

Resource Summary Report

Generated by [NIF](#) on Aug 11, 2026

[w\[*\]; P{w\[+mC\]=UAS-ttk.p69}2](#)

RRID:BDSC_7361

Type: Organism

Proper Citation

RRID:BDSC_7361

Organism Information

URL: <https://n2t.net/bdsc:7361>

Proper Citation: RRID:BDSC_7361

Description: Drosophila melanogaster with name w[*]; P{w[+mC]=UAS-ttk.p69}2 from BDSC.

Species: Drosophila melanogaster

Notes: Cy- balancer present in most flies. Donor: Jean-Maurice Dura, French Institute of Health and Medical Research

Affected Gene: ttk, UAS, w

Genomic Alteration: Chromosome 1, Chromosome 2

Catalog Number: 7361

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_7361, BL7361

Organism Name: w[*]; P{w[+mC]=UAS-ttk.p69}2

Record Creation Time: 20240911T222208+0000

Record Last Update: 20260808T194329+0000

Ratings and Alerts

No rating or validation information has been found for w[*]; P{w[+mC]=UAS-ttk.p69}2.

No alerts have been found for w[*]; P{w[+mC]=UAS-ttk.p69}2.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

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.

Duan J, et al. (2020) Bab2 Functions as an Ecdysone-Responsive Transcriptional Repressor during Drosophila Development. Cell reports, 32(4), 107972.

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