

# Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

## iASeq

RRID:SCR\_000420

Type: Tool

### Proper Citation

iASeq (RRID:SCR\_000420)

### Resource Information

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

**Proper Citation:** iASeq (RRID:SCR\_000420)

**Description:** Software that uses a Bayesian hierarchical mixture model to learn correlation patterns of allele-specificity among multiple proteins.

**Synonyms:** iASeq: integrating multiple sequencing datasets for detecting allele-specific events

**Resource Type:** software resource

**Defining Citation:** [PMID:23194258](#)

**Keywords:** software package, unix/linux, mac os x, windows, r, chip-seq, rna-seq, snp

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

**Resource Name:** iASeq

**Resource ID:** SCR\_000420

**Alternate IDs:** OMICS\_05505

**Record Creation Time:** 20220129T080201+0000

**Record Last Update:** 20260725T190444+0000

### Ratings and Alerts

No rating or validation information has been found for iASeq.

No alerts have been found for iASeq.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.