

Resource Summary Report

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[miRNAMap](#)

RRID:SCR_003156

Type: Tool

Proper Citation

miRNAMap (RRID:SCR_003156)

Resource Information

URL: <http://mirnamap.mbc.nctu.edu.tw>

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Description: A database of experimentally verified microRNAs and miRNA target genes in human, mouse, rat, and other metazoan genomes. In addition to known miRNA targets, three computational tools previously developed, such as miRanda, RNAhybrid and TargetScan, were applied for identifying miRNA targets in 3'-UTR of genes. In order to reduce the false positive prediction of miRNA targets, several criteria are supported for filtering the putative miRNA targets. Furthermore, miRNA expression profiles can provide valuable clues for investigating the properties of miRNAs, such tissue specificity and differential expression in cancer/normal cell. Therefore, we performed the Q-PCR experiments for monitoring the expression profiles of 224 human miRNAs in eighteen major normal tissues in human. The cross-reference between the miRNA expression profiles and the expression profiles of its target genes can provide effective viewpoint to understand the regulatory functions of the miRNA.

Abbreviations: miRNAMap

Resource Type: data or information resource, database

Defining Citation: [PMID:18029362](#), [PMID:16381831](#)

Keywords: microrna, genome, FASEB list

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: miRNAMap

Resource ID: SCR_003156

Alternate IDs: nif-0000-03138, OMICS_00408

Record Creation Time: 20220129T080217+0000

Ratings and Alerts

No rating or validation information has been found for miRNAMap.

No alerts have been found for miRNAMap.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 246 mentions in open access literature.

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

Zhu LL, et al. (2025) Plasma-Derived Small Extracellular Vesicles miR- 182 - 5p Is a Potential Biomarker for Diagnosing Major Depressive Disorder. Molecular neurobiology, 62(9), 11099.

Tout I, et al. (2025) The integrative genomic and functional immunological analyses of colorectal cancer initiating cells to modulate stemness properties and the susceptibility to immune responses. Journal of translational medicine, 23(1), 193.

Cheng J, et al. (2025) MiR- 146b-5p inhibits Candida albicans-induced inflammatory response through targeting HMGB1 in mouse primary peritoneal macrophages. Heliyon, 11(1), e41464.

Li T, et al. (2024) PMAIP1, a novel diagnostic and potential therapeutic biomarker in osteoporosis. Aging, 16(4), 3694.

Yan G, et al. (2024) Hsa_circ_0064636 regulates voltage dependent anion channel 1/ubiquitination factor E4A through miR?326/miR?503?5 in osteosarcoma. Oncology letters, 28(2), 374.

Rong X, et al. (2024) CircMEF2C(2, 3) modulates proliferation and adipogenesis of porcine intramuscular preadipocytes by miR-383/671-3p/MEF2C axis. iScience, 27(5), 109710.

Yan H, et al. (2024) Gastric cancer cell-derived exosomal miRNA-128-3p promotes angiogenesis by targeting SASH1. Frontiers in oncology, 14, 1440996.

Talluri B, et al. (2024) Adult zymosan re-exposure exacerbates the molecular alterations in the brainstem rostral ventromedial medulla of rats with early life zymosan-induced cystitis. Neurobiology of pain (Cambridge, Mass.), 16, 100160.

Wang X, et al. (2023) Integrated aqueous humor ceRNA and miRNA-TF-mRNA network analysis reveals potential molecular mechanisms governing primary open-angle glaucoma

pathogenesis. Indian journal of ophthalmology, 71(2), 553.

Ayaz H, et al. (2023) Mapping CircRNA-miRNA-mRNA regulatory axis identifies hsa_circ_0080942 and hsa_circ_0080135 as a potential theranostic agents for SARS-CoV-2 infection. PloS one, 18(4), e0283589.

Zhang X, et al. (2022) Identification of serum MiRNAs as candidate biomarkers for non-small cell lung cancer diagnosis. BMC pulmonary medicine, 22(1), 479.

El Bezawy R, et al. (2022) miR-550a-3p is a prognostic biomarker and exerts tumor-suppressive functions by targeting HSP90AA1 in diffuse malignant peritoneal mesothelioma. Cancer gene therapy, 29(10), 1394.

Tian D, et al. (2022) Tanshinone IIA protects against chronic obstructive pulmonary disease via exosome-shuttled miR-486-5p. International journal of molecular medicine, 50(1).

Wang L, et al. (2021) CircWAC induces chemotherapeutic resistance in triple-negative breast cancer by targeting miR-142, upregulating WWP1 and activating the PI3K/AKT pathway. Molecular cancer, 20(1), 43.

He Y, et al. (2021) Identification of a circRNA-miRNA-mRNA regulatory network for exploring novel therapeutic options for glioma. PeerJ, 9, e11894.

Kern F, et al. (2021) Validation of human microRNA target pathways enables evaluation of target prediction tools. Nucleic acids research, 49(1), 127.

Wu J, et al. (2021) Analysis of differential expression of long non-coding RNAs in exosomes derived from mature and immature dendritic cells. Molecular medicine reports, 23(2).

Hromadnikova I, et al. (2021) A History of Preterm Delivery Is Associated with Aberrant Postpartal MicroRNA Expression Profiles in Mothers with an Absence of Other Pregnancy-Related Complications. International journal of molecular sciences, 22(8).

Hromadnikova I, et al. (2021) Postnatal Expression Profile of MicroRNAs Associated with Cardiovascular Diseases in 3- to 11-Year-Old Preterm-Born Children. Biomedicines, 9(7).

Peraza-Vega RI, et al. (2021) miR-27b-3p a Negative Regulator of DSB-DNA Repair. Genes, 12(9).