

# Resource Summary Report

Generated by [ASWG](#) on Aug 25, 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:** 20260815T182451+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](#)

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

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.