

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

APPRIIS - Principal Isoforms
 Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.appris.bb

APPRIIS - protein domains
 Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.spade.bb

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

APPRIIS - protein structural information
 Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.matador3d.bb

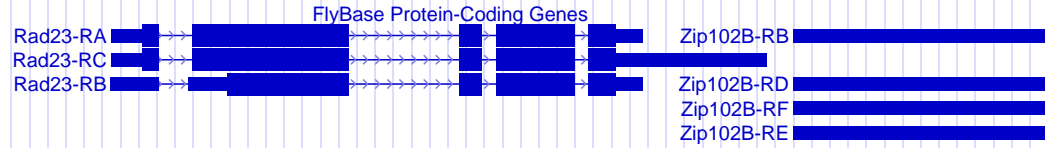
APPRIIS - species conservation
 Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.corsair.bb

APPRIIS - predictions of trans-membrane helices
 Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.thump.bb

APPRIIS - presence and location of signal peptides
 Couldn't open https://apprisws.bioinfo.cnio.es/pub/trackHub/dm6/appris_method.crash.bb

Gap

Gap Locations



miRNA
ncRNA
tRNA
miscRNA

FlyBase Pseudogenes
FlyBase Non-protein Coding Genes

Repeating Elements by RepeatMasker

RepeatMasker

