Resource Summary Report

Generated by NIF on Sep 7, 2026

w[1118]; sna[Sco]/CyO, P{ry[+t7.2]=en1}wg[en11]

RRID:BDSC_1672
Type: Organism

Proper Citation

RRID:BDSC_1672

Organism Information

URL: https://n2t.net/bdsc:1672

Proper Citation: RRID:BDSC_1672

Description: Drosophila melanogaster with name w[1118]; sna[Sco]/CyO, P{ry[+t7.2]=en1}wg[en11] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: David Bilder, Stanford University; Donor's Source: Norbert Perrimon, Harvard Medical School

Affected Gene: Ecol\lacZ, en, wg, sna, w

Genomic Alteration: Chromosome 1, Chromosome 2

Catalog Number: 1672

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_1672, BDSC:1672, BL1672

Organism Name: w[1118]; sna[Sco]/CyO, P{ry[+t7.2]=en1}wg[en11]

Record Creation Time: 20240911T222128+0000

Record Last Update: 20260905T135737+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; sna[Sco]/CyO, P{ry[+t7.2]=en1}wg[en11].

No alerts have been found for w[1118]; sna[Sco]/CyO, P{ry[+t7.2]=en1}wg[en11].

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 .

Agcaoili J, et al. (2024) Drosophila Robo3 guides longitudinal axons partially independently of its cytodomain. microPublication biology, 2024.

Rass M, et al. (2022) The Drosophila functional Smad suppressing element fuss, a homologue of the human Skor genes, retains pro-oncogenic properties of the Ski/Sno family. PloS one, 17(1), e0262360.

Korzelius J, et al. (2019) The WT1-like transcription factor Klumpfuss maintains lineage commitment of enterocyte progenitors in the Drosophila intestine. Nature communications, 10(1), 4123.

Xu K, et al. (2018) Temporospatial induction of homeodomain gene cut dictates natural lineage reprogramming. eLife, 7.