

Resource Summary Report

Generated by [kravitz2](#) on Sep 4, 2026

qtlreaper

RRID:SCR_024199

Type: Tool

Proper Citation

qtlreaper (RRID:SCR_024199)

Resource Information

URL: <https://qtlreaper.sourceforge.net/>

Proper Citation: qtlreaper (RRID:SCR_024199)

Description: Software, written in C and compiled as Python module, for rapidly scanning microarray expression data for Quantitative Trait Loci. Searches for association between each expression trait and all genotypes and evaluates that association by permutation test. Performs bootstrap resampling to estimate confidence region for location of putative QTL.

Resource Type: data analysis software, data processing software, software application, software resource

Keywords: rapidly scanning microarray expression data for QTLs, association between each expression trait and all genotypes, estimate confidence region for location of putative QTL, Quantitative Trait Loci,

Availability: Free, Available for download, Freely available,

Resource Name: qtlreaper

Resource ID: SCR_024199

Old URLs: <https://sources.debian.org/src/qtlreaper/>

Record Creation Time: 20230824T050212+0000

Record Last Update: 20260903T235933+0000

Ratings and Alerts

No rating or validation information has been found for qtlreaper.

No alerts have been found for qtlreaper.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.