

Resource Summary Report

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Df(1)BSC536, w[1118]/FM7h/Dp(2;Y)G, P{w[+mC]=hs-hid}Y

RRID:BDSC_25064

Type: Organism

Proper Citation

RRID:BDSC_25064

Organism Information

URL: <https://n2t.net/bdsc:25064>

Proper Citation: RRID:BDSC_25064

Description: Drosophila melanogaster with name Df(1)BSC536, w[1118]/FM7h/Dp(2;Y)G, P{w[+mC]=hs-hid}Y from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Kevin Cook, Bloomington Drosophila Stock Center

Affected Gene: Acat1, asRNA:CR43615, CARPA, CBP, CG10920, CG11368, CG11369, CG12155, CG12689, CG1409, CG15034, CG15035, CG15036, CG15478, CG1999, CG32718, CG32719, CG32720, CG32726, CG32727, CG34337, CG34338, CG42246, CG42704, CG42705, CG42781, CG43138, CG43198, CheA7a, CHES-1-like, ct, dec, Hira, Ir7a, Ir7b, Ir7c, Ir7d, Ir7e, Ir7f, Ir7g, lncRNA:CR43199, lncRNA:CR43278, lncRNA:CR43279, lncRNA:CR43287, lncRNA:CR43835, lncRNA:CR44357, lncRNA:CR44500, lncRNA:CR44501, lncRNA:CR44502, lncRNA:CR45533, lncRNA:CR45534, lncRNA:CR45665, lncRNA:CR45666, lncRNA:CR45667, Nek2, NELF-B, Pdp, Rab39, rumpel, Tom40, ZC3H18, hid, Hsp70 (generic), w

Genomic Alteration: Chromosome 1, Chromosome Y, Chromosome 2

Catalog Number: 25064

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_25064, BDSC:25064, BL25064

Organism Name: Df(1)BSC536, w[1118]/FM7h/Dp(2;Y)G, P{w[+mC]=hs-hid}Y

Record Creation Time: 20240911T222423+0000

Record Last Update: 20260905T140146+0000

Ratings and Alerts

No rating or validation information has been found for Df(1)BSC536, w[1118]/FM7h/Dp(2;Y)G, P{w[+mC]=hs-hid}Y.

No alerts have been found for Df(1)BSC536, w[1118]/FM7h/Dp(2;Y)G, P{w[+mC]=hs-hid}Y.

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 [dkNET](#) .

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).

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.

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