

Resource Summary Report

Generated by [NIF](#) on Aug 24, 2026

[FlyMine](#)

RRID:SCR_002694

Type: Tool

Proper Citation

FlyMine (RRID:SCR_002694)

Resource Information

URL: <http://www.flymine.org/>

Proper Citation: FlyMine (RRID:SCR_002694)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on January 14,2026. Integrated database of genomic, expression and protein data for Drosophila, Anopheles, C. elegans and other organisms. You can run flexible queries, export results and analyze lists of data. FlyMine presents data in categories, with each providing information on a particular type of data (for example Gene Expression or Protein Interactions). Template queries, as well as the QueryBuilder itself, allow you to perform searches that span data from more than one category. Advanced users can use a flexible query interface to construct their own data mining queries across the multiple integrated data sources, to modify existing template queries or to create your own template queries. Access our FlyMine data via our Application Programming Interface (API). We provide client libraries in the following languages: Perl, Python, Ruby and & Java API

Abbreviations: FlyMine

Resource Type: data or information resource, database

Defining Citation: [PMID:17615057](#)

Keywords: anopheles, genome, c. elegans, drosophila, gene, chromosomal location, genomics, proteomics, gene expression, interaction, homology, function, regulation, protein, phenotype, pathway, disease, publication, FASEB list

Funding: Wellcome Trust 067205;
NHGRI

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: FlyMine

Resource ID: SCR_002694

Alternate IDs: nif-0000-02845

Record Creation Time: 20220129T080214+0000

Record Last Update: 20260821T193650+0000

Ratings and Alerts

No rating or validation information has been found for FlyMine.

No alerts have been found for FlyMine.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 107 mentions in open access literature.

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

McLaughlin CN, et al. (2026) Endocytome profiling uncovers cell-surface protein dynamics underlying neuronal connectivity. *Neuron*, 114(11), 1935.

Li X, et al. (2025) Diacylglycerol metabolism drives host-pathogen responses during enteric infection in *Drosophila*. *bioRxiv : the preprint server for biology*.

Krabbenhoft SD, et al. (2025) Investigating oncoprotein-mediated chromatin dysregulation in *Drosophila melanogaster* uncovers novel modifiers of the developmental impact of H3 K27M and EZHIP. *bioRxiv : the preprint server for biology*.

Boyko AV, et al. (2025) Some molecular aspects of larval development in *Paralithodes camtschaticus*. *PloS one*, 20(4), e0322234.

Thompson A, et al. (2025) Quantifying Transcriptome Turnover on Phylogenies by Modeling Gene Expression as a Binary Trait. *Molecular biology and evolution*, 42(5).

Perlegos AE, et al. (2024) Cell type-specific regulation of m6 A modified RNAs in the aging *Drosophila* brain. *Aging cell*, 23(3), e14076.

Hrdina A, et al. (2024) The endosymbiont *Spiroplasma poulsonii* increases *Drosophila melanogaster* resistance to pathogens by enhancing iron sequestration and melanization. *mBio*, 15(8), e0093624.

Perlegos AE, et al. (2024) TDP-43 impairs sleep in *Drosophila* through Ataxin-2-dependent metabolic disturbance. *Science advances*, 10(2), eadj4457.

Leitão AB, et al. (2024) Recognition of nonself is necessary to activate *Drosophila*'s immune response against an insect parasite. *BMC biology*, 22(1), 89.

Samuels TJ, et al. (2024) Two distinct waves of transcriptome and translome changes drive *Drosophila* germline stem cell differentiation. *The EMBO journal*, 43(8), 1591.

Hikawa N, et al. (2024) Mating-induced increase of kynurenine in *Drosophila* ovary enhances starvation resistance of progeny. *The Journal of biological chemistry*, 300(3), 105663.

Lee JY, et al. (2024) Murine glial protrusion transcripts predict localized *Drosophila* glial mRNAs involved in plasticity. *The Journal of cell biology*, 223(10).

Talross GJS, et al. (2023) The rich non-coding RNA landscape of the *Drosophila* antenna. *Cell reports*, 42(5), 112482.

Mendaluk A, et al. (2023) A genome-wide RNAi screen for genes important for proliferation of cultured *Drosophila* cells at low temperature identifies the Ball/VRK protein kinase. *Chromosoma*, 132(1), 31.

Glaser-Schmitt A, et al. (2023) Dynamics and stage-specificity of between-population gene expression divergence in the *Drosophila melanogaster* larval fat body. *PLoS genetics*, 19(4), e1010730.

Krejčová G, et al. (2023) Macrophage-derived insulin antagonist ImpL2 induces lipoprotein mobilization upon bacterial infection. *The EMBO journal*, 42(23), e114086.

Burghardt E, et al. (2023) Transcriptome analysis reveals temporally regulated genetic networks during *Drosophila* border cell collective migration. *BMC genomics*, 24(1), 728.

Perlegos AE, et al. (2023) dTrmt10A impacts Hsp70 chaperone m6A levels and the stress response in the *Drosophila* brain. *Scientific reports*, 13(1), 22999.

Wippich F, et al. (2023) Nutritional stress-induced regulation of microtubule organization and mRNP transport by HDAC1 controlled α -tubulin acetylation. *Communications biology*, 6(1), 776.

Ismail JN, et al. (2023) Phenotypic and transcriptomic impact of expressing mammalian TET2 in the *Drosophila melanogaster* model. *Epigenetics*, 18(1), 2192375.