

Resource Summary Report

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

PSEUDOMARKER.M

RRID:SCR_009346

Type: Tool

Proper Citation

PSEUDOMARKER.M (RRID:SCR_009346)

Resource Information

URL: <http://www.jax.org/staff/churchill/labsite/software/pseudomarker/index.html>

Proper Citation: PSEUDOMARKER.M (RRID:SCR_009346)

Description: A set of programs written in MATLAB for the analysis of QTL data from inbred line crosses. (entry from Genetic Analysis Software)

Abbreviations: PSEUDOMARKER.M

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, matlab

Resource Name: PSEUDOMARKER.M

Resource ID: SCR_009346

Alternate IDs: nlx_154558

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260905T133250+0000

Ratings and Alerts

No rating or validation information has been found for PSEUDOMARKER.M.

No alerts have been found for PSEUDOMARKER.M.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Johnson KR, et al. (2006) Strain background effects and genetic modifiers of hearing in mice. Brain research, 1091(1), 79.