Resource Summary Report

Generated by dkNET on Sep 12, 2026

miRDeep-P

RRID:SCR_013026
Type: Tool

Proper Citation

miRDeep-P (RRID:SCR_013026)

Resource Information

URL: http://sourceforge.net/projects/mirdp/

Proper Citation: miRDeep-P (RRID:SCR_013026)

Description: A computational tool for analyzing the microRNA (miRNA) transcriptome in plants.

Abbreviations: miRDP

Resource Type: software resource

Defining Citation: PMID:21775303

Availability: GNU General Public License, v3

Resource Name: miRDeep-P

Resource ID: SCR_013026

Alternate IDs: OMICS_00375

Record Creation Time: 20220129T080313+0000

Record Last Update: 20260912T195757+0000

Ratings and Alerts

No rating or validation information has been found for miRDeep-P.

No alerts have been found for miRDeep-P.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET .

Joshi P, et al. (2025) Identification of miRNAs associated with Aspergillus flavus infection and their targets in groundnut (Arachis hypogaea L.). BMC plant biology, 25(1), 345.

Zhang Y, et al. (2020) Integrative expression network analysis of microRNA and gene isoforms in sacred lotus. BMC genomics, 21(1), 429.

Nadiya F, et al. (2017) Data on identification of conserved and novel miRNAs in Elettaria cardamomum. Data in brief, 14, 789.

Feng JL, et al. (2017) High throughput sequencing of small RNAs reveals dynamic microRNAs expression of lipid metabolism during Camellia oleifera and C. meiocarpa seed natural drying. BMC genomics, 18(1), 546.

Seifert F, et al. (2016) Analysis of wheat microspore embryogenesis induction by transcriptome and small RNA sequencing using the highly responsive cultivar "Svilena". BMC plant biology, 16, 97.

Bilichak A, et al. (2015) The elucidation of stress memory inheritance in Brassica rapa plants. Frontiers in plant science, 6, 5.