

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](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](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](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](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](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](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](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