

Resource Summary Report

Generated by [SPARC SAWG](#) on Jul 29, 2026

w[1118]; Df(3R)BSC516/TM2

RRID:BDSC_25020

Type: Organism

Proper Citation

RRID:BDSC_25020

Organism Information

URL: <https://n2t.net/bdsc:25020>

Proper Citation: RRID:BDSC_25020

Description: Drosophila melanogaster with name w[1118]; Df(3R)BSC516/TM2 from BDSC.

Species: Drosophila melanogaster

Notes: This Df stock does not share the genetic background of most Exel and BSC deletions. Donor: Kevin Cook, Bloomington Drosophila Stock Center

Affected Gene: asRNA:CR45024, asRNA:CR45800, asRNA:CR46044, att-ORFB, bon, Cdk2, CG10877, CG10887, CG15922, CG15923, CG16953, CG17199, CG17267, CG31195, CG31199, CG31200, CG31205, CG31206, CG34008, CG4000, CG42322, CG4362, CG4367, CG46441, CG5023, CG5180, CG5191, CG5412, CG5466, cic, Cpr92F, CR42786, Dic2, DPCoAC, eIF3g2, EloB, Gfrl, GluRIID, GluRIIE, Gr92a, H, Ir92a, lncRNA:CR42836, lncRNA:CR43452, lncRNA:CR44400, lncRNA:CR44401, lncRNA:CR45023, lncRNA:CR46043, Lrrk, mir-995, MtnB, MtnC, MtnD, MtnE, Nep4, Nlg4, Oamb, Or92a, Orp8, Pi3K92E, Prosalpha4T1, Pus1, Sag1, Septin2, Sirt2, snoRNA:Psi18S-301, Srp14, Srp72, Stat92E, Syp, Tak11, TFAM, Tmlh, tRNA:Leu-TAA-1-1, VGlut2, w

Genomic Alteration: Chromosome 1, Chromosome 3

Catalog Number: 25020

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_25020, BL25020

Organism Name: w[1118]; Df(3R)BSC516/TM2

Record Creation Time: 20240911T222423+0000

Record Last Update: 20260725T195129+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(3R)BSC516/TM2.

No alerts have been found for w[1118]; Df(3R)BSC516/TM2.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Antel M, et al. (2022) Interchromosomal interaction of homologous Stat92E alleles regulates transcriptional switch during stem-cell differentiation. Nature communications, 13(1), 3981.