

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

[w\[1118\]; Df\(3R\)ED6316,
P{w\[+mW.Scer\FRT.hs3\]=3'.RS5+3.3'}ED6316/TM6C,
cu\[1\] Sb\[1\]](#)

RRID:BDSC_8925

Type: Organism

Proper Citation

RRID:BDSC_8925

Organism Information

URL: <https://n2t.net/bdsc:8925>

Proper Citation: RRID:BDSC_8925

Description: Drosophila melanogaster with name w[1118]; Df(3R)ED6316, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6316/TM6C, cu[1] Sb[1] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: cu, alph, asRNA:CR46109, ATPsyngamma, Bub3, Cap-D2, CG11470, CG14506, CG15506, CG18731, CG1907, CG2006, CG2310, CG2321, CG31041, CG34296, CG45544, CG45546, CG7567, CG7568, CG7582, CG7601, Cnx99A, Diedel, Diedel2, dmrt99B, Dop1R2, Dr, Dup99B, fig, FipoQ, Gnprnat, Gycalpha99B, IntS11, kay, Kul, lncRNA:CR43613, lncRNA:CR44022, lncRNA:CR44953, lncRNA:CR45547, lncRNA:CR45568, lncRNA:CR45569, lncRNA:CR46108, lncRNA:CR46110, Naa40, ND-20L, NdufAF1, Obp99a, Obp99b, Obp99c, Obp99d, Pcd, ppk19, ppk20, ppk21, ppk30, Ptp99A, Spase12, stg, Trc8, Wdr24, yata, Sb, w

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 8925

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_8925, BL8925

Organism Name: w[1118]; Df(3R)ED6316,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6316/TM6C, cu[1] Sb[1]

Record Creation Time: 20240911T222220+0000

Record Last Update: 20260801T194656+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(3R)ED6316,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6316/TM6C, cu[1] Sb[1].

No alerts have been found for w[1118]; Df(3R)ED6316,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED6316/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 2 mentions in open access literature.

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

Zhang W, et al. (2020) Identification of a genetic network for an ecologically relevant behavioural phenotype in *Drosophila melanogaster*. *Molecular ecology*, 29(3), 502.

Hope KA, et al. (2019) A genome-wide enhancer/suppressor screen for Dube3a interacting genes in *Drosophila melanogaster*. *Scientific reports*, 9(1), 2382.