

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 19, 2026

## [miRDeepFinder](#)

RRID:SCR\_004456

Type: Tool

### Proper Citation

miRDeepFinder (RRID:SCR\_004456)

### Resource Information

**URL:** <http://www.leonxie.com/DeepFinder.php>

**Proper Citation:** miRDeepFinder (RRID:SCR\_004456)

**Description:** Provides an comprehensive workflow of analyzing data from plant microRNA (miRNA) deep sequencing.

**Abbreviations:** miRDeepFinder

**Resource Type:** software resource

**Defining Citation:** [PMID:22290409](#)

**Resource Name:** miRDeepFinder

**Resource ID:** SCR\_004456

**Alternate IDs:** OMICS\_00362

**Record Creation Time:** 20220129T080224+0000

**Record Last Update:** 20260912T195613+0000

### Ratings and Alerts

No rating or validation information has been found for miRDeepFinder.

No alerts have been found for miRDeepFinder.

### 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 [PRECISE-TBI](#) .

Liu H, et al. (2019) Small RNA Sequencing Reveals Regulatory Roles of MicroRNAs in the Development of *Meloidogyne incognita*. International journal of molecular sciences, 20(21).

Xie F, et al. (2015) Deep sequencing reveals important roles of microRNAs in response to drought and salinity stress in cotton. Journal of experimental botany, 66(3), 789.