

Original Genome:

D. melanogaster

Original Assembly:

Apr. 2006 (BDGP R5/dm3)

New Genome:

D. melanogaster

New Assembly:

Aug. 2014 (BDGP Release 6 + ISO1 MT/dm6)

Minimum ratio of bases that must remap:

0.95

BED 4 to BED 6 Options

Allow multiple output regions:

Minimum hit size in query:

0

Minimum chain size in target:

0

BED 12 Options

Min ratio of alignment blocks or exons that must map:

1

If thickStart/thickEnd is not mapped, use the closest mapped base:

Paste in data ([BED](#) or chrN:start-end formats):

chrX 2684762 2687041

Submit

Clear

Or upload data from a file ([BED](#) or chrN:start-end in plain text format):

Choose File No file chosen

Submit File