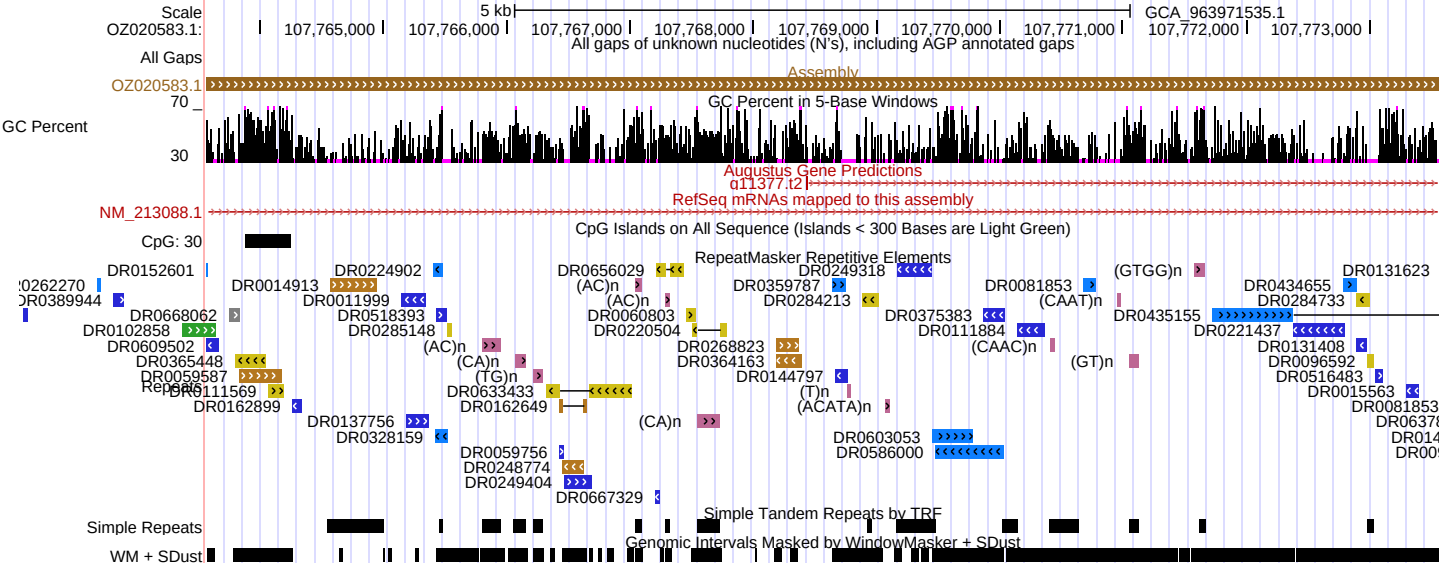
Scale
5 kb
GCA_963971535.1
OZ020583.1:
107,765,000
107,766,000
107,767,000
107,768,000
107,769,000
107,770,000
107,771,000
107,772,000
107,773,000
All Gaps
All gaps of unknown nucleotides (N's), including AGP annotated gaps
OZ020583.1
Assembly
70
GC Percent
30
GC Percent in 5-Base Windows
Augustus Gene Predictions
g11377.t2
NM_213088.1
RefSeq mRNAs mapped to this assembly
CpG: 30
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
DR0152601
DR0224902
DR0656029
DR0249318
(GTGG)n
DR0131623
DR0262270
DR0014913
(AC)n
DR0359787
DR0081853
DR0434655
DR0389944
DR0011999
(AC)n
DR0284213
(CAAT)n
DR0284733
DR0668062
DR0518393
DR0060803
DR0375383
DR0435155
DR0102858
DR0285148
DR0220504
DR0111884
DR0221437
DR0609502
(AC)n
DR0268823
(CAAC)n
DR0131408
DR0365448
(CA)n
DR0364163
(GT)n
DR0096592
DR0059587
(TG)n
DR0144797
DR0516483
Repeats
DR0111569
DR0633433
(T)n
DR0015563
DR0162899
DR0162649
(ACATA)n
DR0081853
DR0137756
(CA)n
DR0637
DR0328159
DR0603053
DR014
DR0586000
DR00
DR0059756
DR0248774
DR0249404
DR0667329
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust