

Resource Summary Report

Generated by [dkNET](#) on Aug 17, 2026

[y\[1\] sc\[*\] v\[1\] sev\[21\]; P{y\[+t7.7\] v\[+t1.8\]=TRiP.HMS00363}attP2](#)

RRID:BDSC_32372

Type: Organism

Proper Citation

RRID:BDSC_32372

Organism Information

URL: <https://n2t.net/bdsc:32372>

Proper Citation: RRID:BDSC_32372

Description: Drosophila melanogaster with name y[1] sc[*] v[1] sev[21]; P{y[+t7.7] v[+t1.8]=TRiP.HMS00363}attP2 from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Transgenic RNAi Project

Affected Gene: Snr1, UAS, sc, sev, v, y

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 32372

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_32372, BL32372

Organism Name: y[1] sc[*] v[1] sev[21]; P{y[+t7.7] v[+t1.8]=TRiP.HMS00363}attP2

Record Creation Time: 20240911T222532+0000

Record Last Update: 20260815T191716+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.HMS00363}attP2.

No alerts have been found for y[1] sc[*] v[1] sev[21]; P{y[+t7.7] v[+t1.8]=TRiP.HMS00363}attP2.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 9 mentions in open access literature.

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

Chang YC, et al. (2024) Identification of secretory autophagy as a mechanism modulating activity-induced synaptic remodeling. *Proceedings of the National Academy of Sciences of the United States of America*, 121(16), e2315958121.

MacPherson RA, et al. (2023) Genetic and Genomic Analyses of *Drosophila melanogaster* Models of Chromatin Modification Disorders. *bioRxiv : the preprint server for biology*.

MacPherson RA, et al. (2023) Genetic and genomic analyses of *Drosophila melanogaster* models of chromatin modification disorders. *Genetics*, 224(4).

Keegan SE, et al. (2023) A chromatin remodelling SWI/SNF subunit, Snr1, regulates neural stem cell determination and differentiation. *Development (Cambridge, England)*, 150(13).

Torres-Campana D, et al. (2022) Three classes of epigenomic regulators converge to hyperactivate the essential maternal gene deadhead within a heterochromatin mini-domain. *PLoS genetics*, 18(1), e1009615.

Molnar C, et al. (2022) Human EWS-FLI protein recapitulates in *Drosophila* the neomorphic functions that induce Ewing sarcoma tumorigenesis. *PNAS nexus*, 1(4), pgac222.

Hu X, et al. (2021) The Osa-Containing SWI/SNF Chromatin-Remodeling Complex Is Required in the Germline Differentiation Niche for Germline Stem Cell Progeny Differentiation. *Genes*, 12(3).

Giraud G, et al. (2021) Developmental Robustness: The Haltere Case in *Drosophila*. *Frontiers in cell and developmental biology*, 9, 713282.

Pillidge Z, et al. (2019) SWI/SNF chromatin remodeling controls Notch-responsive enhancer accessibility. *EMBO reports*, 20(5).