Resource Summary Report

Generated by NIF on Apr 29, 2025

MicroCosm Targets

RRID:SCR_010846

Type: Tool

Proper Citation

MicroCosm Targets (RRID:SCR_010846)

Resource Information

URL: http://www.ebi.ac.uk/enright-srv/microcosm/htdocs/targets/v5/

Proper Citation: MicroCosm Targets (RRID:SCR_010846)

Description: Database of computationally predicted targets for microRNAs across many

species.

Abbreviations: MicroCosm Targets

Synonyms: miRBase Targets

Resource Type: data or information resource, database

Funding:

Resource Name: MicroCosm Targets

Resource ID: SCR_010846

Alternate IDs: OMICS_00400

Alternate URLs: https://www.ebi.ac.uk/

Record Creation Time: 20220129T080301+0000

Record Last Update: 20250429T055446+0000

Ratings and Alerts

No rating or validation information has been found for MicroCosm Targets.

No alerts have been found for MicroCosm Targets.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 341 mentions in open access literature.

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

Wu W, et al. (2025) Identification of disulfidptosis-related subtypes in gastric cancer and GAMT is a key gene during disulfidptosis. Scientific reports, 15(1), 111.

Zhai Y, et al. (2025) Network pharmacology: a crucial approach in traditional Chinese medicine research. Chinese medicine, 20(1), 8.

Shan M, et al. (2025) Significance of KLK7 expression, polymorphisms, and function in sheep horn growth. BMC genomics, 26(1), 78.

Zhuo L, et al. (2025) MAPbrain: a multi-omics atlas of the primate brain. Nucleic acids research, 53(D1), D1055.

Cui J, et al. (2025) Telomere-to-telomere Phragmites australis reference genome assembly with a B chromosome provides insights into its evolution and polysaccharide biosynthesis. Communications biology, 8(1), 73.

Liu N, et al. (2025) RBM39 Enhances Cholangiocarcinoma Growth Through EZH2-mediated WNT7B/?-catenin Pathway. Cellular and molecular gastroenterology and hepatology, 19(1), 101404.

Jo H, et al. (2025) A fetal oncogene NUAK2 is an emerging therapeutic target in glioblastoma. bioRxiv: the preprint server for biology.

Chen Q, et al. (2025) Single-cell transcriptomics unveils multifaceted immune heterogeneity in early-onset versus late-onset cervical cancer. World journal of surgical oncology, 23(1), 12.

Wang H, et al. (2025) Identifying ADME-related gene signature for immune landscape and prognosis in KIRC by single-cell and spatial transcriptome analysis. Scientific reports, 15(1), 1294.

Shahbaz F, et al. (2024) Isolation and in vitro assessment of chicken gut microbes for probiotic potential. Frontiers in microbiology, 15, 1278439.

Cavanaugh NT, et al. (2024) Whole-genome sequencing of a Janthinobacterium sp. isolated from the Patagonian Desert. Microbiology resource announcements, 13(11), e0060024.

Chen S, et al. (2024) GEEES: inferring cell-specific gene-enhancer interactions from multi-modal single-cell data. Bioinformatics (Oxford, England), 40(11).

Hussaini SA, et al. (2024) Pharmacogenetics of Calcineurin inhibitors in kidney transplant recipients: the African gap. A narrative review. Pharmacogenomics, 25(7), 329.

Yuan G, et al. (2024) Genome-wide identification of Shaker K+ channel family in Nicotiana tabacum and functional analysis of NtSKOR1B in response to salt stress. Frontiers in plant science, 15, 1378738.

Ledesma-Bazan S, et al. (2024) Predicting prostate cancer progression with a Multi-IncRNA expression-based risk score and nomogram integrating ISUP grading. Non-coding RNA research, 9(2), 612.

Liu Y, et al. (2024) The causal relationship between human brain morphometry and knee osteoarthritis: a two-sample Mendelian randomization study. Frontiers in genetics, 15, 1420134.

Li J, et al. (2024) Cross-Cohort Gut Microbiome Signatures of Irritable Bowel Syndrome Presentation and Treatment. Advanced science (Weinheim, Baden-Wurttemberg, Germany), 11(41), e2308313.

Wang J, et al. (2024) Mendelian randomization reveals that abnormal lipid metabolism mediates the causal relationship between body mass index and keratoconus. Scientific reports, 14(1), 23698.

Guo X, et al. (2024) Validation of TYK2 and exploration of PRSS36 as drug targets for psoriasis using Mendelian randomization. Scientific reports, 14(1), 23902.

Liu B, et al. (2024) A genetic study to identify pathogenic mechanisms and drug targets for benign prostatic hyperplasia: a multi-omics Mendelian randomization study. Scientific reports, 14(1), 23120.