

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

Functional Residues

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

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

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