

Resource Summary Report

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Df(1)ED7217, w[1118]

P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7217/FM7h

RRID:BDSC_8952

Type: Organism

Proper Citation

RRID:BDSC_8952

Organism Information

URL: <https://n2t.net/bdsc:8952>

Proper Citation: RRID:BDSC_8952

Description: Drosophila melanogaster with name Df(1)ED7217, w[1118] P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7217/FM7h from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: amrt, asRNA:CR43833, CG10993, CG10996, CG11095, CG11134, CG11151, CG11158, CG11162, CG11164, CG12177, CG15760, CG32625, Clic, CtsB, DNAlig4, g, Grip91, inaE, lncRNA:CR42861, Mrgn1, mus101, Nadsyn, NFAT, Nna1, Pdcd4, rdgB, Rtc1, Yp3, w

Genomic Alteration: Chromosome 1

Catalog Number: 8952

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_8952, BL8952

Organism Name: Df(1)ED7217, w[1118] P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7217/FM7h

Record Creation Time: 20240911T222221+0000

Ratings and Alerts

No rating or validation information has been found for Df(1)ED7217, w[1118] P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7217/FM7h.

No alerts have been found for Df(1)ED7217, w[1118] P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED7217/FM7h.

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 [PRECISE-TBI](#) .

Rocha JJ, et al. (2023) Functional unknowns: Systematic screening of conserved genes of unknown function. PLoS biology, 21(8), e3002222.

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.

Hope KA, et al. (2019) A genome-wide enhancer/suppressor screen for Dube3a interacting genes in Drosophila melanogaster. Scientific reports, 9(1), 2382.