Resource Summary Report

Generated by dkNET on Apr 18, 2025

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

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

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

Availability: Free, Freely available, Public, Acknowledgement requested

Resource Name: Dryad Digital Repository

Resource ID: SCR_005910

Alternate IDs: DOI:10.25504/FAIRsharing.wkggtx, DOI:10.5061, 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/dois?prefix=10.18736, https://api.datacite.org/dois?prefix=10.6076,

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: 20250418T055117+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 2278 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Carter-Cusack D, et al. (2025) Wild-type bone marrow cells repopulate tissue resident

macrophages and reverse the impacts of homozygous CSF1R mutation. PLoS genetics, 21(1), e1011525.

Curt JR, et al. (2025) Ambivalent partnership of the Drosophila posterior class Hox protein Abdominal-B with Extradenticle and Homothorax. PLoS genetics, 21(1), e1011355.

Tilahun Zewdu F, et al. (2025) Effectiveness of carbon dioxide cryotherapy for the treatment of cutaneous leishmaniasis: Systematic review and meta-analysis. PLoS neglected tropical diseases, 19(1), e0012741.

Hooper KM, et al. (2025) Short- and long-range roles of UNC-6/Netrin in dorsal-ventral axon guidance in vivo in Caenorhabditis elegans. PLoS genetics, 21(1), e1011526.

Gao Q, et al. (2025) Metabolically healthy overweight/obesity with no metabolic abnormalities and incident hyperglycaemia in Chinese adults: analysis of a retrospective cohort study. BMJ open, 15(1), e087307.

Nascimento Júnior JXD, et al. (2025) Dietary caloric input and tumor growth accelerate senescence and modulate liver and adipose tissue crosstalk. Communications biology, 8(1), 18.

Wang Z, et al. (2025) Sequence specificity of an essential nuclear localization sequence in Mcm3. PLoS genetics, 21(1), e1011499.

Zhou Q, et al. (2025) Association between the atherogenic index of plasma and long-term risk of type 2 diabetes: a 12-year cohort study based on the Japanese population. Cardiovascular diabetology, 24(1), 50.

De Walsche A, et al. (2025) metaGE: Investigating genotype x environment interactions through GWAS meta-analysis. PLoS genetics, 21(1), e1011553.

Baccas M, et al. (2025) SEM-2/SoxC regulates multiple aspects of C. elegans postembryonic mesoderm development. PLoS genetics, 21(1), e1011361.

Mosti F, et al. (2025) Multi-modal investigation reveals pathogenic features of diverse DDX3X missense mutations. PLoS genetics, 21(1), e1011555.

Hebert PDN, et al. (2025) Barcode 100K Specimens: In a Single Nanopore Run. Molecular ecology resources, 25(1), e14028.

Huang YZ, et al. (2025) Association between alcohol consumption and risk of type 2 diabetes in Japan: a population-base longitudinal cohort study. Scientific reports, 15(1), 630.

Ahn J, et al. (2025) Evolutionary lineage-specific genomic imprinting at the ZNF791 locus. PLoS genetics, 21(1), e1011532.

Pimentel MF, et al. (2025) Dual RNA-seq reveals transcriptome changes during Fusarium virguliforme-Trichoderma afroharzianum interactions. PloS one, 20(1), e0310850.

Jiang Y, et al. (2025) Atherogenic index of plasma and cardiovascular outcomes in female patients undergoing percutaneous coronary intervention: insights from a retrospective cohort study. Scientific reports, 15(1), 3112.

Gloriani M, et al. (2025) Sarcoglycans are enriched at the neuromuscular junction in a nerve-dependent manner. Cell death & disease, 16(1), 37.

Rice MM, et al. (2025) Terminology in ecology and evolutionary biology disproportionately harms marginalized groups. PLoS biology, 23(1), e3002933.

Song H, et al. (2025) Mean arterial pressure at the initiation of continuous renal replacement therapy as a prognostic indicator in patients with acute kidney injury. Renal failure, 47(1), 2448582.

Ricou A, et al. (2025) Identification of novel genes responsible for a pollen killer present in local natural populations of Arabidopsis thaliana. PLoS genetics, 21(1), e1011451.