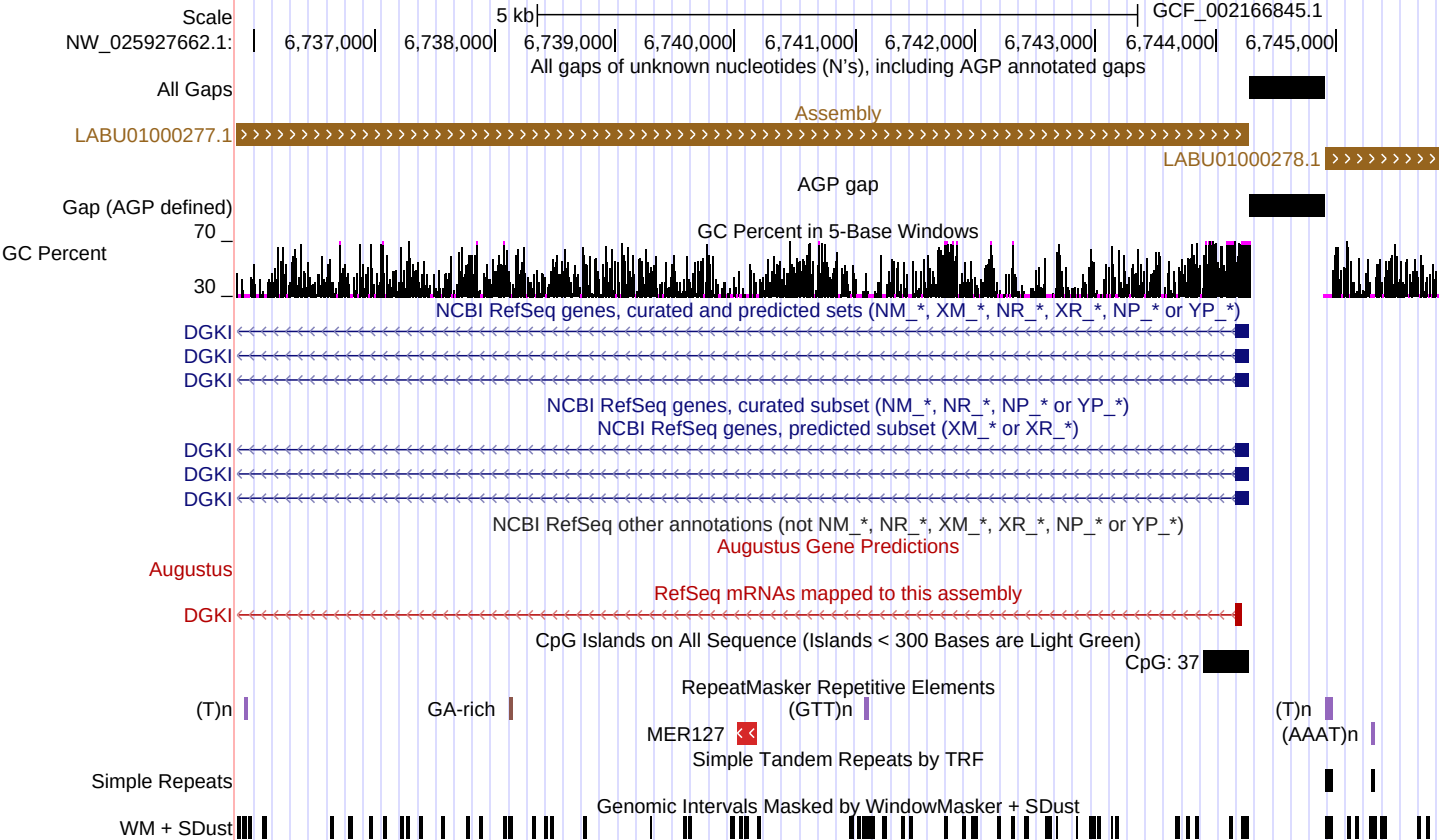
Scale
5 kb
GCF_002166845.1
NW_025927662.1:
6,737,000
6,738,000
6,739,000
6,740,000
6,741,000
6,742,000
6,743,000
6,744,000
6,745,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
LABU01000277.1
LABU01000278.1
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
GC Percent
70
30
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
DGKI
DGKI
DGKI
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
DGKI
DGKI
DGKI
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
DGKI
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
CpG: 37
RepeatMasker Repetitive Elements
(T)n
GA-rich
(GTT)n
(T)n
MER127
(AAAT)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust