

Resource Summary Report

Generated by [SPARC SAWG](#) on Jul 29, 2026

[w\[1118\]; Df\(2L\)ED441,
P{w\[+mW.Scer\FRT.hs3\]=3'.RS5+3.3'}ED441/SM6a](#)

RRID:BDSC_24652

Type: Organism

Proper Citation

RRID:BDSC_24652

Organism Information

URL: <https://n2t.net/bdsc:24652>

Proper Citation: RRID:BDSC_24652

Description: Drosophila melanogaster with name w[1118]; Df(2L)ED441, P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED441/SM6a from BDSC.

Species: Drosophila melanogaster

Notes: Donor: DrosDel Project

Affected Gene: Aatf, AkhR, Atac1, Caper, CG11322, CG13775, CG17375, CG17376, CG17377, CG17378, CG18304, CG31908, CG31910, CG32829, CG3430, CG43232, CG43321, CG43322, CG43610, CG44271, Coprox, CR34663, CR43142, Daa02, Fgop2, Gart, Gas41, Heatr1, Hmgcl, homer, Hrb27C, ihog, Liprin-alpha, lncRNA:CR44081, lncRNA:CR44326, lncRNA:CR44611, lncRNA:CR44914, lncRNA:CR44915, LUBEL, meng, milt, mir-932, MME1, Mnn1, Mst27D, Nha1, Nlg2, nop5, nrv1, nrv2, Nuf2, Pcp, Polr1F, Pvf2, Rab30, Rat1, Rca1, SA1, Sem1, sens-2, sip2, snoRNA:CD67, snRNP-U1-70K, Sumo, tRNA:Arg-CCT-1-1, TTLL3A, TTLL3B, Ugt307A1, uif, Wee1, x16, w

Genomic Alteration: Chromosome 1, Chromosome 2

Catalog Number: 24652

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_24652, BL24652

Organism Name: w[1118]; Df(2L)ED441,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED441/SM6a

Record Creation Time: 20240911T222419+0000

Record Last Update: 20260725T195125+0000

Ratings and Alerts

No rating or validation information has been found for w[1118]; Df(2L)ED441,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED441/SM6a.

No alerts have been found for w[1118]; Df(2L)ED441,
P{w[+mW.Scer\FRT.hs3]=3'.RS5+3.3'}ED441/SM6a.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

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

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