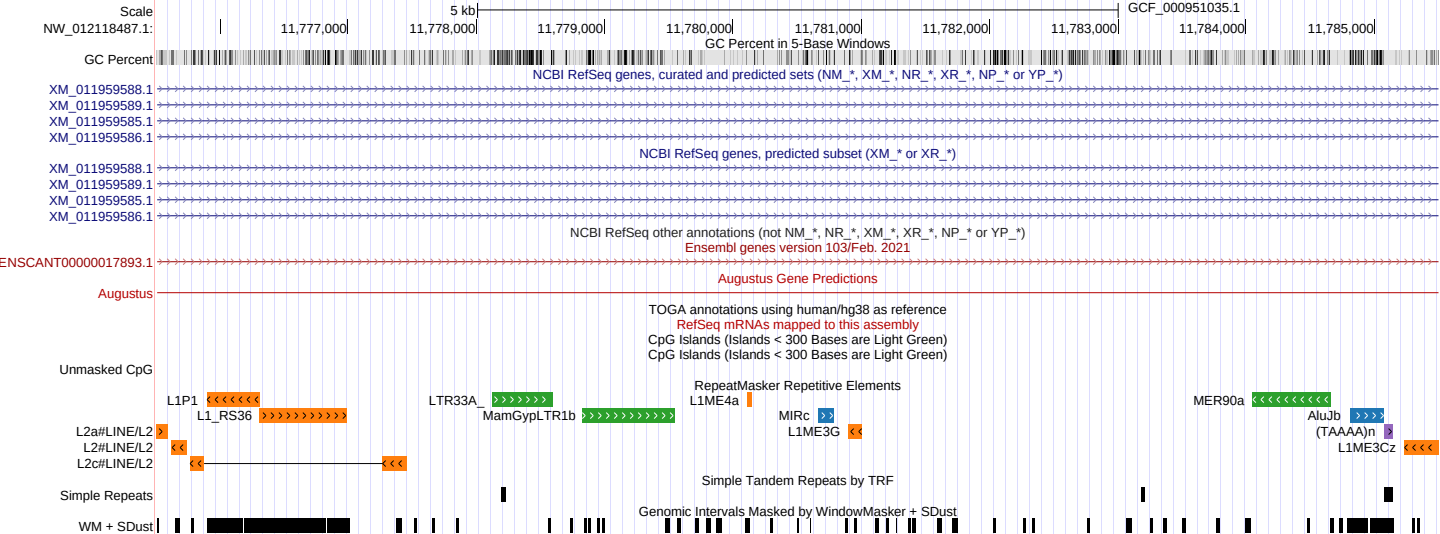
Scale
5 kb
GCF_000951035.1
NW_012118487.1:
11,777,000
11,778,000
11,779,000
11,780,000
11,781,000
11,782,000
11,783,000
11,784,000
11,785,000
GC Percent
GC Percent in 5-Base Windows
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
XM_011959588.1
XM_011959589.1
XM_011959585.1
XM_011959586.1
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
XM_011959588.1
XM_011959589.1
XM_011959585.1
XM_011959586.1
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Ensembl genes version 103/Feb. 2021
ENSCANT00000017893.1
Augustus Gene Predictions
Augustus
TOGA annotations using human/hg38 as reference
RefSeq mRNAs mapped to this assembly
CpG Islands (Islands < 300 Bases are Light Green)
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
L1P1
L1_RS36
LTR33A_
MamGypLTR1b
L1ME4a
MIRc
L1ME3G
MER90a
AluJb
(TAAAA)n
L1ME3Cz
L2a#LINE/L2
L2#LINE/L2
L2c#LINE/L2
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust