

# Resource Summary Report

Generated by ASWG on Sep 20, 2026

## DGRP-399

RRID:BDSC\_25192

Type: Organism

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### Proper Citation

RRID:BDSC\_25192

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### Organism Information

**URL:** <https://n2t.net/bdsc:25192>

**Proper Citation:** RRID:BDSC\_25192

**Description:** Drosophila melanogaster with name DGRP-399 from BDSC.

**Species:** Drosophila melanogaster

**Notes:** Drosophila Genetic Reference Panel strain - sequenced isogenic reference wild type strain (aka 'Raleigh' or 'Mackay' line), core 40 set. Donor: Trudy Mackay, North Carolina State University

**Genomic Alteration:** Chromosome wt

**Catalog Number:** 25192

**Database:** Bloomington Drosophila Stock Center (BDSC)

**Database Abbreviation:** BDSC

**Availability:** available

**Alternate IDs:** BDSC\_25192, BDSC:25192, BL25192

**Organism Name:** DGRP-399

**Record Creation Time:** 20240911T222424+0000

**Record Last Update:** 20260919T202809+0000

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### Ratings and Alerts

No rating or validation information has been found for DGRP-399.

No alerts have been found for DGRP-399.

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## Data and Source Information

**Source:** [Integrated Animals](#)

**Source Database:** Bloomington Drosophila Stock Center (BDSC)

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## Usage and Citation Metrics

We found 17 mentions in open access literature.

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

Okuyama T, et al. (2026) Genetic heterogeneity alleviates behavioral trade-off between exploration and vigilance in *Drosophila*. *iScience*, 29(6), 116171.

Shiran MG, et al. (2026) Efficient RFLP-based Protocol for Routine Authentication of *Drosophila*. *microPublication biology*, 2026.

Koury S, et al. (2026) An extension of the Cx(Co) m model of crossover patterning to account for experimental mortality in *Drosophila melanogaster*. *bioRxiv : the preprint server for biology*.

Marconcini M, et al. (2026) Genome-wide association studies identify new candidate genes and tissues underlying resistance to a natural toxin in drosophilids. *G3 (Bethesda, Md.)*, 16(4).

Bak NK, et al. (2026) Host genetics and diet jointly shape the microbiome of *Drosophila melanogaster* but do not predict lifespan or age-related traits. *Biogerontology*, 27(3).

Backlund AE, et al. (2025) Mating status drives fitness trade-offs in exercised female *Drosophila*. *PloS one*, 20(10), e0327080.

Sato DX, et al. (2025) Multifaceted and extensive behavioral trajectories of genomically diverse *Drosophila* lines. *Scientific data*, 12(1), 400.

Yamamoto A, et al. (2024) The genetic basis of variation in *Drosophila melanogaster* mating behavior. *iScience*, 27(5), 109837.

Jansen G, et al. (2024) Tolerance thresholds underlie responses to DNA damage during germline development. *Genes & development*, 38(13-14), 631.

Tang C, et al. (2022) Genetic dissection of innate immune memory in *Drosophila melanogaster*. *Frontiers in immunology*, 13, 857707.

Spierer AN, et al. (2021) Natural variation in the regulation of neurodevelopmental genes modifies flight performance in *Drosophila*. *PLoS genetics*, 17(3), e1008887.

Chowdhury B, et al. (2021) The Divider Assay is a high-throughput pipeline for aggression analysis in *Drosophila*. *Communications biology*, 4(1), 85.

Patel SP, et al. (2021) Identification of genetic modifiers of lifespan on a high sugar diet in the *Drosophila* Genetic Reference Panel. *Heliyon*, 7(6), e07153.

Litovchenko M, et al. (2021) Extensive tissue-specific expression variation and novel regulators underlying circadian behavior. *Science advances*, 7(5).

Everett LJ, et al. (2020) Gene expression networks in the *Drosophila* Genetic Reference Panel. *Genome research*, 30(3), 485.

Okada H, et al. (2019) Sex-dependent and sex-independent regulatory systems of size variation in natural populations. *Molecular systems biology*, 15(11), e9012.

Langley CH, et al. (2012) Genomic variation in natural populations of *Drosophila melanogaster*. *Genetics*, 192(2), 533.