

# Resource Summary Report

Generated by [ASWG](#) on Aug 4, 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 toolkit, software resource

**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:** 20260803T040910+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 [ASWG](#) .

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