

Resource Summary Report

Generated by [ASWG](#) on Aug 7, 2026

[w\[1118\]; Df\(2L\)ED385,
P{w\[+mW.Scer\FRT.hs3\]=3'.RS5+3.3'}ED385/SM6a](#)

RRID:BDSC_9341

Type: Organism

Proper Citation

RRID:BDSC_9341

Organism Information

URL: <https://n2t.net/bdsc:9341>

Proper Citation: RRID:BDSC_9341

Description: Drosophila melanogaster with name w[1118]; Df(2L)ED385, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED385/SM6a from BDSC.

Species: Drosophila melanogaster

Notes: May be segregating y[1]. Donor: DrosDel Project

Affected Gene: AANATL2, Acox3, Arpc4, asRNA:CR43866, B9d2, CG13982, CG13983, CG13984, CG13989, CG13991, CG13994, CG13995, CG13996, CG31639, CG31642, CG31643, CG34179, CG34180, CG42369, CG43922, CG45075, CG9098, CG9107, CG9109, CG9117, CG9135, CG9147, CG9150, CG9154, CG9175, CG9222, CG9497, CG9498, CG9500, chic, Cpr, Ddr, DIP-epsilon, eIF4A, Fbw5, Fic, frj, Gal, H2.0, ifc, IPIP, Kdm5, Kr-h1, Kr-h2, lncRNA:CR43801, lncRNA:CR44773, lncRNA:CR44777, lncRNA:CR44778, lncRNA:CR44823, lncRNA:CR44824, lncRNA:CR45719, lncRNA:CR46410, lncRNA:CR9162, mir-966, mtm, Muc26B, ND-51, Nepl4, Nepl5, Nepl6, PDZ-GEF, Pez, Pfas, Pfdn1, ppk14, ppk7, Rchy1, retm, slam, slmo, smal, stai, tctn, Tig, Tsp26A, Ugt37B1, WDR79, w

Genomic Alteration: Chromosome 1, Chromosome 2

Catalog Number: 9341

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_9341, BL9341

Organism Name: w[1118]; Df(2L)ED385,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED385/SM6a

Record Creation Time: 20240911T222224+0000

Record Last Update: 20260801T194702+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(2L)ED385,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED385/SM6a.

No alerts have been found for w[1118]; Df(2L)ED385,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED385/SM6a.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 8 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).

Marion SB, et al. (2026) Transposable elements contribute substantially to naturally occurring genetic lethality in *Drosophila melanogaster*. *PLoS biology*, 24(3), e3003467.

Deger JM, et al. (2025) Revealing the nervous system requirements of Alzheimer's disease risk genes in *Drosophila*. *bioRxiv* : the preprint server for biology.

Deger JM, et al. (2025) Revealing the nervous system requirements of Alzheimer disease risk genes in *Drosophila*. *American journal of human genetics*, 112(12), 2870.

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.

Mehsen H, et al. (2018) PP2A-B55 promotes nuclear envelope reformation after mitosis in *Drosophila*. *The Journal of cell biology*, 217(12), 4106.