

Resource Summary Report

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

[massiR](#)

RRID:SCR_001157

Type: Tool

Proper Citation

massiR (RRID:SCR_001157)

Resource Information

URL: <http://bioconductor.org/packages/devel/bioc/html/massiR.html>

Proper Citation: massiR (RRID:SCR_001157)

Description: Software that predicts the sex of samples in gene expression microarray datasets.

Synonyms: massiR: MicroArray Sample Sex Identifier, MicroArray Sample Sex Identifier

Resource Type: software resource

Defining Citation: [PMID:24659105](#)

Keywords: standalone software, mac os x, unix/linux, windows, r, classification, clustering, gene expression, microarray, quality control, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: massiR

Resource ID: SCR_001157

Alternate IDs: biotools:massir, OMICS_03638

Alternate URLs: <https://bio.tools/massir>

Record Creation Time: 20220129T080205+0000

Record Last Update: 20260905T132430+0000

Ratings and Alerts

No rating or validation information has been found for massiR.

No alerts have been found for massiR.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.