

Resource Summary Report

Generated by T1D on Sep 17, 2026

w[1118]; Df(3R)ED7665,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7665/TM6C,
cu[1] Sb[1]

RRID:BDSC_8685

Type: Organism

Proper Citation

RRID:BDSC_8685

Organism Information

URL: <https://n2t.net/bdsc:8685>

Proper Citation: RRID:BDSC_8685

Description: Drosophila melanogaster with name w[1118]; Df(3R)ED7665, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7665/TM6C, cu[1] Sb[1] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: cu, Alh, alpha-Est1, alpha-Est10, alpha-Est2, alpha-Est3, alpha-Est4, alpha-Est4aPsi, alpha-Est5, alpha-Est6, alpha-Est7, alpha-Est8, alpha-Est9, alphaTub84D, Ar5, ArgRS-m, asRNA:CR42738, asRNA:CR45909, asRNA:CR46128, asRNA:CR46129, CD98hc, CG10029, CG10053, CG10055, CG10086, CG10098, CG10435, CG10445, CG10903, CG10919, CG11035, CG11052, CG11286, CG1142, CG1227, CG1288, CG14598, CG14605, CG14606, CG14607, CG14609, CG14610, CG17816, CG17944, CG18268, CG18744, CG18745, CG18746, CG18747, CG18748, CG2616, CG2641, CG2656, CG2698, CG2993, CG3014, CG31248, CG31473, CG31493, CG31496, CG3223, CG34384, CG42544, CG42650, CG42656, CG43060, CG43061, CG43254, CG43290, CG43618, CG45545, CG7910, CR10032, CR34681, CR42875, CR46300, dsx, Dtw2, e(y)2b, Elmod, Elov17, EMC1, Fer1, gfzf, Gld, Heatr5, ImpE3, Ir84a, kipf, Kmn2, lap, lds, lncRNA:CR43685, lncRNA:CR44260, lncRNA:CR44330, lncRNA:CR44332, lncRNA:CR44333, lncRNA:CR44334, lncRNA:CR44335, lncRNA:CR44337, lncRNA:CR44338, lncRNA:CR44934, lncRNA:CR44935, lncRNA:CR44936, lncRNA:CR45560, lncRNA:CR45906, lncRNA:CR46127, lncRNA:CR46130, lncRNA:CR46142, lncRNA:CR46143, lncRNA:CR46224, lncRNA:CR46225, lncRNA:CR46226, lncRNA:CR46299, lncRNA:TS14, mAChR-B, MAGE, Mics1, mir-4950, Mlp84B, mRpS9, Mst84Da, Mst84Db, Mst84Dc, Mst84Dd,

MstProx, Nlg1, Nlg3, Noc3, nxf4, Obp84a, Os-C, Pbp95, PIG-H, Prat, pyd3, RacGAP84C, Rfk, rn, Rtnl2, sas, Sas-4, scaRNA:MeU2-C41, scaRNA:MeU5-C46, Sfp84E, sgll, snRNA:7SKa, Sp7, Syt4, Taf7, thw, tRNA:Gly-TCC-1-1, tRNA:Gly-TCC-1-2, tRNA:Val-CAC-2-1, tRNA:Val-CAC-2-2, tRNA:Val-CAC-2-3, tRNA:Val-CAC-2-4, Ubc84D, unc-45, wa-cup, wtrw, Sb, w

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 8685

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_8685, BDSC:8685, BL8685

Organism Name: w[1118]; Df(3R)ED7665,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7665/TM6C, cu[1] Sb[1]

Record Creation Time: 20240911T222218+0000

Record Last Update: 20260912T203149+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(3R)ED7665,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7665/TM6C, cu[1] Sb[1].

No alerts have been found for w[1118]; Df(3R)ED7665,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7665/TM6C, cu[1] Sb[1].

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 [T1D](#).

Wang Y, et al. (2023) hkb is required for DIP-?? expression and target recognition in the Drosophila neuromuscular circuit. bioRxiv : the preprint server for biology.

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.

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