Resource Summary Report

Generated by NIF on Aug 29, 2026

Df(1)BSC761, w[1118] P+PBac{w[+mC]=XP3.WH3}BSC761/FM7h

RRID:BDSC_26858
Type: Organism

Proper Citation

RRID:BDSC_26858

Organism Information

URL: https://n2t.net/bdsc:26858

Proper Citation: RRID:BDSC_26858

Description: Drosophila melanogaster with name Df(1)BSC761, w[1118] P+PBac{w[+mC]=XP3.WH3}BSC761/FM7h from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Kevin Cook, Bloomington Drosophila Stock Center

Affected Gene: Arp2, CanA-14F, CG13014, CG9902, CG9903, Cnx14D, para, Pp2B-14D, Smc3, Ubc7, w

Genomic Alteration: Chromosome 1

Catalog Number: 26858

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_26858, BL26858

Organism Name: Df(1)BSC761, w[1118] P+PBac{w[+mC]=XP3.WH3}BSC761/FM7h

Record Creation Time: 20240911T222440+0000

Record Last Update: 20260829T185916+0000

Ratings and Alerts

No rating or validation information has been found for Df(1)BSC761, w[1118] P+PBac{w[+mC]=XP3.WH3}BSC761/FM7h.

No alerts have been found for Df(1)BSC761, w[1118] P+PBac{w[+mC]=XP3.WH3}BSC761/FM7h.

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 NIF .

Ribeiro V, et al. (2026) Genome-wide screen for deficiencies modifying Cyclin G-induced developmental instability in Drosophila melanogaster. Genetics, 232(3).

Sawyer JK, et al. (2020) Exploiting codon usage identifies intensity-specific modifiers of Ras/MAPK signaling in vivo. PLoS genetics, 16(12), e1009228.

Pischedda A, et al. (2020) The Loci of Behavioral Evolution: Evidence That Fas2 and tilB Underlie Differences in Pupation Site Choice Behavior between Drosophila melanogaster and D.??simulans. Molecular biology and evolution, 37(3), 864.

Li X, et al. (2020) The Mediator CDK8-Cyclin C complex modulates Dpp signaling in Drosophila by stimulating Mad-dependent transcription. PLoS genetics, 16(5), e1008832.