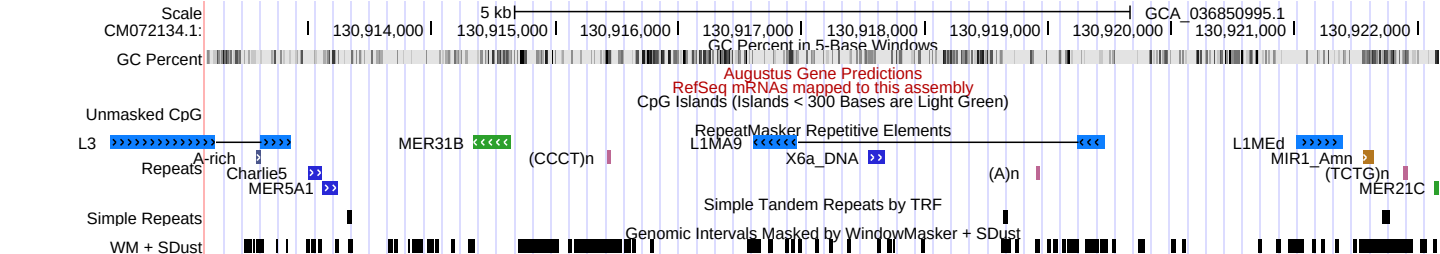
Scale
5 kb
GCA_036850995.1
CM072134.1:
130,914,000
130,915,000
130,916,000
130,917,000
130,918,000
130,919,000
130,920,000
130,921,000
130,922,000
GC Percent in 5-Base Windows
GC Percent
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
L3
MER31B
L1MA9
L1MEd
A-rich
(CCCT)n
X6a_DNA
MIR1_Amn
Repeats
Charlie5
(A)n
(TCTG)n
MER5A1
MER21C
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust