

Resource Summary Report

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[Df\(1\)ED7331, w\[1118\]](#)

[P{w\[+mW.Scer\FRT.hs3\]=3'.RS5+3.3'}ED7331/FM7h](#)

RRID:BDSC_9219

Type: Organism

Proper Citation

RRID:BDSC_9219

Organism Information

URL: <https://n2t.net/bdsc:9219>

Proper Citation: RRID:BDSC_9219

Description: Drosophila melanogaster with name Df(1)ED7331, w[1118] P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7331/FM7h from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: Ac13E, Ahcy, Alg14, Alp11, Arp6, asRNA:CR43299, betaNACtes5, CCT6, cerv, CG11655, CG11679, CG12379, CG12708, CG15599, CG15601, CG15602, CG15641, CG15642, CG15643, CG15646, CG33172, CG42299, CG42300, CG43672, CG43673, CG6294, CG6299, CG6324, CG6340, CG7860, CG7872, CG8097, CG8117, CG8119, CG8128, CG8191, CG8206, CG8260, CG8509, CG8944, CG8952, CG9213, CG9215, CG9240, CG9281, Chsy, CR44383, Efhc1.1, Gmap, Graf, Grip128, Hap40, HUWE1, lncRNA:CR43132, lncRNA:CR44353, lncRNA:CR44354, lncRNA:CR44384, lncRNA:CR44423, lncRNA:CR44888, lncRNA:CR44893, lncRNA:CR45523, lncRNA:CR45524, lncRNA:CR45526, MagR, mh, mir-12, mir-283, mir-304, mRpl3, mRps30, Mvd, Paf-AHalpha, PGRP-LE, Pis, Polr3A, Rhp, sd, shtd, sog, sordd1, sordd2, w

Genomic Alteration: Chromosome 1

Catalog Number: 9219

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_9219, BL9219

Organism Name: Df(1)ED7331, w[1118]
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7331/FM7h

Record Creation Time: 20240911T222223+0000

Record Last Update: 20260801T194705+0000

Ratings and Alerts

No rating or validation information has been found for Df(1)ED7331, w[1118]
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7331/FM7h.

No alerts have been found for Df(1)ED7331, w[1118]
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7331/FM7h.

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 [NIF](#) .

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

Messer CL, et al. (2025) A deficiency screen of the X chromosome for Rap1 GTPase dominant interacting genes in *Drosophila* border cell migration. *G3* (Bethesda, Md.), 15(5).

Pischedda A, et al. (2020) The Loci of Behavioral Evolution: Evidence That Fas2 and tilB Underlie Differences in Pupation Site Choice Behavior between *Drosophila melanogaster* and *D.??simulans*. *Molecular biology and evolution*, 37(3), 864.

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.

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

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