Resource Summary Report

Generated by kravitz2 on Aug 1, 2026

miRTargetLink

RRID:SCR_026858
Type: Tool

Proper Citation

miRTargetLink (RRID:SCR_026858)

Resource Information

URL: https://ccb-compute.cs.uni-saarland.de/mirtargetlink2/

Proper Citation: miRTargetLink (RRID:SCR_026858)

Description: Web application provides users with visualization interface to explore and analyze interaction networks between miRNAs and target genes.

Synonyms: miRTargetLink 2.0

Resource Type: software resource, data access protocol, web service

Defining Citation: PMID:27089332, DOI:10.1093/nar/gkab297

Keywords: visualization interface, explore and analyze interaction networks, miRNAs and target genes, miRNA, interaction network, gene,

Funding: EU FP7 project BestAgeing ;
Saarland University.

Availability: Free, Freely available

Resource Name: miRTargetLink

Resource ID: SCR_026858

Record Creation Time: 20250430T055337+0000

Record Last Update: 20260801T042948+0000

Ratings and Alerts

No rating or validation information has been found for miRTargetLink.

No alerts have been found for miRTargetLink.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 10 mentions in open access literature.

Listed below are recent publications. The full list is available at kravitz2 .

Lobo J, et al. (2026) MicroRNA profiling of testicular Leydig cell tumors identifies a microRNA signature associated with malignancy and miR-196b-5p as a potentially useful biomarker. The Journal of pathology, 268(1), 54.

Kyurkchiyan S, et al. (2025) Co-expression of miRNA players in advanced laryngeal carcinoma - Insights into the roles of miR-93-5p, miR-145-5p, and miR-210-3p. Biomolecules & biomedicine, 25(5), 1052.

Giovannetti F, et al. (2025) Conjunctival MicroRNA Expression Signature in Primary Sjögren's Syndrome Dry Eye: A NanoString-Based Bioinformatic Analysis. Investigative ophthalmology & visual science, 66(4), 80.

Sanyal D, et al. (2025) Decoding molecular signatures and identifying therapeutic targets in triple-negative breast cancer subtypes using omics and network analysis. Scientific reports, 15(1), 44469.

Kim YJ, et al. (2025) Exosomal microRNA Panels for Detecting Early-Stage Non-Small Cell Lung Cancer. Diagnostics (Basel, Switzerland), 15(21).

Kargutkar N, et al. (2025) Decoding HbF reactivation by hydroxyurea in hemoglobinopathy patients through microRNA signatures. Annals of hematology, 104(4), 2159.

Sharma U, et al. (2025) Irradiation of muscle precursor cells impairs the proliferative and angiogenic functions of their extracellular vesicles. Scientific reports, 15(1), 30563.

Smith K, et al. (2024) Comprehensive analysis of differentially expressed miRNAs in hepatocellular carcinoma: Prognostic, predictive significance and pathway insights. PloS one, 19(4), e0296198.

Abdo SM, et al. (2023) Bio-diagnostic performances of microRNAs set related to DNA damage response pathway among hepatitis C virus-associated hepatocellular carcinoma patients. Journal, genetic engineering & biotechnology, 21(1), 85.

de Pedro MÁ, et al. (2023) Menstrual blood-derived stromal cells: insights into their secretome in acute hypoxia conditions. Molecular medicine (Cambridge, Mass.), 29(1), 48.