

Resource Summary Report

Generated by SPARC SAWG on Aug 1, 2026

ade4

RRID:SCR_024259

Type: Tool

Proper Citation

ade4 (RRID:SCR_024259)

Resource Information

URL: <https://cran.r-project.org/web/packages/ade4/index.html>

Proper Citation: ade4 (RRID:SCR_024259)

Description: Software R package for multivariate data analysis. Used for analysis of one-table, two-table, three-table and K-table.

Resource Type: software resource, software toolkit

Keywords: multivariate data analysis,

Availability: Free, Available for download, Freely available,

Resource Name: ade4

Resource ID: SCR_024259

Alternate URLs: <https://sources.debian.org/src/r-cran-ade4/>

Record Creation Time: 20230830T050217+0000

Record Last Update: 20260801T042921+0000

Ratings and Alerts

No rating or validation information has been found for ade4.

No alerts have been found for ade4.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 17 mentions in open access literature.

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

Braga J, et al. (2025) Infant craniofacial diversity in Early Pleistocene Homo. *Nature communications*, 16(1), 4796.

Menozzi E, et al. (2025) Exploring the relationship between GBA1 host genotype and gut microbiome in the GBA1 L444P/WT mouse model: implications for Parkinson's disease pathogenesis. *Frontiers in neuroscience*, 19, 1546203.

Sadia CG, et al. (2024) The impact of agrochemical pollutant mixtures on the selection of insecticide resistance in the malaria vector *Anopheles gambiae*: insights from experimental evolution and transcriptomics. *Malaria journal*, 23(1), 69.

Li Z, et al. (2023) Integrated Human Skin Bacteria Genome Catalog Reveals Extensive Unexplored Habitat-Specific Microbiome Diversity and Function. *Advanced science* (Weinheim, Baden-Wurttemberg, Germany), 10(28), e2300050.

Nodari R, et al. (2023) Computational protocol to perform a spatiotemporal reconstruction of an epidemic. *STAR protocols*, 4(4), 102548.

Stewart A, et al. (2021) Altitudinal Zonation of Green Algae Biodiversity in the French Alps. *Frontiers in plant science*, 12, 679428.

Gopalan-Nair R, et al. (2021) Convergent Rewiring of the Virulence Regulatory Network Promotes Adaptation of *Ralstonia solanacearum* on Resistant Tomato. *Molecular biology and evolution*, 38(5), 1792.

Xiang L, et al. (2020) Gut Microbiotic Features Aiding the Diagnosis of Acute Ischemic Stroke. *Frontiers in cellular and infection microbiology*, 10, 587284.

Brauer CJ, et al. (2020) Recent and rapid anthropogenic habitat fragmentation increases extinction risk for freshwater biodiversity. *Evolutionary applications*, 13(10), 2857.

Chen R, et al. (2019) Transplantation of fecal microbiota rich in short chain fatty acids and butyric acid treat cerebral ischemic stroke by regulating gut microbiota. *Pharmacological research*, 148, 104403.

Feyrer LJ, et al. (2019) Evolutionary impacts differ between two exploited populations of northern bottlenose whale (*Hyperoodon ampullatus*). *Ecology and evolution*, 9(23), 13567.

Prussing C, et al. (2018) Decreasing proportion of *Anopheles darlingi* biting outdoors between long-lasting insecticidal net distributions in peri-Iquitos, Amazonian Peru. *Malaria journal*, 17(1), 86.

Colella S, et al. (2018) Bacteriocyte Reprogramming to Cope With Nutritional Stress in a Phloem Sap Feeding Hemipteran, the Pea Aphid *Acyrtosiphon pisum*. *Frontiers in physiology*, 9, 1498.

Berthier D, et al. (2015) A comparison of phenotypic traits related to trypanotolerance in five west african cattle breeds highlights the value of shorthorn taurine breeds. PLoS one, 10(5), e0126498.

Hodgson JA, et al. (2014) Early back-to-Africa migration into the Horn of Africa. PLoS genetics, 10(6), e1004393.

Klein CC, et al. (2012) Exploration of the core metabolism of symbiotic bacteria. BMC genomics, 13, 438.

Vercken E, et al. (2010) Glacial refugia in pathogens: European genetic structure of anther smut pathogens on *Silene latifolia* and *Silene dioica*. PLoS pathogens, 6(12), e1001229.