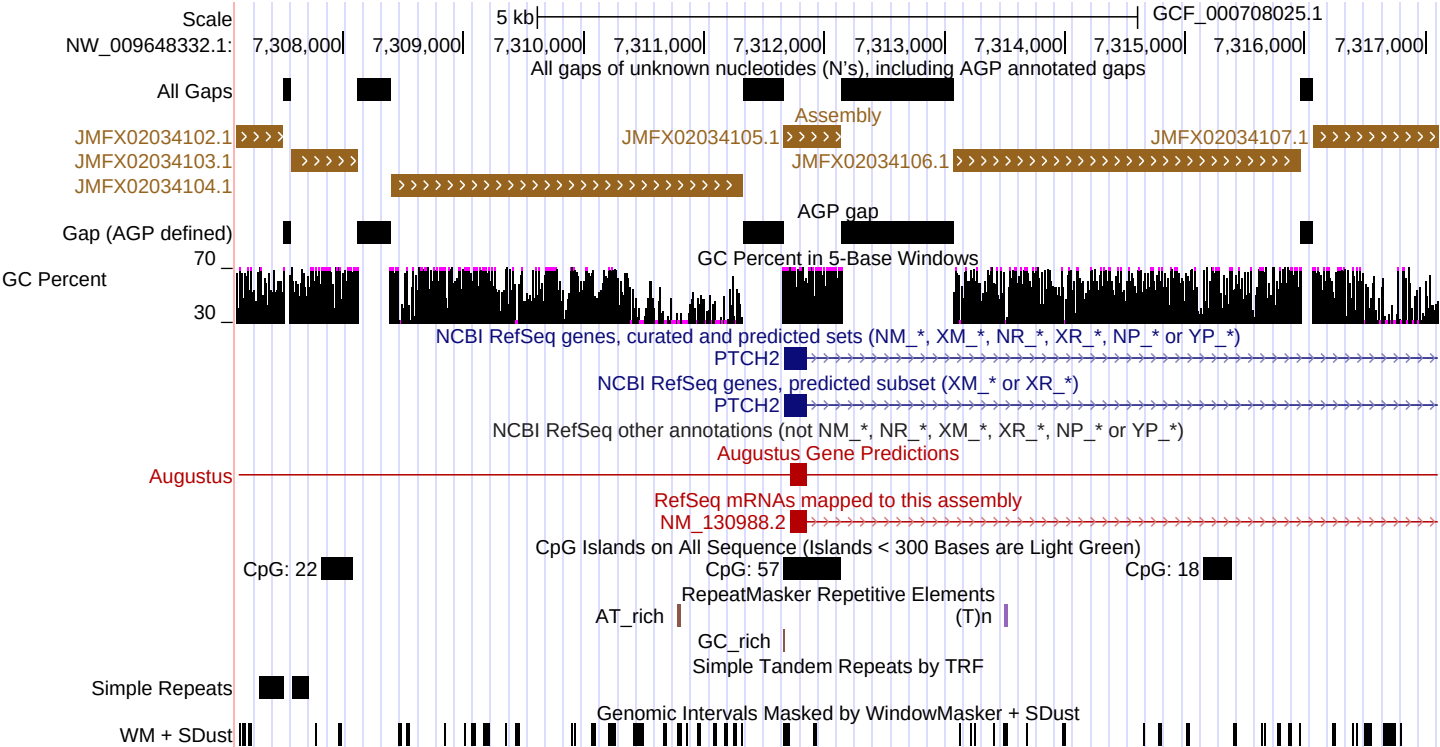
Scale
5 kb
GCF_000708025.1
NW_009648332.1:
7,308,000
7,309,000
7,310,000
7,311,000
7,312,000
7,313,000
7,314,000
7,315,000
7,316,000
7,317,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JMFX02034102.1
JMFX02034105.1
JMFX02034107.1
JMFX02034103.1
JMFX02034106.1
JMFX02034104.1
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_*)
PTCH2
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
PTCH2
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
NM_130988.2
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
CpG: 22
CpG: 57
CpG: 18
RepeatMasker Repetitive Elements
AT_rich
(T)n
GC_rich
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust