Resource Summary Report

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Omics Discovery Index

RRID:SCR_010494
Type: Tool

Proper Citation

Omics Discovery Index (RRID:SCR_010494)

Resource Information

URL: http://www.omicsdi.org/

Proper Citation: Omics Discovery Index (RRID:SCR_010494)

Description: Portal for dataset discovery across a heterogeneous, distributed group of transcriptomics, genomics, proteomics and metabolomics data resources. These resources span eight repositories in three continents and six organisations, including both open and controlled access data resources.

Abbreviations: OmicsDI, DDI, DDICC

Synonyms: Omics Discovery Index (OmicsDI)

Resource Type: database, data or information resource, portal

Keywords: dataset search, knowledge framework, knowledge discovery

Funding: NIA 1U24AI117966-01;
NIGMS 1U54GM114833-01

Availability: Free, Public

Resource Name: Omics Discovery Index

Resource ID: SCR_010494

Alternate IDs: nlx_158507, SCR_014747

Record Creation Time: 20220129T080259+0000

Record Last Update: 20260808T190316+0000

Ratings and Alerts

No rating or validation information has been found for Omics Discovery Index.

No alerts have been found for Omics Discovery Index.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 33 mentions in open access literature.

Listed below are recent publications. The full list is available at nidm-terms .

Chianese U, et al. (2026) Glycolytic heterogeneity drives metabolic-targeted therapy in pancreatic ductal adenocarcinoma. Signal transduction and targeted therapy, 11(1), 25.

Deutsch EW, et al. (2026) The ProteomeXchange consortium in 2026: making proteomics data FAIR. Nucleic acids research, 54(D1), D459.

Cui R, et al. (2025) Prognostic and immunotherapeutic potential of disulfidptosis-associated signature in pancreatic cancer. Frontiers in immunology, 16, 1568976.

Iperi C, et al. (2025) BiomiX, a user-friendly bioinformatic tool for democratized analysis and integration of multiomics data. BMC bioinformatics, 26(1), 8.

Swaminathan G, et al. (2024) Exploiting open source omics data to advance pancreas research. Journal of pancreatology, 7(1), 21.

Deutsch EW, et al. (2023) The ProteomeXchange consortium at 10 years: 2023 update. Nucleic acids research, 51(D1), D1539.

Arenas GA, et al. (2022) Transcriptional and Epigenomic Markers of the Arterial-Venous and Micro/Macro-Vascular Endothelial Heterogeneity within the Umbilical-Placental Bed. International journal of molecular sciences, 23(19).

Boro? D, et al. (2022) Recent Multiomics Approaches in Endometrial Cancer. International journal of molecular sciences, 23(3).

Rincón-Riveros A, et al. (2022) Identification of Two Exosomal miRNAs in Circulating Blood of Cancer Patients by Using Integrative Transcriptome and Network Analysis. Non-coding RNA, 8(3).

Dato S, et al. (2021) Omics in a Digital World: The Role of Bioinformatics in Providing New Insights Into Human Aging. Frontiers in genetics, 12, 689824.

Kumar Das J, et al. (2021) Data science in unveiling COVID-19 pathogenesis and diagnosis: evolutionary origin to drug repurposing. Briefings in bioinformatics, 22(2), 855.

Rauner M, et al. (2021) Perspective of the GEMSTONE Consortium on Current and Future Approaches to Functional Validation for Skeletal Genetic Disease Using Cellular, Molecular and Animal-Modeling Techniques. Frontiers in endocrinology, 12, 731217.

Parolo S, et al. (2021) Reconstruction of the Cytokine Signaling in Lysosomal Storage Diseases by Literature Mining and Network Analysis. Frontiers in cell and developmental biology, 9, 703489.

Denoyelle L, et al. (2021) VarGoats project: a dataset of 1159 whole-genome sequences to dissect Capra hircus global diversity. Genetics, selection, evolution : GSE, 53(1), 86.

Fanidis D, et al. (2021) Fibromine is a multi-omics database and mining tool for target discovery in pulmonary fibrosis. Scientific reports, 11(1), 21712.

Subramanian I, et al. (2020) Multi-omics Data Integration, Interpretation, and Its Application. Bioinformatics and biology insights, 14, 1177932219899051.

Paananen J, et al. (2020) An omics perspective on drug target discovery platforms. Briefings in bioinformatics, 21(6), 1937.

Bono H, et al. (2020) All of gene expression (AOE): An integrated index for public gene expression databases. PloS one, 15(1), e0227076.

Palermo A, et al. (2020) Cloud-based archived metabolomics data: A resource for in-source fragmentation/annotation, meta-analysis and systems biology. Analytical science advances, 1(1), 70.

Krassowski M, et al. (2020) State of the Field in Multi-Omics Research: From Computational Needs to Data Mining and Sharing. Frontiers in genetics, 11, 610798.