

Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

Dryad Digital Repository

RRID:SCR_005910

Type: Tool

Proper Citation

Dryad Digital Repository (RRID:SCR_005910)

Resource Information

URL: <https://datadryad.org>

Proper Citation: Dryad Digital Repository (RRID:SCR_005910)

Description: International, curated, digital repository that makes the data underlying scientific publications discoverable, freely reusable, and citable. Particularly data for which no specialized repository exists. Provides the infrastructure for, and promotes the re-use of, data underlying the scholarly literature. Governed by a nonprofit membership organization. Membership is open to any stakeholder organization, including but not limited to journals, scientific societies, publishers, research institutions, libraries, and funding organizations. Most data are associated with peer-reviewed articles, although data associated with non-peer reviewed publications from reputable academic sources, such as dissertations, are also accepted. Used to validate published findings, explore new analysis methodologies, repurpose data for research questions unanticipated by the original authors, and perform synthetic studies. UC system is member organization of Dryad general subject data repository.

Synonyms: , The Dryad Digital Repository, Dryad Digital Repository, Dryad

Resource Type: data or information resource, data repository, database, service resource, storage service resource

Keywords: international, digital, repository, curated, data, collection, scientific, medical, publication, dataset, FASEB list

Funding: European Commission ;
Institute for Museum and Library Services ;
JISC ;
NSF

Resource Name: Dryad Digital Repository

Resource ID: SCR_005910

Alternate IDs: DOI:10.25504/FAIRsharing.wkggtx, DOI:10.5061, r3d100000044, DOI:10.15146, DOI:10.17616/R34S33, nlx_149486

Alternate URLs: <https://doi.org/10.17616/R34S33>, <https://doi.org/10.5061/>, <https://doi.org/10.15146>, <https://dx.doi.org/10.5061/>, <https://dx.doi.org/10.15146>, <https://fairsharing.org/10.25504/FAIRsharing.wkggtx>, <https://api.datacite.org/doi?prefix=10.18736>, <https://api.datacite.org/doi?prefix=10.6076>, <https://doi.org/10.17616/R34S33>

Old URLs: <http://www.datadryad.org/>

License: Creative Commons Zero

License URLs: <http://datadryad.org/pages/policies>

Record Creation Time: 20220129T080233+0000

Record Last Update: 20260905T132543+0000

Ratings and Alerts

No rating or validation information has been found for Dryad Digital Repository.

No alerts have been found for Dryad Digital Repository.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2790 mentions in open access literature.

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

Ding C, et al. (2026) Glioblastoma-Secreted C1QL1 Orchestrates Tumor Microtube Expansion and Neural Synaptic Pruning to Drive Malignant Synapse Formation and Recurrence. Cancer discovery, 16(6), 1176.

Graham SM, et al. (2026) Acceptability, Feasibility, and Safety of the Shauriana Intervention for Young Kenyan Gay, Bisexual, and Other Men who Have Sex with Men, and Exploration of Potential Impact on Sexual Health Outcomes. AIDS and behavior, 30(5), 1267.

Wada-Katsumata A, et al. (2026) Bed bug preferences for host odor or aggregation odor are differentially modulated by physiological state in various odorscapes. Pest management science, 82(2), 1426.

Qin J, et al. (2026) Oocyte vitrification disrupts zygotic genome activation in embryos by impairing maternal spliceosome translation and Crxos splicing. PLoS genetics, 22(4),

e1012121.

Sharma S, et al. (2026) Worm orthologues of cytokinesis-associated proteins CIT and ASPM regulate neuronal microtubule dynamics and polarity in *C. elegans*. *PLoS genetics*, 22(4), e1012106.

A Batista R, et al. (2026) Dominance modifiers at the *Arabidopsis* self-incompatibility locus retain proto-miRNA features and act through non-canonical pathways. *PLoS genetics*, 22(4), e1012127.

Ramírez-Zavala B, et al. (2026) Inducible gene deletion reveals essentiality of protein kinases and a septation initiation network in *Candida albicans*. *PLoS genetics*, 22(4), e1012118.

Nußbaum P, et al. (2026) CsmR controls both, motility and cell shape, in *Haloferax volcanii*. *PLoS genetics*, 22(6), e1012198.

Domanico MC, et al. (2026) Naphthalene-DNA Adduct Formation in a Lung Airway Explant Model: The Role of Bioactivation and Naphthalene Metabolites. *Chemical research in toxicology*, 39(3), 339.

Yin G, et al. (2026) Nonlinear association of residual cholesterol to high-density lipoprotein cholesterol ratio with diabetes mellitus: a retrospective cohort study. *Scientific reports*, 16(1).

Dabak T, et al. (2026) The genetic basis of mimicry in the snowy bumble bee (*Bombus niveatus*) in Anatolia with insights from a color polymorphic gynandromorph. *PLoS genetics*, 22(3), e1012060.

Yue Y, et al. (2026) Genetic and functional characterization of AMH Signaling in Zebrafish - Evidence for Roles of *Amh-Bmpr2a-Bmpr1bb* Pathway in Controlling Gonadal Homeostasis. *PLoS genetics*, 22(3), e1011958.

Kafetzopoulos I, et al. (2026) DNMT1 loss leads to hypermethylation of a subset of late replicating domains by DNMT3A. *PLoS genetics*, 22(4), e1012098.

Schlomann BH, et al. (2026) Spatial microenvironments tune immune response dynamics in the *Drosophila* larval fat body. *PLoS genetics*, 22(2), e1012029.

Tian Y, et al. (2026) A dose-response relationship between low-density lipoprotein cholesterol levels within the normal range and the incidence of diabetes mellitus: a retrospective cohort study. *Frontiers in endocrinology*, 17, 1765884.

Castro DL, et al. (2026) The rewiring of a terminal selector regulatory cascade generates convergent neuronal laterality. *PLoS genetics*, 22(2), e1011782.

Voss AK, et al. (2026) KAT6A is essential for developmental control gene expression in neural stem and progenitor cells. *PLoS genetics*, 22(5), e1012133.

Xing Y, et al. (2026) Nonlinear associations between the cholesterol, high-density lipoprotein, and glucose index and type 2 diabetes risk in Chinese prediabetic adults: a cohort study. *Scientific reports*, 16(1).

Coleman DL, et al. (2026) G protein regulatory network shapes magnitude and kinetics of behavioral responses in an engineered opioid receptor model. *The Journal of biological chemistry*, 302(5), 111425.

Hénault M, et al. (2026) Towards a unified model of aneuploid karyotype dynamics. *PLoS genetics*, 22(6), e1012210.