

Resource Summary Report

Generated by [NIF](#) on Aug 27, 2026

[eisa](#)

RRID:SCR_012883

Type: Tool

Proper Citation

eisa (RRID:SCR_012883)

Resource Information

URL: <http://www.bioconductor.org/packages/release/bioc/html/eisa.html>

Proper Citation: eisa (RRID:SCR_012883)

Description: A biclustering method; it finds correlated blocks (transcription modules) in gene expression (or other tabular) data.

Abbreviations: eisa

Synonyms: eisa - Expression data analysis via the Iterative Signature Algorithm

Resource Type: software resource

Keywords: bio.tools

Availability: Free

Resource Name: eisa

Resource ID: SCR_012883

Alternate IDs: OMICS_01801, biotools:eisa

Alternate URLs: <https://bio.tools/eisa>

Record Creation Time: 20220129T080313+0000

Record Last Update: 20260815T182450+0000

Ratings and Alerts

No rating or validation information has been found for eisa.

No alerts have been found for eisa.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Pitoy A, et al. (2024) Isatuximab-dexamethasone-pomalidomide combination effects on serum M protein and PFS in myeloma: Development of a joint model using phase I/II data. CPT: pharmacometrics & systems pharmacology, 13(12), 2087.

Munz N, et al. (2021) Exon-Intron Differential Analysis Reveals the Role of Competing Endogenous RNAs in Post-Transcriptional Regulation of Translation. Non-coding RNA, 7(2).