

Resource Summary Report

Generated by [NIF](#) on Sep 22, 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, 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: 20260919T202953+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 [NIF](#) .

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).