

# Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

## iterativeBMAurv

RRID:SCR\_001254

Type: Tool

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### Proper Citation

iterativeBMAurv (RRID:SCR\_001254)

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### Resource Information

**URL:** <http://www.bioconductor.org/packages/release/bioc/html/iterativeBMAurv.html>

**Proper Citation:** iterativeBMAurv (RRID:SCR\_001254)

**Description:** Software package providing a variable selection method for applying survival analysis to microarray data.

**Abbreviations:** iterativeBMAurv

**Synonyms:** The Iterative Bayesian Model Averaging (BMA) Algorithm For Survival Analysis, iterativeBMAurv - The Iterative Bayesian Model Averaging (BMA) Algorithm For Survival Analysis

**Resource Type:** software resource

**Defining Citation:** [PMID:19245714](#)

**Keywords:** microarray

**Availability:** GNU General Public License, v2 or newer

**Resource Name:** iterativeBMAurv

**Resource ID:** SCR\_001254

**Alternate IDs:** OMICS\_02086

**Record Creation Time:** 20220129T080206+0000

**Record Last Update:** 20260808T185727+0000

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### Ratings and Alerts

No rating or validation information has been found for iterativeBMAsurv.

No alerts have been found for iterativeBMAsurv.

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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.