

# Resource Summary Report

Generated by [dkNET](#) on Sep 5, 2026

## R/QTLBIM

RRID:SCR\_009375

Type: Tool

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### Proper Citation

R/QTLBIM (RRID:SCR\_009375)

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### Resource Information

**URL:** <http://pages.stat.wisc.edu/~yandell/qtl/software/qtlbim/>

**Proper Citation:** R/QTLBIM (RRID:SCR\_009375)

**Description:** Software library for QTL Bayesian Interval Mapping that provides a Bayesian model selection approach to map multiple interacting QTL. It works on experimentally inbred lines and performs a genome-wide search to locate multiple potential QTL. The package can handle continuous, binary and ordinal traits. (entry from Genetic Analysis Software)

**Resource Type:** software application, software library, software resource, software toolkit

**Keywords:** gene, genetic, genomic, r, bio.tools

**Resource Name:** R/QTLBIM

**Resource ID:** SCR\_009375

**Alternate IDs:** nlx\_154597, biotools:qtlbim

**Alternate URLs:** <http://www.ssg.uab.edu/qtlbim/index.jsp>, <https://cran.r-project.org/src/contrib/Archive/qtlbim/>, <https://bio.tools/qtlbim>

**Old URLs:** <http://www.qtlbim.org/>

**Record Creation Time:** 20220129T080252+0000

**Record Last Update:** 20260903T235008+0000

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### Ratings and Alerts

No rating or validation information has been found for R/QTLBIM.

No alerts have been found for R/QTLBIM.

## Data and Source Information

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

## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [dkNET](#) .

Chung W, et al. (2014) Mixed-effects models for GAW18 longitudinal blood pressure data. BMC proceedings, 8(Suppl 1), S87.

Salom?? PA, et al. (2011) Genetic architecture of flowering-time variation in Arabidopsis thaliana. Genetics, 188(2), 421.