

Resource Summary Report

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RAiD

RRID:SCR_023480

Type: Tool

Proper Citation

RAiD (RRID:SCR_023480)

Resource Information

URL: <https://www.raid.org.au/>

Proper Citation: RAiD (RRID:SCR_023480)

Description: Project that aims to provide persistent, unique and resolvable identifiers for research projects based on global handle system, funded by Australian Research Data Commons.

Synonyms: Research Activity Identifier

Resource Type: data or information resource, portal, community building portal, project portal

Keywords: PID, academic, persistent, unique, resolvable, identifiers, research projects, global handle system,

Funding: Australian Research Data Commons

Availability: Free

Resource Name: RAiD

Resource ID: SCR_023480

Alternate URLs: <https://app.demo.raid.org.au/raids>, <https://raid.org/10.82841>, <https://doi.org/10.82841>, <https://dx.doi.org/10.82841>

Record Creation Time: 20230422T050205+0000

Record Last Update: 20260808T190255+0000

Ratings and Alerts

No rating or validation information has been found for RAiD.

No alerts have been found for RAiD.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 22 mentions in open access literature.

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

Zhang M, et al. (2024) CBIL-VHPLI: a model for predicting viral-host protein-lncRNA interactions based on machine learning and transfer learning. Scientific reports, 14(1), 17549.

Sun L, et al. (2024) Differential gene expression and miRNA regulatory network in coronary slow flow. Scientific reports, 14(1), 8419.

Li G, et al. (2023) Exploring molecular markers and drug candidates for colorectal cancer through comprehensive bioinformatics analysis. Aging, 15(14), 7038.

Liu S, et al. (2022) Analysis of Multifactor-Driven Myopia Disease Modules to Guide Personalized Treatment and Drug Development. Computational and mathematical methods in medicine, 2022, 5262259.

Guo C, et al. (2021) Gene Expression Characteristics of Liver Tissue Reveal the Underlying Pathogenesis of Hepatocellular Carcinoma. BioMed research international, 2021, 9458328.

Yuan L, et al. (2021) A machine learning framework that integrates multi-omics data predicts cancer-related lncRNAs. BMC bioinformatics, 22(1), 332.

Zhang K, et al. (2021) Systematic analysis of molecular mechanisms of heart failure through the pathway and network-based approach. Life sciences, 265, 118830.

Lu C, et al. (2021) Breast cancer cell-derived extracellular vesicles transfer miR-182-5p and promote breast carcinogenesis via the CMTM7/EGFR/AKT axis. Molecular medicine (Cambridge, Mass.), 27(1), 78.

Sheng Z, et al. (2021) Identification of key non-coding RNAs and transcription factors regulators and their potential drugs for steroid-induced femoral head necrosis. Genomics, 113(2), 490.

Hu Y, et al. (2021) Bioinformatics analysis of genes related to iron death in diabetic nephropathy through network and pathway levels based approaches. PloS one, 16(11), e0259436.

Cao Y, et al. (2021) miR-590-3 and SP1 Promote Neuronal Apoptosis in Patients with Alzheimer's Disease via AMPK Signaling Pathway. *Contrast media & molecular imaging*, 2021, 6010362.

Yin X, et al. (2020) Identification of key modules and genes associated with breast cancer prognosis using WGCNA and ceRNA network analysis. *Aging*, 13(2), 2519.

Song J, et al. (2020) Identification of hydatidosis-related modules and key regulatory genes. *PeerJ*, 8, e9280.

Xu Z, et al. (2020) Development of Multiscale Transcriptional Regulatory Network in Esophageal Cancer Based on Integrated Analysis. *BioMed research international*, 2020, 5603958.

Fu X, et al. (2020) Construction of Glycometabolism- and Hormone-Related lncRNA-Mediated Feedforward Loop Networks Reveals Global Patterns of lncRNAs and Drug Repurposing in Gestational Diabetes. *Frontiers in endocrinology*, 11, 93.

Cui H, et al. (2020) Inferences of Individual Drug Response-Related Long Non-coding RNAs Based on Integrating Multi-omics Data in Breast Cancer. *Molecular therapy. Nucleic acids*, 20, 128.

Liang R, et al. (2020) Mechanism and Molecular Network of RBM8A-Mediated Regulation of Oxaliplatin Resistance in Hepatocellular Carcinoma. *Frontiers in oncology*, 10, 585452.

Li R, et al. (2020) CUL1-Mediated Organelle Fission Pathway Inhibits the Development of Chronic Obstructive Pulmonary Disease. *Computational and mathematical methods in medicine*, 2020, 5390107.

Shen SN, et al. (2020) Silencing lncRNAs PVT1 Upregulates miR-145 and Confers Inhibitory Effects on Viability, Invasion, and Migration in EC. *Molecular therapy. Nucleic acids*, 19, 668.

Wang F, et al. (2020) STAT3 speeds up progression of osteoarthritis through NF- κ B signaling pathway. *Experimental and therapeutic medicine*, 19(1), 722.