

Resource Summary Report

Generated by [ASWG](#) on Aug 7, 2026

Df(1)C128/FM6

RRID:BDSC_949

Type: Organism

Proper Citation

RRID:BDSC_949

Organism Information

URL: <https://n2t.net/bdsc:949>

Proper Citation: RRID:BDSC_949

Description: Drosophila melanogaster with name Df(1)C128/FM6 from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Caltech Stock Center

Affected Gene: asRNA:CR45535, asRNA:CR45620, bys, CG15330, CG15332, CG15333, CG1571, CG2233, CG2256, CG2258, Dhdds, dpr14, fs(1)h, Gclc, hdm, Ldsdh1, mahe, mys, orion, RpS14a, RpS14b, sn, snz, spidey, sws, Tbh, UbcE2H, Ubr3, Upf2, Ykt6

Genomic Alteration: Chromosome 1

Catalog Number: 949

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_949, BL949

Organism Name: Df(1)C128/FM6

Record Creation Time: 20240911T222125+0000

Record Last Update: 20260801T194536+0000

Ratings and Alerts

No rating or validation information has been found for Df(1)C128/FM6.

No alerts have been found for Df(1)C128/FM6.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

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

Messer CL, et al. (2025) A deficiency screen of the X chromosome for Rap1 GTPase dominant interacting genes in *Drosophila* border cell migration. *G3* (Bethesda, Md.), 15(5).

Boulanger A, et al. (2021) Axonal chemokine-like Orion induces astrocyte infiltration and engulfment during mushroom body neuronal remodeling. *Nature communications*, 12(1), 1849.

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.

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