

Resource Summary Report

Generated by T1D on Sep 6, 2026

w[1118]; Df(4)ED6364,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}Abcd1[ED6364]/l(4)1

RRID:BDSC_9421

Type: Organism

Proper Citation

RRID:BDSC_9421

Organism Information

URL: <https://n2t.net/bdsc:9421>

Proper Citation: RRID:BDSC_9421

Description: Drosophila melanogaster with name w[1118]; Df(4)ED6364, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}Abcd1[ED6364]/l(4)102EFf[1] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: Abcd1, Ank, anne, Arl4, asRNA:CR43957, asRNA:CR45198, CG32006, CG33978, gvl, lncRNA:CR43956, lncRNA:CR46488, pan, RpS3A, snoRNA:CD68, l(4)102EFf, w

Genomic Alteration: Chromosome 1, Chromosome 4

Catalog Number: 9421

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_9421, BDSC:9421, BL9421

Organism Name: w[1118]; Df(4)ED6364, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}Abcd1[ED6364]/l(4)102EFf[1]

Record Creation Time: 20240911T222224+0000

Record Last Update: 20260905T135900+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(4)ED6364, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}Abcd1[ED6364]/l(4)102EFf[1].

No alerts have been found for w[1118]; Df(4)ED6364, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}Abcd1[ED6364]/l(4)102EFf[1].

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Ribeiro V, et al. (2026) Genome-wide screen for deficiencies modifying Cyclin G-induced developmental instability in *Drosophila melanogaster*. *Genetics*, 232(3).

Sawyer JK, et al. (2020) Exploiting codon usage identifies intensity-specific modifiers of Ras/MAPK signaling in vivo. *PLoS genetics*, 16(12), e1009228.

Li X, et al. (2020) The Mediator CDK8-Cyclin C complex modulates Dpp signaling in *Drosophila* by stimulating Mad-dependent transcription. *PLoS genetics*, 16(5), e1008832.

Hope KA, et al. (2019) A genome-wide enhancer/suppressor screen for Dube3a interacting genes in *Drosophila melanogaster*. *Scientific reports*, 9(1), 2382.