

# Resource Summary Report

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## w[1118]; Df(2R)BSC357/CyO

RRID:BDSC\_24381

Type: Organism

### Proper Citation

RRID:BDSC\_24381

### Organism Information

**URL:** <https://n2t.net/bdsc:24381>

**Proper Citation:** RRID:BDSC\_24381

**Description:** Drosophila melanogaster with name w[1118]; Df(2R)BSC357/CyO from BDSC.

**Species:** Drosophila melanogaster

**Notes:** Donor: Kevin Cook, Bloomington Drosophila Stock Center

**Affected Gene:** Achl, Adgf-E, Asx, Blos1, CG10104, CG10139, CG12860, CG12861, CG12862, CG12863, CG12868, CG12869, CG17385, CG30075, CG30076, CG30197, CG30479, CG30480, CG33468, CG33469, CG34184, CG34185, CG34442, CG34443, CG34444, CG43919, Cpr51A, Cpsf160, Dh44-R1, Had2, L, LamC, lncRNA:CR42715, lncRNA:CR44369, lncRNA:CR44462, lncRNA:CR45277, mspo, Mtkl, Obp50a, Obp50b, Obp50c, Obp50d, Obp50e, phyl, Polr1A, PRAS40, scaRNA:PsiU5-44, Sec61beta, Sin1, snoRNA:CD24, snoRNA:CD40, snoRNA:CD42, snoRNA:CD45, snoRNA:CD46, snoRNA:CD53a, snoRNA:CD53b, snoRNA:CD53c, snoRNA:CD54a, snoRNA:CD54b, snoRNA:CD61, Su(var)2-HP2, Targ, Tfb1, tra2, ttv, Uhg5, w

**Genomic Alteration:** Chromosome 1, Chromosome 2

**Catalog Number:** 24381

**Database:** Bloomington Drosophila Stock Center (BDSC)

**Database Abbreviation:** BDSC

**Availability:** available

**Alternate IDs:** BDSC\_24381, BL24381

**Organism Name:** w[1118]; Df(2R)BSC357/CyO

**Record Creation Time:** 20240911T222417+0000

**Record Last Update:** 20260801T194931+0000

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## Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(2R)BSC357/CyO.

No alerts have been found for w[1118]; Df(2R)BSC357/CyO.

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## Data and Source Information

**Source:** [Integrated Animals](#)

**Source Database:** Bloomington Drosophila Stock Center (BDSC)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Patterson M, et al. (2025) The M.3.2 mutation in Drosophila melanogaster mapped as a novel allele of ?? tout-velu. microPublication biology, 2025.

Son W, et al. (2020) The Classic Lobe Eye Phenotype of Drosophila Is Caused by Transposon Insertion-Induced Misexpression of a Zinc-Finger Transcription Factor. Genetics, 216(1), 117.