

Resource Summary Report

Generated by [ASWG](#) on Aug 6, 2026

w[1118]; Df(2R)BSC408/CyO

RRID:BDSC_24912

Type: Organism

Proper Citation

RRID:BDSC_24912

Organism Information

URL: <https://n2t.net/bdsc:24912>

Proper Citation: RRID:BDSC_24912

Description: Drosophila melanogaster with name w[1118]; Df(2R)BSC408/CyO from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Kevin Cook, Bloomington Drosophila Stock Center

Affected Gene: Alp6, asRNA:CR45281, brp, Camta, ced-6, CG13955, CG1814, CG1888, CG33757, CG33758, lncRNA:CR43651, lncRNA:CR45321, mir-14, Not1, Or45b, Pdk, prel, Wnt2, won, won2, w

Genomic Alteration: Chromosome 1, Chromosome 2

Catalog Number: 24912

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_24912, BL24912

Organism Name: w[1118]; Df(2R)BSC408/CyO

Record Creation Time: 20240911T222422+0000

Record Last Update: 20260801T194938+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(2R)BSC408/CyO.

No alerts have been found for w[1118]; Df(2R)BSC408/CyO.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Chen J, et al. (2024) Astrocyte growth is driven by the Tre1/S1pr1 phospholipid-binding G protein-coupled receptor. *Neuron*, 112(1), 93.

Vuilleumier R, et al. (2022) Dichotomous cis-regulatory motifs mediate the maturation of the neuromuscular junction by retrograde BMP signaling. *Nucleic acids research*, 50(17), 9748.

Sato K, et al. (2019) Calmodulin-binding transcription factor shapes the male courtship song in *Drosophila*. *PLoS genetics*, 15(7), e1008309.

Komori H, et al. (2018) Multilayered gene control drives timely exit from the stem cell state in uncommitted progenitors during *Drosophila* asymmetric neural stem cell division. *Genes & development*, 32(23-24), 1550.