

# Resource Summary Report

Generated by [dkNET](#) on Aug 3, 2026

## RmiR

RRID:SCR\_001069

Type: Tool

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### Proper Citation

RmiR (RRID:SCR\_001069)

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### 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 resource, data processing software, sequence analysis software, software application, data analysis software

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

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**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:** 20260803T040302+0000

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### Ratings and Alerts

No rating or validation information has been found for RmiR.

No alerts have been found for RmiR.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.