Resource Summary Report

Generated by RRID on Apr 9, 2025

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

Funding:

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: 20250214T183219+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 RRID.

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).