Resource Summary Report

Generated by dkNET on May 19, 2025

RmiR

RRID:SCR_001069

Type: Tool

Proper Citation

RmiR (RRID:SCR_001069)

Resource Information

URL: http://www.bioconductor.org/packages/release/bioc/html/RmiR.html

Proper Citation: RmiR (RRID:SCR_001069)

Description: R package that contains functions to merge microRNA and respective targets

using different databases.

Abbreviations: RmiR

Resource Type: software application, data analysis software, data processing software,

sequence analysis software, software resource

Keywords: micro rna, mrna, function, target, r, sequence analysis software

Funding:

Availability: Open source, Available for installation

Resource Name: RmiR

Resource ID: SCR_001069

Alternate IDs: OMICS_00789

Alternate URLs: https://rdrr.io/bioc/RmiR/

Record Creation Time: 20220129T080205+0000

Record Last Update: 20250517T055454+0000

Ratings and Alerts

No rating or validation information has been found for RmiR.

No alerts have been found for RmiR.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.