

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

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APPRIS - protein domains

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APPRIS - functional residues

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APPRIS - protein structural information

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APPRIS - species conservation

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APPRIS - predictions of trans-membrane helices

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APPRIS - presence and location of signal peptides

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Functional Residues

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