

# Resource Summary Report

Generated by [dkNET](#) on Sep 8, 2026

## TargetMiner

RRID:SCR\_010573

Type: Tool

### Proper Citation

TargetMiner (RRID:SCR\_010573)

### Resource Information

**URL:** [http://www.isical.ac.in/~bioinfo\\_miu/targetminer20.htm](http://www.isical.ac.in/~bioinfo_miu/targetminer20.htm)

**Proper Citation:** TargetMiner (RRID:SCR\_010573)

**Description:** A robust tool for microRNA target prediction with systematic identification of negative examples.

**Abbreviations:** TargetMiner

**Resource Type:** analysis service resource, data analysis service, production service resource, service resource, software resource

**Availability:** Free for academic use

**Resource Name:** TargetMiner

**Resource ID:** SCR\_010573

**Alternate IDs:** OMICS\_00419

**Record Creation Time:** 20220129T080259+0000

**Record Last Update:** 20260905T132643+0000

### Ratings and Alerts

No rating or validation information has been found for TargetMiner.

No alerts have been found for TargetMiner.

### Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 89 mentions in open access literature.

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

Chen Y, et al. (2026) Correlation between the expression levels of miR-199a-3p and FN1 in serum and aqueous humor and the severity of type 2 diabetic retinopathy in a Chinese population. *BMC ophthalmology*, 26(1).

Du Y, et al. (2025) Unraveling enhanced liver regeneration in ALPPS: Integrating multi-omics profiling and in vivo CRISPR in mouse models. *Hepatology communications*, 9(3).

Lin S, et al. (2025) Predicting microRNA target genes using pan-cancer correlation patterns. *BMC genomics*, 26(1), 77.

Liu C, et al. (2025) High Glucose-Induced Senescent Fibroblasts-Derived Exosomal miR-497 Inhibits Wound Healing by Regulating Endothelial Cellular Autophagy via ATG13. *Analytical cellular pathology (Amsterdam)*, 2025, 8890200.

Chen M, et al. (2024) miRSNP rs188493331: A key player in genetic control of microRNA-induced pathway activation in hypertrophic scars and keloids. *Skin research and technology : official journal of International Society for Bioengineering and the Skin (ISBS) [and] International Society for Digital Imaging of Skin (ISDIS) [and] International Society for Skin Imaging (ISSI)*, 30(5), e13686.

Nelson HM, et al. (2024) Transfer of miR-100 and miR-125b increases 3D growth and invasiveness in recipient cancer cells. *Extracellular vesicles and circulating nucleic acids*, 5(3), 397.

Xu J, et al. (2024) Correlation analysis of disulfidptosis-related gene signatures with clinical prognosis and immunotherapy response in sarcoma. *Scientific reports*, 14(1), 7158.

Piccinno E, et al. (2024) Identification of a Novel miR-195-5p/PNN Axis in Colorectal Cancer. *International journal of molecular sciences*, 25(11).

An Z, et al. (2024) Enhanced expression of miR-20a driven by nanog exacerbated the degradation of extracellular matrix in thoracic aortic dissection. *Non-coding RNA research*, 9(4), 1040.

Zheng S, et al. (2023) Extracellular shuttling miR-21 contributes to esophageal cancers and human umbilical vein endothelial cell communication in the tumor microenvironment and promotes tumor angiogenesis by targeting phosphatase and tensin homolog. *Thoracic cancer*, 14(31), 3119.

Li X, et al. (2023) Integrative pan-cancer analysis of cuproplasia-associated genes for the genomic and clinical characterization of 33 tumors. *Chinese medical journal*, 136(21), 2621.

Wang H, et al. (2023) A novel circRNA, hsa\_circ\_0069382, regulates gastric cancer progression. *Cancer cell international*, 23(1), 35.

Liu S, et al. (2023) Identification of hub genes associated with spermatogenesis by bioinformatics analysis. *Scientific reports*, 13(1), 18435.

Li Q, et al. (2023) Identification of macrophage-related genes in sepsis-induced ARDS using bioinformatics and machine learning. *Scientific reports*, 13(1), 9876.

Lin JJ, et al. (2023) Antitumor activity of miR-188-3p in gastric cancer is achieved by targeting CBL expression and inactivating the AKT/mTOR signaling. *World journal of gastrointestinal oncology*, 15(8), 1384.

Huang X, et al. (2023) Identification of basement membrane-related biomarkers associated with the diagnosis of osteoarthritis based on machine learning. *BMC medical genomics*, 16(1), 198.

Han X, et al. (2023) Machine learning developed a PI3K/Akt pathway-related signature for predicting prognosis and drug sensitivity in ovarian cancer. *Aging*, 15(20), 11162.

Ying TH, et al. (2022) Angelol-A exerts anti-metastatic and anti-angiogenic effects on human cervical carcinoma cells by modulating the phosphorylated-ERK/miR-29a-3p that targets the MMP2/VEGFA axis. *Life sciences*, 296, 120317.

Tu B, et al. (2022) Identification of a five-miRNA signature as a novel potential prognostic biomarker in patients with nasopharyngeal carcinoma. *Hereditas*, 159(1), 3.

Yang B, et al. (2022) CircRNA has\_circ\_0017109 promotes lung tumor progression via activation of Wnt/??-catenin signaling due to modulating miR-671-5p/FZD4 axis. *BMC pulmonary medicine*, 22(1), 443.