

Resource Summary Report

Generated by [NIF](#) on Aug 9, 2026

EMImR

RRID:SCR_027327

Type: Tool

Proper Citation

EMImR (RRID:SCR_027327)

Resource Information

URL: <https://github.com/omicscodeathon/emimr>

Proper Citation: EMImR (RRID:SCR_027327)

Description: Software application for identifying transcriptomic and epigenomic changes.

Resource Type: software resource, software application, source code, data analysis software, time-series analysis software, 1d time-series analysis software, data processing software

Keywords: identifying transcriptomic and epigenomic changes, transcriptomic changes, epigenomic changes,

Availability: Free, Available for download, Freely available

Resource Name: EMImR

Resource ID: SCR_027327

License: Artistic-2.0 license

Record Creation Time: 20250818T062958+0000

Record Last Update: 20260808T190806+0000

Ratings and Alerts

No rating or validation information has been found for EMImR.

No alerts have been found for EMImR.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Ben Aribi H, et al. (2025) EMIImR: a Shiny application for identifying transcriptomic and epigenomic changes. GigaByte (Hong Kong, China), 2025, gigabyte168.