

Resource Summary Report

Generated by [dkNET](#) on Aug 14, 2026

[y\[1\] v\[1\]; P{y\[+t7.7\] v\[+t1.8\]=TRiP.JF02515}attP2](#)

RRID:BDSC_29353

Type: Organism

Proper Citation

RRID:BDSC_29353

Organism Information

URL: <https://n2t.net/bdsc:29353>

Proper Citation: RRID:BDSC_29353

Description: Drosophila melanogaster with name y[1] v[1]; P{y[+t7.7] v[+t1.8]=TRiP.JF02515}attP2 from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Transgenic RNAi Project

Affected Gene: Eip74EF, UAS, v, y

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 29353

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_29353, BL29353

Organism Name: y[1] v[1]; P{y[+t7.7] v[+t1.8]=TRiP.JF02515}attP2

Record Creation Time: 20240911T222504+0000

Record Last Update: 20260808T194703+0000

Ratings and Alerts

No rating or validation information has been found for y[1] v[1]; P{y[+t7.7] v[+t1.8]=TRiP.JF02515}attP2.

No alerts have been found for y[1] v[1]; P{y[+t7.7] v[+t1.8]=TRiP.JF02515}attP2.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Lane EA, et al. (2026) Cholinergic signaling modulates intestinal pathophysiology in a Drosophila model of cystic fibrosis. PLoS genetics, 22(2), e1012048.

Lane E, et al. (2025) Cholinergic Signaling Modulates Intestinal Pathophysiology in a Drosophila Model of Cystic Fibrosis. bioRxiv : the preprint server for biology.

Villalobos-Cantor S, et al. (2025) Genetic basis of maneb-induced dopaminergic neurodegeneration in Drosophila. G3 (Bethesda, Md.), 15(9).

Buffry AD, et al. (2023) Characterisation of the role and regulation of Ultrabithorax in sculpting fine-scale leg morphology. Frontiers in cell and developmental biology, 11, 1119221.

Chalmers MR, et al. (2023) Eip74EF is a dominant modifier for ALS-FTD-linked VCPR152H phenotypes in the Drosophila eye model. BMC research notes, 16(1), 30.

Xu K, et al. (2018) Temporospatial induction of homeodomain gene cut dictates natural lineage reprogramming. eLife, 7.