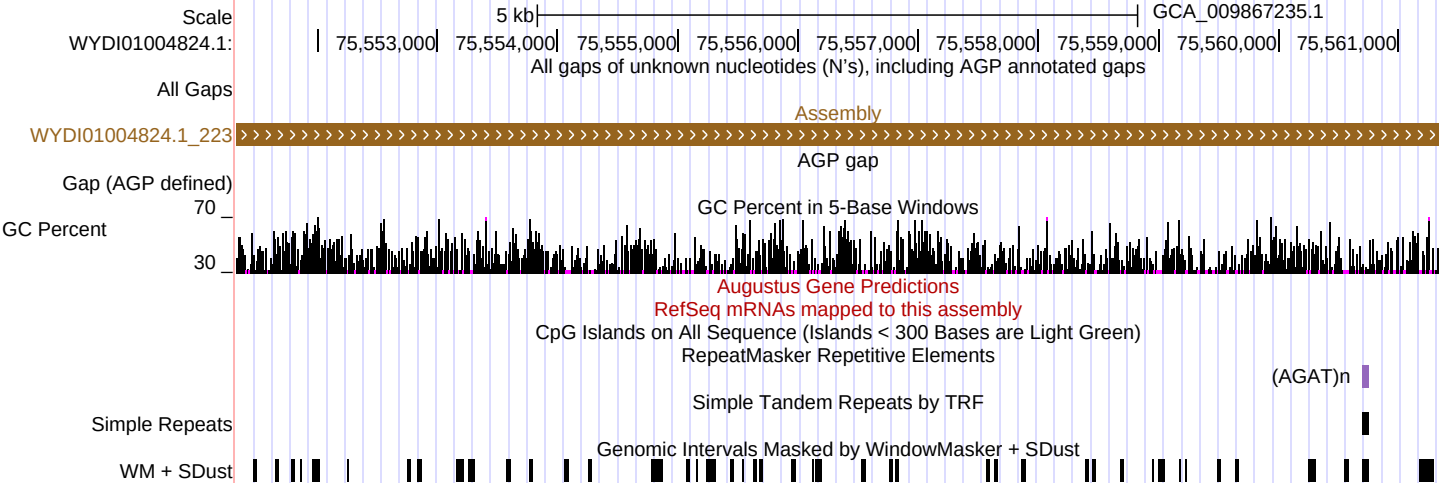
Scale
5 kb
GCA_009867235.1
WYDI01004824.1:
75,553,000
75,554,000
75,555,000
75,556,000
75,557,000
75,558,000
75,559,000
75,560,000
75,561,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
WYDI01004824.1_223
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(AGAT)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust