

Resource Summary Report

Generated by [ASWG](#) on Aug 25, 2026

[w\[1118\]; Df\(3R\)ED2,
P{w\[+mW.Scer\FRT.hs3\]=3'.RS5+3.3'}koko\[ED2\]/TM6C,
cu\[1\] Sb\[1\]](#)

RRID:BDSC_6962

Type: Organism

Proper Citation

RRID:BDSC_6962

Organism Information

URL: <https://n2t.net/bdsc:6962>

Proper Citation: RRID:BDSC_6962

Description: Drosophila melanogaster with name w[1118]; Df(3R)ED2, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}koko[ED2]/TM6C, cu[1] Sb[1] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: cu, asRNA:CR45028, ATPsynD, cdi, CG10864, CG14285, CG14286, CG14292, CG14294, CG14297, CG14298, CG14300, CG14301, CG14302, CG14304, CG31122, CG31224, CG31226, CG31229, CG31230, CG31231, CG34282, CG34283, CG34284, CG42613, CG43210, CG43732, CG46318, CG6005, CG6013, CG6026, CG7685, CG7691, CG7694, CG7695, CG7702, CG7706, CG7714, CG7715, CG7720, ChAT, ChT, Cpsf73, CstF64, dind, dnk, EndoA, Epg5, euc, fray, fru, GatA, Gos28, gukh, gwl, hemo, koko, lncRNA:CR45633, lncRNA:CR45996, lncRNA:CR46041, lncRNA:CR46265, Mekk1, Mpc1, mRpl55, Muc91C, NP15.6, Octalpha2R, P5cr, PGS1, Prp18, qin, Rh2, Sgsh, Smu1, snRNP-U1-C, tRNA:Pro-TGG-1-5, Tssk, VACHT, Vti1b, Wdr37, Xrp1, Sb, w

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 6962

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_6962, BL6962

Organism Name: w[1118]; Df(3R)ED2,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}koko[ED2]/TM6C, cu[1] Sb[1]

Record Creation Time: 20240911T222204+0000

Record Last Update: 20260815T191255+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(3R)ED2,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}koko[ED2]/TM6C, cu[1] Sb[1].

No alerts have been found for w[1118]; Df(3R)ED2,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}koko[ED2]/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 [ASWG](#) .

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).

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.