

Resource Summary Report

Generated by [dkNET](#) on Aug 31, 2026

[y\[1\] v\[1\]; P{y\[+t7.7\] v\[+t1.8\]=TRiP.GL01544}attP40](#)

RRID:BDSC_43200

Type: Organism

Proper Citation

RRID:BDSC_43200

Organism Information

URL: <https://n2t.net/bdsc:43200>

Proper Citation: RRID:BDSC_43200

Description: Drosophila melanogaster with name y[1] v[1]; P{y[+t7.7] v[+t1.8]=TRiP.GL01544}attP40 from BDSC.

Species: Drosophila melanogaster

Notes: May be segregating CyO Donor: Transgenic RNAi Project

Affected Gene: csul, UAS, v, y

Genomic Alteration: Chromosome 1, Chromosome 2

Catalog Number: 43200

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_43200, BL43200

Organism Name: y[1] v[1]; P{y[+t7.7] v[+t1.8]=TRiP.GL01544}attP40

Record Creation Time: 20240911T222711+0000

Record Last Update: 20260829T190230+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.GL01544}attP40.

No alerts have been found for y[1] v[1]; P{y[+7.7] v[+1.8]=TRiP.GL01544}attP40.

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](#) .

Akkouche A, et al. (2025) Binding of heterochromatin protein Rhino to a subset of piRNA clusters depends on a combination of two histone marks. Nature structural & molecular biology, 32(8), 1517.

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

Curnutte HA, et al. (2023) Proteins rather than mRNAs regulate nucleation and persistence of Oskar germ granules in Drosophila. Cell reports, 42(7), 112723.

Vrettos N, et al. (2021) Modulation of Aub-TDRD interactions elucidates piRNA amplification and germline formation. Life science alliance, 4(3).

Lo Piccolo L, et al. (2019) The lncRNA hsr?? regulates arginine dimethylation of human FUS to cause its proteasomal degradation in Drosophila. Journal of cell science, 132(20).