

Resource Summary Report

Generated by [dkNET](#) on Aug 16, 2026

[y\[1\] v\[1\]; P{y\[+t7.7\]
v\[+t1.8\]=TRiP.HMS01570}attP2/TM6C, Sb\[1\] Tb\[1\]](#)

RRID:BDSC_36682
Type: Organism

Proper Citation

RRID:BDSC_36682

Organism Information

URL: <https://n2t.net/bdsc:36682>

Proper Citation: RRID:BDSC_36682

Description: Drosophila melanogaster with name y[1] v[1]; P{y[+t7.7]
v[+t1.8]=TRiP.HMS01570}attP2/TM6C, Sb[1] Tb[1] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Transgenic RNAi Project

Affected Gene: nej, UAS, Sb, Tb, v, y

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 36682

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_36682, BL36682

Organism Name: y[1] v[1]; P{y[+t7.7] v[+t1.8]=TRiP.HMS01570}attP2/TM6C, Sb[1] Tb[1]

Record Creation Time: 20240911T222615+0000

Record Last Update: 20260815T191805+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.HMS01570}attP2/TM6C, Sb[1] Tb[1].

No alerts have been found for y[1] v[1]; P{y[+t7.7] v[+t1.8]=TRiP.HMS01570}attP2/TM6C, Sb[1] Tb[1].

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](#) .

Cho CY, et al. (2025) A mitotic bookmark coordinates transcription and replication in Drosophila embryos. Nucleic acids research, 53(22).

Cho CY, et al. (2025) A mitotic bookmark coordinates transcription and replication. bioRxiv : the preprint server for biology.

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.

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.

Ciabrelli F, et al. (2023) CBP and Gcn5 drive zygotic genome activation independently of their catalytic activity. Science advances, 9(16), eadf2687.

Janssens DH, et al. (2017) An Hdac1/Rpd3-Poised Circuit Balances Continual Self-Renewal and Rapid Restriction of Developmental Potential during Asymmetric Stem Cell Division. Developmental cell, 40(4), 367.