Resource Summary Report

Generated by NIF on Sep 10, 2026

combi

RRID:SCR_024986
Type: Tool

Proper Citation

combi (RRID:SCR_024986)

Resource Information

URL: https://bioconductor.org/packages/combi/

Proper Citation: combi (RRID:SCR_024986)

Description: Software R package for simultaneous exploration of multiple datasets. Compositional omics model based visual integration.Used to integrate omics data for visualization, with special focus on compositionality.

Synonyms: Compositional Omics Model-Based Integration

Resource Type: software resource, software toolkit

Defining Citation: PMID:33575602

Keywords: integrate omics data for visualization, simultaneous exploration, multiple datasets, visual integration, joint visualization, including sample variables in analysis,

Funding: Johnson and Johnson

Availability: Free, Available for download, Freely available

Resource Name: combi

Resource ID: SCR_024986

License: GPL v2

Record Creation Time: 20240202T050227+0000

Record Last Update: 20260905T133447+0000

Ratings and Alerts

No rating or validation information has been found for combi.

No alerts have been found for combi.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Arikan M, et al. (2024) gNOMO2: a comprehensive and modular pipeline for integrated multi-omics analyses of microbiomes. GigaScience, 13.