Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 13, 2025

MeRIP-PF

RRID:SCR_010904
Type: Tool

Proper Citation

MeRIP-PF (RRID:SCR_010904)

Resource Information

URL: http://software.big.ac.cn/MeRIP-PF.html

Proper Citation: MeRIP-PF (RRID:SCR_010904)

Description: A high-efficiency and easy-to-use analysis pipeline for MeRIP-Seq peak-finding at high resolution, which compares distributions of reads between immunoprecipitation sample and control sample.

Abbreviations: MeRIP-PF

Synonyms: MeRIP-Seq Peak-Finding Program

Resource Type: software resource

Keywords: bio.tools

Funding:

Resource Name: MeRIP-PF

Resource ID: SCR_010904

Alternate IDs: biotools:merip-pf, OMICS_00571

Alternate URLs: https://bio.tools/merip-pf

Record Creation Time: 20220129T080301+0000

Record Last Update: 20250410T070029+0000

Ratings and Alerts

No rating or validation information has been found for MeRIP-PF.

No alerts have been found for MeRIP-PF.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Yang Y, et al. (2023) Analysis approaches for the identification and prediction of N6-methyladenosine sites. Epigenetics, 18(1), 2158284.

Wang L, et al. (2022) m6A RNA methylation impairs gene expression variability and reproductive thermotolerance in Arabidopsis. Genome biology, 23(1), 244.

Liu L, et al. (2020) Bioinformatics approaches for deciphering the epitranscriptome: Recent progress and emerging topics. Computational and structural biotechnology journal, 18, 1587.