

Resource Summary Report

Generated by [NIF](#) on Sep 9, 2026

[w\[1118\]; Df\(3R\)ED6361,
P{w\[+mW.Scer\FRT.hs3\]=3'.RS5+3.3'}ED6361/TM6C,
cu\[1\] Sb\[1\]](#)

RRID:BDSC_24143

Type: Organism

Proper Citation

RRID:BDSC_24143

Organism Information

URL: <https://n2t.net/bdsc:24143>

Proper Citation: RRID:BDSC_24143

Description: Drosophila melanogaster with name w[1118]; Df(3R)ED6361, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6361/TM6C, cu[1] Sb[1] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: cu, Acf, asRNA:CR33945, awd, betaGlu, CG11539, CG11550, CG11563, CG1792, CG1896, CG1910, CG1971, CG2003, CG2126, CG2150, CG2187, CG42233, CG42693, CR46418, CstF50, CycG, eEF1alpha2, faf, FeCH, heph, Ir100a, kek6, krz, lncRNA:CR34046, lncRNA:CR43436, lncRNA:CR44091, lncRNA:CR44263, lncRNA:CR45052, lncRNA:CR45556, lncRNA:CR45564, lncRNA:CR45565, Map205, Mccc1, Med, mir-1014, mod, nero, pasha, Pos, RhoGAP100F, Rift, salt, sip3, Smvt, Tbca, ttk, Zmynd8, Sb, w

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 24143

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_24143, BDSC:24143, BL24143

Organism Name: w[1118]; Df(3R)ED6361,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6361/TM6C, cu[1] Sb[1]

Record Creation Time: 20240911T222415+0000

Record Last Update: 20260905T140135+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(3R)ED6361,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6361/TM6C, cu[1] Sb[1].

No alerts have been found for w[1118]; Df(3R)ED6361,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6361/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 7 mentions in open access literature.

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

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

Ell CM, et al. (2024) A genome-engineered tool set for *Drosophila* TGF- β /BMP signaling studies. *Development (Cambridge, England)*, 151(22).

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

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.