

Resource Summary Report

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ggbiplot

RRID:SCR_025581

Type: Tool

Proper Citation

ggbiplot (RRID:SCR_025581)

Resource Information

URL: <https://cran.r-project.org/web/packages/ggbiplot/readme/README.html>

Proper Citation: ggbiplot (RRID:SCR_025581)

Description: Software R package to provide ggplot2 implementation of biplot, simultaneous plot of scores for observations and vectors for variables for principal component-like analyses.

Resource Type: source code, software resource, software toolkit

Keywords: ggplot2 implementation of biplot, vectors for variables, variables for principal component-like analyses, simultaneous plot of scores, scores for observations,

Availability: Free, Available for download, Freely available

Resource Name: ggbiplot

Resource ID: SCR_025581

Alternate URLs: <https://github.com/vqv/ggbiplot>

Record Creation Time: 20240806T053245+0000

Record Last Update: 20260728T043726+0000

Ratings and Alerts

No rating or validation information has been found for ggbiplot.

No alerts have been found for ggbiplot.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Chowdhury MMH, et al. (2024) Impact of a tailored exercise regimen on physical capacity and plasma proteome profile in post-COVID-19 condition. *Frontiers in physiology*, 15, 1416639.

Zhao Y, et al. (2024) Distinct molecular profiles drive multifaceted characteristics of colorectal cancer metastatic seeds. *The Journal of experimental medicine*, 221(5).

Toh J, et al. (2024) Multi-modal analysis reveals tumor and immune features distinguishing EBV-positive and EBV-negative post-transplant lymphoproliferative disorders. *Cell reports. Medicine*, 5(12), 101851.

Miller S, et al. (2024) Does ecology shape geographical parthenogenesis? Evidence from the facultatively parthenogenetic stick insect *Megacrana batesii*. *Ecology and evolution*, 14(8), e70145.

Peris D, et al. (2023) Macroevolutionary diversity of traits and genomes in the model yeast genus *Saccharomyces*. *Nature communications*, 14(1), 690.