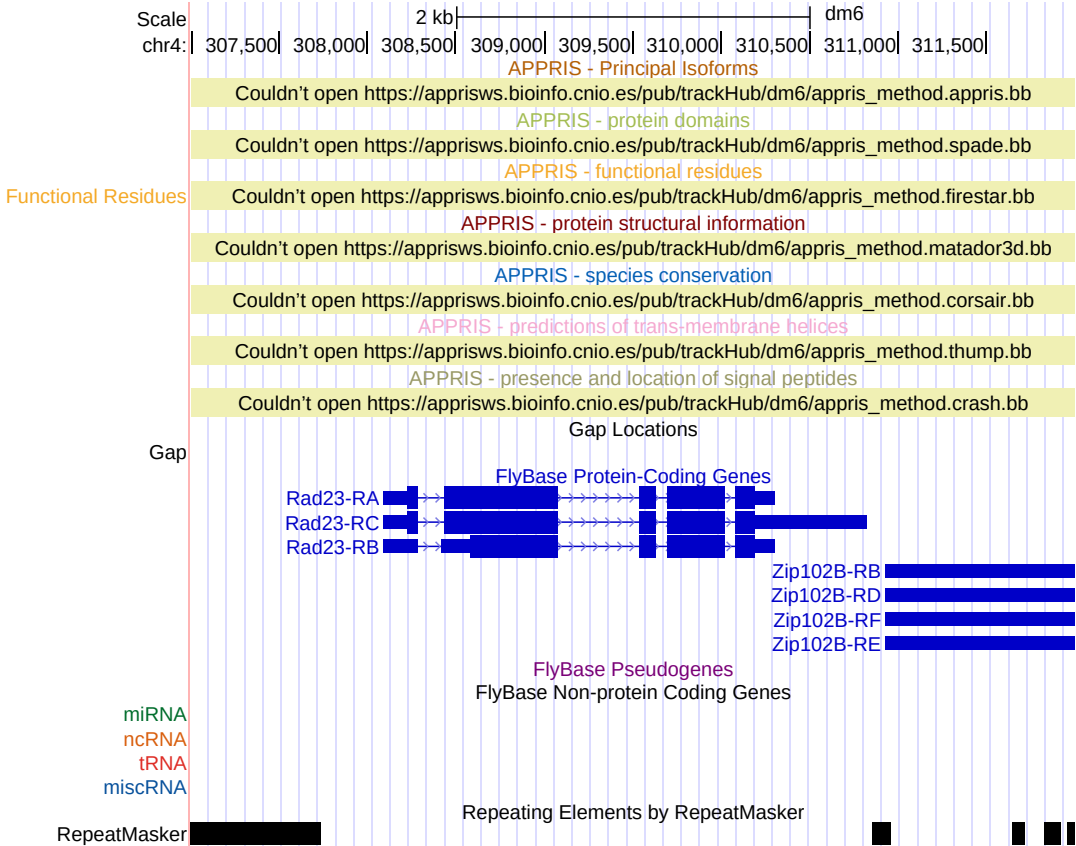
Scale
chr4:
2 kb
dm6
307,500
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APPRIS - Principal Isoforms
Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.appris.bb
APPRIS - protein domains
Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.spade.bb
APPRIS - functional residues
Functional Residues
Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.firestar.bb
APPRIS - protein structural information
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APPRIS - species conservation
Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.corsair.bb
APPRIS - predictions of trans-membrane helices
Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.thump.bb
APPRIS - presence and location of signal peptides
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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