Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 15, 2025

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 toolkit, source code, software resource

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

Funding:

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: 20250412T060810+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 FDI Lab - SciCrunch.org.

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 Megacrania batesii. Ecology and evolution, 14(8), e70145.

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.

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