

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: software resource, software toolkit, source code

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: 20260821T194316+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

Usage and Citation Metrics

We found 11 mentions in open access literature.

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

Ding C, et al. (2026) Glioblastoma-Secreted C1QL1 Orchestrates Tumor Microtube Expansion and Neural Synaptic Pruning to Drive Malignant Synapse Formation and Recurrence. *Cancer discovery*, 16(6), 1176.

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.

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.

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.

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

Zhao Y, et al. (2023) Colonic stem cells from normal tissues adjacent to tumor drive inflammation and fibrosis in colorectal cancer. *Cell communication and signaling : CCS*, 21(1), 186.

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

Paret C, et al. (2022) GD2 Expression in Medulloblastoma and Neuroblastoma for Personalized Immunotherapy: A Matter of Subtype. *Cancers*, 14(24).

Shen H, et al. (2022) Whole blood transcriptome profiling identifies candidate genes associated with alopecia in male giant pandas (*Ailuropoda melanoleuca*). *BMC genomics*, 23(1), 297.

Palma F, et al. (2022) In vitro and in silico parameters for precise cgMLST typing of *Listeria monocytogenes*. *BMC genomics*, 23(1), 235.

Stamatakis K, et al. (2021) Cyclooxygenase 2 Effector Genes as Potential Inflammation-Related Biomarkers for Colorectal Cancer Circulating Tumor Cells Detection by Liquid Biopsy. *Frontiers in pharmacology*, 12, 806395.