

# Resource Summary Report

Generated by [dkNET](#) on Aug 24, 2026

## [miRSeqNovel](#)

RRID:SCR\_013257

Type: Tool

### Proper Citation

miRSeqNovel (RRID:SCR\_013257)

### Resource Information

**URL:** <http://sourceforge.net/projects/mirseq/files/>

**Proper Citation:** miRSeqNovel (RRID:SCR\_013257)

**Description:** An R/Bioconductor based workflow for novel miRNA prediction from deep sequencing data.

**Abbreviations:** miRSeqNovel

**Resource Type:** software resource

**Availability:** Free, Public, Non-commercial

**Resource Name:** miRSeqNovel

**Resource ID:** SCR\_013257

**Alternate IDs:** OMICS\_00381

**Record Creation Time:** 20220129T080315+0000

**Record Last Update:** 20260815T182437+0000

### Ratings and Alerts

No rating or validation information has been found for miRSeqNovel.

No alerts have been found for miRSeqNovel.

### Data and Source Information

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

## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Wang Y, et al. (2024) HPV16-miRNAs exert oncogenic effects through enhancers in human cervical cancer. *Cancer cell international*, 24(1), 172.

Sablok G, et al. (2015) isomiRs: Increasing Evidences of isomiRs Complexity in Plant Stress Functional Biology. *Frontiers in plant science*, 6, 949.