

# Resource Summary Report

Generated by [ASWG](#) on Aug 3, 2026

## ggbiplot

RRID:SCR\_025581

Type: Tool

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### Proper Citation

ggbiplot (RRID:SCR\_025581)

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### Resource Information

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

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**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 toolkit, software resource

**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:** 20260803T040900+0000

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### Ratings and Alerts

No rating or validation information has been found for ggbiplot.

No alerts have been found for ggbiplot.

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### Data and Source Information

Source: [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 5 mentions in open access literature.

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

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.