### **Resource Summary Report**

Generated by dkNET on Apr 25, 2025

# **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

**Funding:** 

Availability: GNU General Public License, v2

Resource Name: iASeq

Resource ID: SCR\_000420

Alternate IDs: OMICS\_05505

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

Record Last Update: 20250420T013946+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.