

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

R/QTLBIM

RRID:SCR_009375

Type: Tool

Proper Citation

R/QTLBIM (RRID:SCR_009375)

Resource Information

URL: <http://pages.stat.wisc.edu/~yandell/ctl/software/qlbim/>

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:qlbim

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

Old URLs: <http://www.qlbim.org/>

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260905T132632+0000

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 [Kravitz](#) .

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.