

Resource Summary Report

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[w\[1118\]; Df\(3R\)ED6255,
P{w\[+mW.Scer\FRT.hs3\]=3'.RS5+3.3'}ED6255/TM6C,
cu\[1\] Sb\[1\]](#)

RRID:BDSC_9210

Type: Organism

Proper Citation

RRID:BDSC_9210

Organism Information

URL: <https://n2t.net/bdsc:9210>

Proper Citation: RRID:BDSC_9210

Description: Drosophila melanogaster with name w[1118]; Df(3R)ED6255, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6255/TM6C, cu[1] Sb[1] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: cu, asRNA:CR44049, asRNA:CR45209, ball, CAH9, CG14252, CG14253, CG14257, CG14258, CG14259, CG14260, CG14262, CG17191, CG17192, CG17991, CG18766, CG31064, CG31068, CG31075, CG31076, CG31077, CG34292, CG34293, CG42813, CG42814, CG43935, CG46313, CG46339, CG5500, CG5521, CG5815, CG5880, CG5882, CG5909, CG5934, CG5938, CG5984, CG6051, CG6059, CG6271, CG6277, CG6283, CG6295, CG6296, CG6330, Cpr97Ea, Cpr97Eb, CR34689, eater, Ets97D, gb, Gr97a, grass, His2Av, IntS12, Jhbp11, l(3)mbt, Lerp, lncRNA:CR34006, lncRNA:CR46073, lncRNA:CR46085, Men-b, mir-1012, mrt, ms(3)K81, Nep5, Nkap, pll, PolA2, ppk31, Rb97D, ro, RYa-R, Ser, Spag1, SPARC, spn-D, spz, T48, TflIA-L, TI, tRNA:Gln-TTG-1-2, tRNA:Gln-TTG-1-3, Tsp97E, TTLL6B, TwdIC, TwdIT, woc, Sb, w

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 9210

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_9210, BDSC:9210, BL9210

Organism Name: w[1118]; Df(3R)ED6255,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6255/TM6C, cu[1] Sb[1]

Record Creation Time: 20240911T222223+0000

Record Last Update: 20260919T202523+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(3R)ED6255,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6255/TM6C, cu[1] Sb[1].

No alerts have been found for w[1118]; Df(3R)ED6255,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6255/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 8 mentions in open access literature.

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

Ribeiro V, et al. (2026) Genome-wide screen for deficiencies modifying Cyclin G-induced developmental instability in *Drosophila melanogaster*. *Genetics*, 232(3).

Kumar N, et al. (2026) Identification of a cytosol-to-nucleus feedback loop that regulates neuronal microtubule nucleation. *The Journal of cell biology*, 225(3).

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.

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.