

# Resource Summary Report

Generated by [NIF](#) on Sep 20, 2026

[y\[1\] sc\[\\*\] v\[1\] sev\[21\]; P{y\[+t7.7\] v\[+t1.8\]=TRiP.GL01004}attP40](#)

RRID:BDSC\_36799

Type: Organism

## Proper Citation

RRID:BDSC\_36799

## Organism Information

**URL:** <https://n2t.net/bdsc:36799>

**Proper Citation:** RRID:BDSC\_36799

**Description:** Drosophila melanogaster with name y[1] sc[\*] v[1] sev[21]; P{y[+t7.7] v[+t1.8]=TRiP.GL01004}attP40 from BDSC.

**Species:** Drosophila melanogaster

**Notes:** May be segregating CyO. Donor: Transgenic RNAi Project

**Affected Gene:** Hcf, UAS, sc, sev, v, y

**Genomic Alteration:** Chromosome 1, Chromosome 2

**Catalog Number:** 36799

**Database:** Bloomington Drosophila Stock Center (BDSC)

**Database Abbreviation:** BDSC

**Availability:** available

**Alternate IDs:** BDSC\_36799, BDSC:36799, BL36799

**Organism Name:** y[1] sc[\*] v[1] sev[21]; P{y[+t7.7] v[+t1.8]=TRiP.GL01004}attP40

**Record Creation Time:** 20240911T222616+0000

**Record Last Update:** 20260919T203049+0000

## Ratings and Alerts

No rating or validation information has been found for y[1] sc[\*] v[1] sev[21]; P{y[+t7.7] v[+t1.8]=TRiP.GL01004}attP40.

No alerts have been found for y[1] sc[\*] v[1] sev[21]; P{y[+t7.7] v[+t1.8]=TRiP.GL01004}attP40.

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## Data and Source Information

**Source:** [Integrated Animals](#)

**Source Database:** Bloomington Drosophila Stock Center (BDSC)

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## Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Malin JA, et al. (2024) Spatial patterning controls neuron numbers in the Drosophila visual system. *Developmental cell*, 59(9), 1132.

Kakemura B, et al. (2024) Blockade of Crk eliminates Yki/YAP-activated tumors via JNK-mediated apoptosis in Drosophila. *Communications biology*, 7(1), 1196.

Huang W, et al. (2022) Lpt, trr, and Hcf regulate histone mono- and dimethylation that are essential for Drosophila heart development. *Developmental biology*, 490, 53.

Ibarra-Morales D, et al. (2021) Histone variant H2A.Z regulates zygotic genome activation. *Nature communications*, 12(1), 7002.