

Resource Summary Report

Generated by [RRID](#) on Aug 4, 2026

iterativeBMAurv

RRID:SCR_001254

Type: Tool

Proper Citation

iterativeBMAurv (RRID:SCR_001254)

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: 20260801T190126+0000

Ratings and Alerts

No rating or validation information has been found for iterativeBMAsurv.

No alerts have been found for iterativeBMAsurv.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.