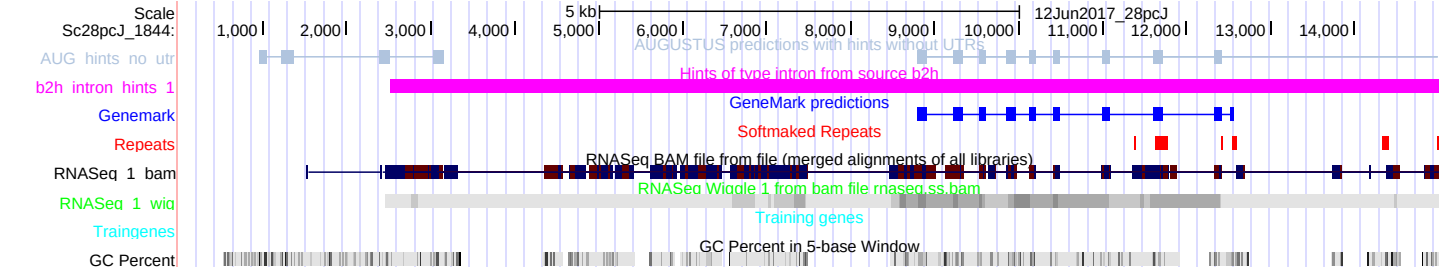
Scale
Sc28pcJ_1844:
5 kb
12Jun2017_28pcJ
1,000
2,000
3,000
4,000
5,000
6,000
7,000
8,000
9,000
10,000
11,000
12,000
13,000
14,000
AUG hints no utr
AUGUSTUS predictions with hints without UTRs
b2h intron hints 1
Hints of type intron from source b2h
Genemark
GeneMark predictions
Repeats
Softmaked Repeats
RNASeq 1 bam
RNASeq BAM file from file (merged alignments of all libraries)
RNASeq 1 wig
RNASeq Wiggle 1 from bam file rnaseq.ss.bam
Traingenes
Training genes
GC Percent
GC Percent in 5-base Window