## **Resource Summary Report**

Generated by <u>dkNET</u> on Apr 24, 2025

# **iterativeBMAsurv**

RRID:SCR\_001254 Type: Tool

**Proper Citation** 

iterativeBMAsurv (RRID:SCR\_001254)

#### **Resource Information**

URL: http://www.bioconductor.org/packages/release/bioc/html/iterativeBMAsurv.html

Proper Citation: iterativeBMAsurv (RRID:SCR\_001254)

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

Abbreviations: iterativeBMAsurv

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

Resource Type: software resource

Defining Citation: PMID:19245714

Keywords: microarray

**Funding:** 

Availability: GNU General Public License, v2 or newer

Resource Name: iterativeBMAsurv

Resource ID: SCR\_001254

Alternate IDs: OMICS\_02086

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