

Resource Summary Report

Generated by [dkNET](#) on Aug 23, 2026

[y\[1\] v\[1\]; P{y\[+t7.7\] v\[+t1.8\]=TRiP.GL00184}attP2](#)

RRID:BDSC_35283

Type: Organism

Proper Citation

RRID:BDSC_35283

Organism Information

URL: <https://n2t.net/bdsc:35283>

Proper Citation: RRID:BDSC_35283

Description: Drosophila melanogaster with name y[1] v[1]; P{y[+t7.7] v[+t1.8]=TRiP.GL00184}attP2 from BDSC.

Species: Drosophila melanogaster

Notes: May be segregating TM3, Sb[1] and/or sc[*] and sev[21]. Donor: Transgenic RNAi Project

Affected Gene: Mps1, UAS, sc, sev, v, y

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 35283

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_35283, BL35283

Organism Name: y[1] v[1]; P{y[+t7.7] v[+t1.8]=TRiP.GL00184}attP2

Record Creation Time: 20240911T222602+0000

Record Last Update: 20260815T191755+0000

Ratings and Alerts

No rating or validation information has been found for y[1] v[1]; P{y[+7.7] v[+1.8]=TRiP.GL00184}attP2.

No alerts have been found for y[1] v[1]; P{y[+7.7] v[+1.8]=TRiP.GL00184}attP2.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at [dkNET](#) .

Gohel P, et al. (2026) A non-transcriptional mitotic function of POU/Oct factors ensures spindle assembly and chromosome segregation. *Journal of cell science*, 139(5).

Gilliland WD, et al. (2024) A Cytological F1 RNAi Screen for Defects in *Drosophila melanogaster* Female Meiosis. *bioRxiv : the preprint server for biology*.

Br??s R, et al. (2021) Aneuploidy facilitates dysplastic and tumorigenic phenotypes in the *Drosophila* gut. *Biology open*, 10(11).

Wang LI, et al. (2019) Sister centromere fusion during meiosis I depends on maintaining cohesins and destabilizing microtubule attachments. *PLoS genetics*, 15(5), e1008072.

Silva RD, et al. (2018) Absence of the Spindle Assembly Checkpoint Restores Mitotic Fidelity upon Loss of Sister Chromatid Cohesion. *Current biology : CB*, 28(17), 2837.