# **Resource Summary Report**

Generated by dkNET on May 18, 2025

# **BAC**

RRID:SCR\_001067

Type: Tool

## **Proper Citation**

BAC (RRID:SCR\_001067)

#### **Resource Information**

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

Proper Citation: BAC (RRID:SCR\_001067)

**Description:** R software package that uses a Bayesian hierarchical model to detect enriched

regions from ChIP-chip experiments.

**Abbreviations: BAC** 

**Resource Type:** software application, data analysis software, data processing software,

sequence analysis software, software resource

**Keywords:** software, bayesian, chip, hierarchy, sequence analysis software

**Funding:** 

Availability: Open source, Available for installation

Resource Name: BAC

Resource ID: SCR\_001067

Alternate IDs: OMICS\_00801

**Record Creation Time:** 20220129T080205+0000

**Record Last Update**: 20250517T055454+0000

## **Ratings and Alerts**

No rating or validation information has been found for BAC.

No alerts have been found for BAC.

## **Data and Source Information**

Source: SciCrunch Registry

# **Usage and Citation Metrics**

We have not found any literature mentions for this resource.