

# Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

## MULTIMAPPER/OUTBRED

RRID:SCR\_009306

Type: Tool

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### Proper Citation

MULTIMAPPER/OUTBRED (RRID:SCR\_009306)

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### Resource Information

**URL:** <http://www.rni.helsinki.fi/~mjs/>

**Proper Citation:** MULTIMAPPER/OUTBRED (RRID:SCR\_009306)

**Description:** Bayesian QTL mapping software for analysing backcross and F2 data from designed crossing experiments of outbred lines (entry from Genetic Analysis Software)

**Abbreviations:** MULTIMAPPER/OUTBRED

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

**Defining Citation:** [PMID:10101181](#)

**Keywords:** gene, genetic, genomic, c

**Resource Name:** MULTIMAPPER/OUTBRED

**Resource ID:** SCR\_009306

**Alternate IDs:** nlx\_154496

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

**Record Last Update:** 20260727T040458+0000

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### Ratings and Alerts

No rating or validation information has been found for MULTIMAPPER/OUTBRED.

No alerts have been found for MULTIMAPPER/OUTBRED.

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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.