

Resource Summary Report

Generated by SPARC SAWG on Sep 15, 2026

iBBiG

RRID:SCR_012882

Type: Tool

Proper Citation

iBBiG (RRID:SCR_012882)

Resource Information

URL: <http://www.bioconductor.org/packages/release/bioc/html/iBBiG.html>

Proper Citation: iBBiG (RRID:SCR_012882)

Description: A bi-clustering algorithm which is optimizes for binary data analysis.

Abbreviations: iBBiG

Synonyms: Iterative Binary Biclustering of Genesets

Resource Type: software resource

Defining Citation: [PMID:22789589](#)

Availability: Free

Resource Name: iBBiG

Resource ID: SCR_012882

Alternate IDs: OMICS_01802

Record Creation Time: 20220129T080313+0000

Record Last Update: 20260912T195754+0000

Ratings and Alerts

No rating or validation information has been found for iBBiG.

No alerts have been found for iBBiG.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

Li H, et al. (2020) TERT mutations correlate with higher TMB value and unique tumor microenvironment and may be a potential biomarker for anti-CTLA4 treatment. Cancer medicine, 9(19), 7151.

Thorsson V, et al. (2018) The Immune Landscape of Cancer. Immunity, 48(4), 812.

Uzun A, et al. (2016) Targeted Sequencing and Meta-Analysis of Preterm Birth. PloS one, 11(5), e0155021.