

Resource Summary Report

Generated by [ASWG](#) on Aug 24, 2026

[Df\(1\)ED6565,
P{w\[+mW.Scer\FRT.hs3\]=3'.RS5+3.3'}ED6565
w\[1118\]/FM7h](#)

RRID:BDSC_9299

Type: Organism

Proper Citation

RRID:BDSC_9299

Organism Information

URL: <https://n2t.net/bdsc:9299>

Proper Citation: RRID:BDSC_9299

Description: Drosophila melanogaster with name Df(1)ED6565, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6565 w[1118]/FM7h from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: Actn, arm, asRNA:CR43637, asRNA:CR43864, asRNA:CR44381, asRNA:CR44382, asRNA:CR44690, asRNA:CR44896, asRNA:CR45479, asRNA:CR45929, bcn92, CG14052, CG14053, CG14054, CG16903, CG17776, CG17777, CG2865, CG2924, CG3071, CG3078, CG3091, CG3191, CG32801, CG3457, CG3587, CG3630, CG3857, CG4025, CG4194, CG4281, CG4313, CG4325, CR34672, CR46447, crn, csw, Cyp4ae1, Cyp4d1, Cyp4d14, Cyp4d2, D2hgdh, east, Edem1, eIF2Bepsilon, FarO, fs(1)K10, Grp170, Hr4, kz, Ldaf1, lncRNA:CR45478, lncRNA:CR45480, Mct1, Mettl1, mip130, mir-2496, mir-3641, mir-4954, moody, ND-B14.5A, Nmd3, Ocr1, Or2a, pcx, Pgd, ph-d, ph-p, PIG-K, pn, Raf, Rbcn-3B, Rtca, Sik2, SmydA-8, temp, Unc-76, unmet, usp, Vinc, Vml, wapl, w

Genomic Alteration: Chromosome 1

Catalog Number: 9299

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_9299, BL9299

Organism Name: Df(1)ED6565, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6565 w[1118]/FM7h

Record Creation Time: 20240911T222223+0000

Record Last Update: 20260815T191316+0000

Ratings and Alerts

No rating or validation information has been found for Df(1)ED6565, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6565 w[1118]/FM7h.

No alerts have been found for Df(1)ED6565, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6565 w[1118]/FM7h.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

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.

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.

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

Komori H, et al. (2018) Multilayered gene control drives timely exit from the stem cell state in uncommitted progenitors during *Drosophila* asymmetric neural stem cell division. *Genes & development*, 32(23-24), 1550.