

Resource Summary Report

Generated by T1D on Aug 9, 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](https://pubmed.ncbi.nlm.nih.gov/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: 20260808T185722+0000

Ratings and Alerts

No rating or validation information has been found for iASeq.

No alerts have been found for iASeq.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.