

# Resource Summary Report

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## Algorithms and Framework for Nonnegative Matrix Factorization

RRID:SCR\_023124

Type: Tool

### Proper Citation

Algorithms and Framework for Nonnegative Matrix Factorization (RRID:SCR\_023124)

### Resource Information

**URL:** <https://cran.r-project.org/package=NMF>

**Proper Citation:** Algorithms and Framework for Nonnegative Matrix Factorization (RRID:SCR\_023124)

**Description:** Software R package for nonnegative matrix factorization. Implements set of already published algorithms and seeding methods, and provides framework to test, develop and plug new/custom algorithms.

**Abbreviations:** NMF

**Synonyms:** Non-negative Matrix Factorization

**Resource Type:** software resource, software toolkit

**Defining Citation:** [DOI:10.1186/1471-2105-11-367](https://doi.org/10.1186/1471-2105-11-367)

**Keywords:** Non-negative Matrix Factorization, nonnegative matrix factorization,

**Funding:** South-African National Bioinformatics Network ;  
Science Foundation Ireland

**Availability:** Free, Available for download, Freely available

**Resource Name:** Algorithms and Framework for Nonnegative Matrix Factorization

**Resource ID:** SCR\_023124

**Record Creation Time:** 20230116T062750+0000

**Record Last Update:** 20250422T060318+0000

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## Ratings and Alerts

No rating or validation information has been found for Algorithms and Framework for Nonnegative Matrix Factorization.

No alerts have been found for Algorithms and Framework for Nonnegative Matrix Factorization.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Zhang J, et al. (2024) Protocol to infer and analyze miRNA sponge modules in heterogeneous data using miRSM 2.0. STAR protocols, 5(4), 103317.

Ambeskovic A, et al. (2024) Exon-Skipping-Based Subtyping of Colorectal Cancers. Gastroenterology.

Berard AR, et al. (2023) Vaginal epithelial dysfunction is mediated by the microbiome, metabolome, and mTOR signaling. Cell reports, 42(5), 112474.

Huang C, et al. (2023) Identification of S1PR4 as an immune modulator for favorable prognosis in HNSCC through machine learning. iScience, 26(9), 107693.