

Scale chr4: | 307,500 | 308,000 | 308,500 | 309,000 | 309,500 | 310,000 | 310,500 | 311,000 | 311,500 |

2 kb

dm6

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

Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.firestar.bb

APPRIS - protein structural information

Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.matador3d.bb

APPRIS - species conservation

Species Conserv

Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.corsair.bb

APPRIS - predictions of trans-membrane helices

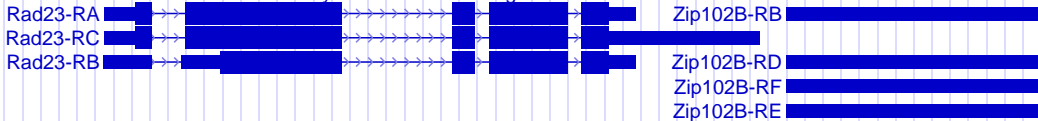
Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.thump.bb

APPRIS - presence and location of signal peptides

Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.crash.bb

Gap Locations

FlyBase Protein-Coding Genes



FlyBase Pseudogenes

FlyBase Non-protein Coding Genes

miRNA
ncRNA
tRNA
miscRNA

Repeating Elements by RepeatMasker

RepeatMasker

