

# Resource Summary Report

Generated by [NIF](#) on Aug 31, 2026

## BAC

RRID:SCR\_001067

Type: Tool

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### Proper Citation

BAC (RRID:SCR\_001067)

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

**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:** 20260829T182035+0000

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### Ratings and Alerts

No rating or validation information has been found for BAC.

No alerts have been found for BAC.

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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.