

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

MULTIMAPPER/OUTBRED

RRID:SCR_009306

Type: Tool

Proper Citation

MULTIMAPPER/OUTBRED (RRID:SCR_009306)

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: 20260905T133249+0000

Ratings and Alerts

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

No alerts have been found for MULTIMAPPER/OUTBRED.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.