

Resource Summary Report

Generated by T1D on Sep 18, 2026

P{w[+mC]=UAS-pros.L}L3a, w[*]

RRID:BDSC_32244

Type: Organism

Proper Citation

RRID:BDSC_32244

Organism Information

URL: <https://n2t.net/bdsc:32244>

Proper Citation: RRID:BDSC_32244

Description: Drosophila melanogaster with name P{w[+mC]=UAS-pros.L}L3a, w[*] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Joy Tea & Liqun Luo, Stanford University; Donor's Source: Chris Doe, University of Oregon

Affected Gene: pros, UAS, w

Genomic Alteration: Chromosome 1

Catalog Number: 32244

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_32244, BDSC:32244, BL32244

Organism Name: P{w[+mC]=UAS-pros.L}L3a, w[*]

Record Creation Time: 20240911T222531+0000

Record Last Update: 20260912T203611+0000

Ratings and Alerts

No rating or validation information has been found for P{w[+mC]=UAS-pros.L}L3a, w[*].

No alerts have been found for P{w[+mC]=UAS-pros.L}L3a, w[*].

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 [T1D](#) .

Hu Y, et al. (2025) The TRIM-NHL RNA-binding protein MEI-P26 modulates the size of Drosophila Type I neuroblast lineages. *Genetics*, 229(3).

Guo X, et al. (2024) Cell-fate conversion of intestinal cells in adult Drosophila midgut by depleting a single transcription factor. *Nature communications*, 15(1), 2656.

Guo X, et al. (2022) A hierarchical transcription factor cascade regulates enteroendocrine cell diversity and plasticity in Drosophila. *Nature communications*, 13(1), 6525.

Han MH, et al. (2020) NF- κ B disinhibition contributes to dendrite defects in fly models of neurodegenerative diseases. *The Journal of cell biology*, 219(12).

Bravo González-Blas C, et al. (2020) Identification of genomic enhancers through spatial integration of single-cell transcriptomics and epigenomics. *Molecular systems biology*, 16(5), e9438.

Alyagor I, et al. (2018) Combining Developmental and Perturbation-Seq Uncovers Transcriptional Modules Orchestrating Neuronal Remodeling. *Developmental cell*, 47(1), 38.