

Resource Summary Report

Generated by [Kravitz](#) on Sep 14, 2026

w[1118]; Df(3R)Exel8178/TM6B, Tb[+]

RRID:BDSC_7993

Type: Organism

Proper Citation

RRID:BDSC_7993

Organism Information

URL: <https://n2t.net/bdsc:7993>

Proper Citation: RRID:BDSC_7993

Description: Drosophila melanogaster with name w[1118]; Df(3R)Exel8178/TM6B, Tb[+] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Exelixis, Inc.

Affected Gene: asRNA:CR44021, asRNA:CR46093, BRWD3, CG13613, CG13614, CG13615, CG13616, CG13622, CG17780, CG17781, CG17782, CG17784, CG18528, CG31128, CG33339, CG33340, CG33341, CG33342, CG34290, CG42811, CG42812, CG5715, CG5746, CG5762, CG5768, CG6356, CG6432, CG6454, crb, Dis3, DZIP1, Golgin84, jar, Jhbp12, lncRNA:CR43455, lncRNA:CR45649, lncRNA:CR45662, lncRNA:CR45950, mir-1015, mir-1017, Ms, Nab2, nAChRalpha1, nAChRalpha2, nAChRbeta2, Orct, Orct2, Pp1alpha-96A, Rrp5, shps, Uck, Tb, w

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 7993

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_7993, BDSC:7993, BL7993

Organism Name: w[1118]; Df(3R)Exel8178/TM6B, Tb[+]

Record Creation Time: 20240911T222213+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(3R)Exel8178/TM6B, Tb[+].

No alerts have been found for w[1118]; Df(3R)Exel8178/TM6B, Tb[+].

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Deger JM, et al. (2025) Revealing the nervous system requirements of Alzheimer's disease risk genes in Drosophila. bioRxiv : the preprint server for biology.

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.

Lapart JA, et al. (2019) Dzip1 and Fam92 form a ciliary transition zone complex with cell type specific roles in Drosophila. eLife, 8.