

Resource Summary Report

Generated by RRID on Aug 5, 2026

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: data processing software, data analysis software, software resource, sequence analysis software, software application

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

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: BAC

Resource ID: SCR_001067

Alternate IDs: OMICS_00801

Record Creation Time: 20220129T080205+0000

Record Last Update: 20260805T044017+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.