## **Resource Summary Report**

Generated by dkNET on Apr 25, 2025

# **TargetScore**

RRID:SCR\_012933

Type: Tool

## **Proper Citation**

TargetScore (RRID:SCR\_012933)

#### **Resource Information**

**URL:** http://www.bioconductor.org/packages/devel/bioc/html/TargetScore.html

**Proper Citation:** TargetScore (RRID:SCR\_012933)

**Description:** Software to infer the posterior distributions of microRNA targets by probabilistically modelling the likelihood microRNA-overexpression fold-changes and sequence-based scores.

Abbreviations: TargetScore

Synonyms: TargetScore: Infer microRNA targets using microRNA-overexpression data and

sequence information

**Resource Type:** software resource

**Defining Citation: PMID:24135265** 

**Funding:** 

**Availability:** Free

Resource Name: TargetScore

Resource ID: SCR\_012933

Alternate IDs: OMICS\_00421

**Record Creation Time:** 20220129T080313+0000

**Record Last Update:** 20250420T014625+0000

### **Ratings and Alerts**

No rating or validation information has been found for TargetScore.

No alerts have been found for TargetScore.

#### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 3 mentions in open access literature.

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

Yan G, et al. (2022) BET inhibition induces vulnerability to MCL1 targeting through upregulation of fatty acid synthesis pathway in breast cancer. Cell reports, 40(11), 111304.

Su H, et al. (2019) Identification of potential biomarkers of lung adenocarcinoma brain metastases via microarray analysis of cDNA expression profiles. Oncology letters, 17(2), 2228.

Wu J, et al. (2019) MicroRNA target gene prediction of ischemic stroke by using variational Bayesian inference for Gauss mixture model. Experimental and therapeutic medicine, 17(4), 2734.