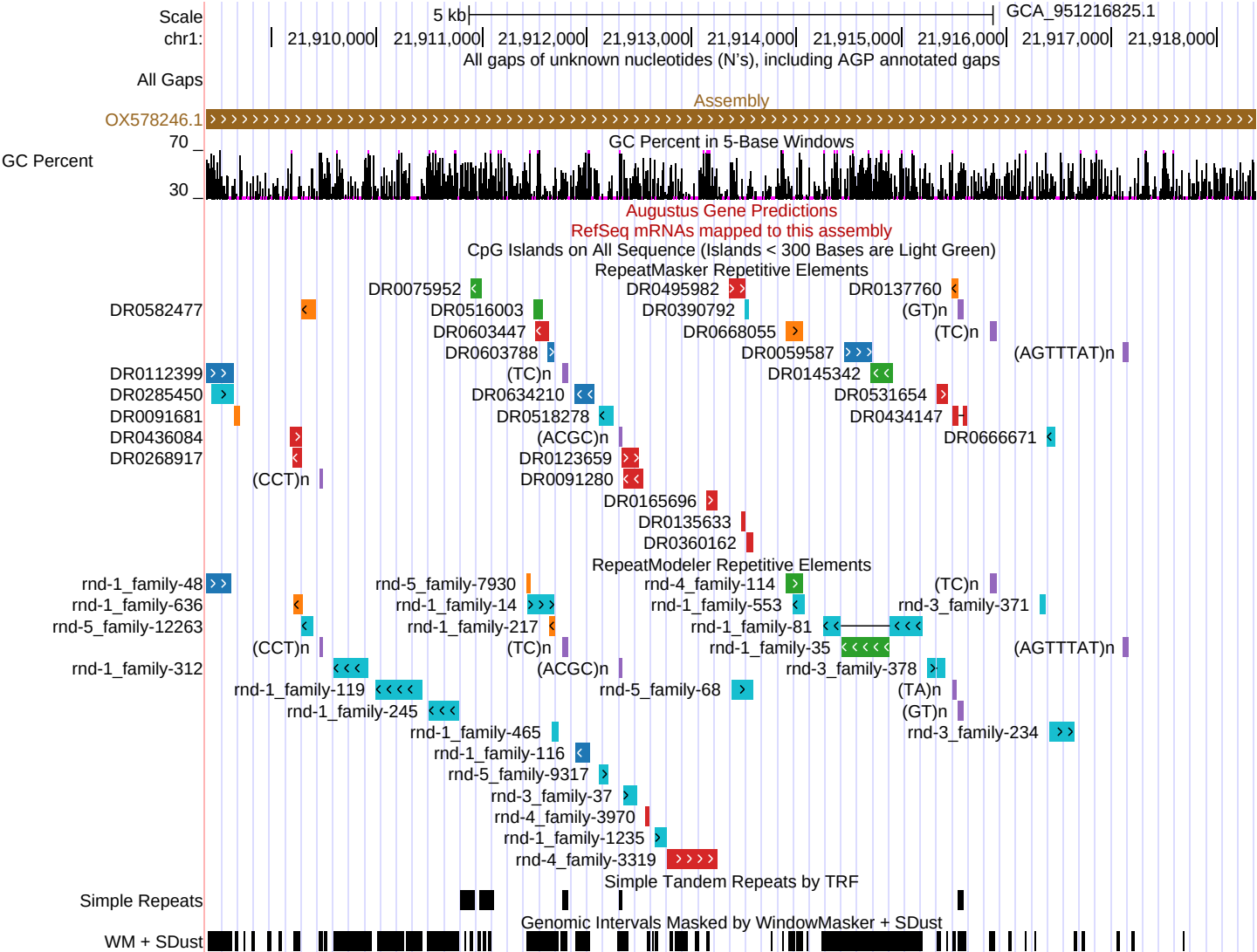
Scale
5 kb
GCA_951216825.1
chr1:
21,910,000
21,911,000
21,912,000
21,913,000
21,914,000
21,915,000
21,916,000
21,917,000
21,918,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
OX578246.1
GC Percent in 5-Base Windows
GC Percent
70
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
DR0075952
DR0495982
DR0137760
DR0582477
DR0516003
DR0390792
(GT)n
DR0603447
DR0668055
(TC)n
DR0603788
DR0059587
(AGTTTAT)n
DR0112399
(TC)n
DR0145342
DR0285450
DR0634210
DR0531654
DR0091681
DR0518278
DR0434147
DR0436084
(ACGC)n
DR0666671
DR0268917
DR0123659
(CCT)n
DR0091280
DR0165696
DR0135633
DR0360162
RepeatModeler Repetitive Elements
rnd-1_family-48
rnd-5_family-7930
rnd-4_family-114
(TC)n
rnd-1_family-636
rnd-1_family-14
rnd-1_family-553
rnd-3_family-371
rnd-5_family-12263
rnd-1_family-217
rnd-1_family-81
(CCT)n
(TC)n
rnd-1_family-35
(AGTTTAT)n
rnd-1_family-312
(ACGC)n
rnd-3_family-378
rnd-1_family-119
rnd-5_family-68
(TA)n
rnd-1_family-245
(GT)n
rnd-1_family-465
rnd-3_family-234
rnd-1_family-116
rnd-5_family-9317
rnd-3_family-37
rnd-4_family-3970
rnd-1_family-1235
rnd-4_family-3319
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust