Scale
2 kb
dm6
chr4:
307,500
308,000
308,500
309,000
309,500
310,000
310,500
311,000
311,500
FBS predictions in Drosophila melanogaster (dm6) for all profiles in the JASPAR CORE insects collection (2020) (11981 items filtered ou
FBS predictions in Drosophila melanogaster (dm6) for all profiles in the JASPAR CORE insects collection (2022) (12108 items filtered ou
Fruitfly MNR/DNR/TNR repeats plus hinges (breakpoints)
Gap Locations
Gap
FlyBase Protein-Coding Genes
Rad23-RA
Rad23-RC
Rad23-RB
Zip102B-RB
Zip102B-RD
Zip102B-RF
Zip102B-RE
FlyBase Pseudogenes
FlyBase Non-protein Coding Genes
miRNA
ncRNA
tRNA
miscRNA
Repeating Elements by RepeatMasker
RepeatMasker