

# Resource Summary Report

Generated by [dkNET](#) on Sep 10, 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:** 1d time-series analysis software, data analysis software, data processing software, software application, software resource, source code, time-series analysis 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:** 20260905T133545+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](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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