

Resource Summary Report

Generated by [NIF](#) 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 [NIF](#) .

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.