

# Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

## 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

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

**Resource Name:** iBMQ

**Resource ID:** SCR\_000481

**Alternate IDs:** OMICS\_04601

**License:** Artistic License, v2

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

**Record Last Update:** 20260912T195510+0000

### Ratings and Alerts

No rating or validation information has been found for iBMQ.

No alerts have been found for iBMQ.

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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.