

# Resource Summary Report

Generated by ASWG on Aug 4, 2026

Df(1)ED411,  
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED411  
w[1118]/FM7j, Bar[1]

RRID:BDSC\_8031

Type: Organism

## Proper Citation

RRID:BDSC\_8031

## Organism Information

**URL:** <https://n2t.net/bdsc:8031>

**Proper Citation:** RRID:BDSC\_8031

**Description:** Drosophila melanogaster with name Df(1)ED411, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED411 w[1118]/FM7j, Bar[1] from BDSC.

**Species:** Drosophila melanogaster

**Notes:** Donor: DrosDel Project

**Affected Gene:** Bar, AANATL7, boi, brv3, CG13760, egh, eIF3g1, Klp3A, mir-4955, Pdfr, Pi4KIIIalpha, Seipin, sgg, Smyd3, trol, Tsp3A, Vha36-3, wds, Zw10, w

**Genomic Alteration:** Chromosome 1

**Catalog Number:** 8031

**Database:** Bloomington Drosophila Stock Center (BDSC)

**Database Abbreviation:** BDSC

**Availability:** available

**Alternate IDs:** BDSC\_8031, BL8031

**Organism Name:** Df(1)ED411, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED411 w[1118]/FM7j, Bar[1]

**Record Creation Time:** 20240911T222213+0000

**Record Last Update:** 20260801T194644+0000

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## Ratings and Alerts

No rating or validation information has been found for Df(1)ED411, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED411 w[1118]/FM7j, Bar[1].

No alerts have been found for Df(1)ED411, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED411 w[1118]/FM7j, Bar[1].

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## Data and Source Information

**Source:** [Integrated Animals](#)

**Source Database:** Bloomington Drosophila Stock Center (BDSC)

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## Usage and Citation Metrics

We found 6 mentions in open access literature.

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

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).

Guss EJ, et al. (2023) Loss of the extracellular matrix protein Perlecan disrupts axonal and synaptic stability during Drosophila development. eLife, 12.

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.

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.

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.