## **Resource Summary Report**

Generated by NIF on May 25, 2025

# ggsignif

RRID:SCR\_023047

Type: Tool

## **Proper Citation**

ggsignif (RRID:SCR\_023047)

#### **Resource Information**

URL: https://CRAN.R-project.org/package=ggsignif

**Proper Citation:** ggsignif (RRID:SCR\_023047)

**Description:** Software package to indicate if two groups are significantly different. Used to add significance brackets to ggplots.

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

**Keywords:** indicate if two groups are significantly different, add significance brackets

**Funding:** 

Availability: Free, Available for download, Freely available

Resource Name: ggsignif

Resource ID: SCR\_023047

Alternate URLs: https://github.com/const-ae/ggsignif

License: GPL v3

**Record Creation Time:** 20221215T050202+0000

**Record Last Update:** 20250525T031935+0000

## **Ratings and Alerts**

No rating or validation information has been found for ggsignif.

No alerts have been found for ggsignif.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 9 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.

Olweny G, et al. (2025) Protocol for identifying Mycobacterium tuberculosis infection status through airway microbiome profiling. STAR protocols, 6(1), 103574.

Selma-Royo M, et al. (2024) Birthmode and environment-dependent microbiota transmission dynamics are complemented by breastfeeding during the first year. Cell host & microbe, 32(6), 996.

Yang B, et al. (2024) Identification of ferroptosis-related gene signature for tuberculosis diagnosis and therapy efficacy. iScience, 27(7), 110182.

Giovannetti M, et al. (2024) SIN-3 transcriptional coregulator maintains mitochondrial homeostasis and polyamine flux. iScience, 27(5), 109789.

Zhao J, et al. (2023) Changes in m6A RNA methylation are associated with male sterility in wolfberry. BMC plant biology, 23(1), 456.

Ares-Arroyo M, et al. (2023) Origins of transfer establish networks of functional dependencies for plasmid transfer by conjugation. Nucleic acids research, 51(7), 3001.

Elizabeth Deeter M, et al. (2023) Accelerated abdominal lipid depletion from pesticide treatment alters honey bee pollen foraging strategy, but not onset, in worker honey bees. The Journal of experimental biology, 226(7).

Midha AD, et al. (2023) Organ-specific fuel rewiring in acute and chronic hypoxia redistributes glucose and fatty acid metabolism. Cell metabolism, 35(3), 504.

Salimi A, et al. (2023) InterOpt: Improved gene expression quantification in qPCR experiments using weighted aggregation of reference genes. iScience, 26(10), 107945.