

Resource Summary Report

Generated by [kravitz2](#) on Sep 16, 2026

Df(1)ED7005, w[1118]

P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7005/FM7h

RRID:BDSC_9153

Type: Organism

Proper Citation

RRID:BDSC_9153

Organism Information

URL: <https://n2t.net/bdsc:9153>

Proper Citation: RRID:BDSC_9153

Description: Drosophila melanogaster with name Df(1)ED7005, w[1118] P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7005/FM7h from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: alpha-Man-Ia, asRNA:CR43959, asRNA:CR44894, asRNA:CR45629, ATPsyndelta, CG12640, CG12643, CG12645, CG15296, CG15306, CG15308, CG17841, CG2887, CG2889, CG2898, CG2909, CG2961, CG2962, CG2972, CG32679, CG32681, CG32683, CG32686, CG32687, CG34408, CG43740, CG43902, CR34674, CR34675, flw, Gip, Hk, l(1)G0289, lncRNA:CR32690, lncRNA:CR43899, Naxe, Neb-cGP, nocte, Or9a, PPP4R2r, Psf3, RabX2, RhoU, snoRNA:CD11c, spri, Ypel, ZAP3, w

Genomic Alteration: Chromosome 1

Catalog Number: 9153

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_9153, BDSC:9153, BL9153

Organism Name: Df(1)ED7005, w[1118]

P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7005/FM7h

Record Creation Time: 20240911T222222+0000

Record Last Update: 20260912T203154+0000

Ratings and Alerts

No rating or validation information has been found for Df(1)ED7005, w[1118]
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7005/FM7h.

No alerts have been found for Df(1)ED7005, w[1118]
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7005/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 [kravitz2](#) .

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.

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.