# **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on Apr 15, 2025

# 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

Funding:

Availability: Free, Public, Non-commercial

Resource Name: miRSeqNovel

Resource ID: SCR\_013257

Alternate IDs: OMICS\_00381

Record Creation Time: 20220129T080315+0000

Record Last Update: 20250410T070348+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: <u>SciCrunch Registry</u>

## **Usage and Citation Metrics**

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

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.