### **Resource Summary Report**

Generated by dkNET on Apr 25, 2025

## **iBMQ**

RRID:SCR\_000481

Type: Tool

### **Proper Citation**

iBMQ (RRID:SCR\_000481)

#### **Resource Information**

URL: http://www.bioconductor.org/packages/release/bioc/html/iBMQ.html

Proper Citation: iBMQ (RRID:SCR\_000481)

Description: Software for integrated Bayesian Modeling of eQTL data. It implements a joint

hierarchical Bayesian model where all genes and SNPs are modeled concurrently.

Synonyms: iBMQ - integrated Bayesian Modeling of eQTL data

**Resource Type:** software resource

**Defining Citation: PMID:23958729** 

**Keywords:** standalone software, mac os x, unix/linux, windows, r, gene expression,

microarray, preprocessing, snp

**Funding:** 

Availability: Artistic License, v2

Resource Name: iBMQ

Resource ID: SCR\_000481

Alternate IDs: OMICS\_04601

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

Record Last Update: 20250420T013950+0000

### Ratings and Alerts

No rating or validation information has been found for iBMQ.

No alerts have been found for iBMQ.

### Data and Source Information

Source: SciCrunch Registry

# Usage and Citation Metrics

We have not found any literature mentions for this resource.