

Resource Summary Report

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FactoMineR

RRID:SCR_014602

Type: Tool

Proper Citation

FactoMineR (RRID:SCR_014602)

Resource Information

URL: <http://factominer.free.fr/index.html>

Proper Citation: FactoMineR (RRID:SCR_014602)

Description: Software R package for multivariate analysis which takes into account different types of data structure. Data can be organized in groups of variable, groups of individuals, or into hierarchy of variables.

Synonyms: factominer

Resource Type: data processing software, data analysis software, software application, software resource

Keywords: multivariate, analysis, data, structure, organized, group, variable, individual, hierarchy

Resource Name: FactoMineR

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Alternate URLs: <https://cran.r-project.org/web/packages/FactoMineR/index.html>,
<https://github.com/husson/FactoMineR>

Record Creation Time: 20220129T080321+0000

Record Last Update: 20260801T042732+0000

Ratings and Alerts

No rating or validation information has been found for FactoMineR.

No alerts have been found for FactoMineR.

Data and Source Information

Usage and Citation Metrics

We found 2349 mentions in open access literature.

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

Wang Y, et al. (2026) Machine learning-based unsupervised phenotypic clustering analysis of patients with IgA nephropathy: Distinct therapeutic responses of different groups. Chinese medical journal, 139(1), 83.

Hua L, et al. (2026) Myeloid-Derived CD38 Mediates Age-Related Endometrial Aging Through NAD⁺ Depletion. Aging cell, 25(1), e70356.

Makhmut A, et al. (2026) Spatial proteomics of ovarian cancer precursors delineates early disease changes and drug targets. Molecular systems biology, 22(1), 7.

Škop V, et al. (2026) Partial non-adherence to antidiabetic therapy undermines diabetes management and correlates with treatment complexity: A cross-sectional study using blood plasma analysis. Diabetes, obesity & metabolism, 28(1), 186.

Lazzari N, et al. (2025) Stemness and hybrid epithelial-mesenchymal profiles guide peritoneal dissemination of malignant mesothelioma and pseudomyxoma peritonei. International journal of cancer, 156(1), 201.

Michel L, et al. (2025) Unraveling variability in young children's brain responses to varied sensory stimulations. iScience, 28(11), 113773.

Mohanty R, et al. (2025) Distinct cerebrovascular pathways underlying Alzheimer's disease-related neurodegeneration. Acta neuropathologica, 150(1), 64.

Bleve A, et al. (2025) ROR γ Bridges Cancer-Driven Lipid Dysmetabolism and Myeloid Immunosuppression. Cancer discovery, 15(7), 1505.

Gonzalez-Lozano MA, et al. (2025) EndoMAP.v1 charts the structural landscape of human early endosome complexes. Nature, 643(8070), 252.

Ayabe H, et al. (2025) Cellular crosstalk mediated by TGF- β drives epithelial-mesenchymal transition in patient-derived multi-compartment biliary organoids. Nature communications, 16(1), 6575.

Jousse C, et al. (2025) Identification of a specific set of genes predicting obesity before phenotype appearance. iScience, 28(5), 112377.

Zhang G, et al. (2025) Genomic and transcriptomic analyses of Heteropoda venatoria reveal the expansion of P450 family for starvation resistance in spiders. GigaScience, 14.

Salié H, et al. (2025) Spatial single-cell profiling and neighbourhood analysis reveal the determinants of immune architecture connected to checkpoint inhibitor therapy outcome in hepatocellular carcinoma. Gut, 74(3), 451.

He W, et al. (2025) Semi-quantitative elemental imaging of corrosion products from bioabsorbable Mg vascular implants in vivo. *Bioactive materials*, 43, 225.

Serdar M, et al. (2025) Prenatal inflammation exacerbates hyperoxia-induced neonatal brain injury. *Journal of neuroinflammation*, 22(1), 57.

Renzetti S, et al. (2025) Soluble fibres modulate dough rheology and gluten structure via hydrogen bond density and Flory-Huggins water interaction parameter. *Current research in food science*, 10, 100991.

Tian Y, et al. (2025) Revealing the Transcriptional and Metabolic Characteristics of Sebocytes Based on the Donkey Cell Transcriptome Atlas. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 12(16), e2413819.

Guerrero-Lagunes LA, et al. (2025) The Bioprospecting of *Bixa orellana* L. for the Selection of Characters with Biological Activity. *Metabolites*, 15(2).

Lindberg H, et al. (2025) In-Depth Analysis of Disease Manifestations in Antineutrophil Cytoplasmic Antibody-Associated Vasculitides Identifies Distinct Clinical Phenotypes. *ACR open rheumatology*, 7(3), e70009.

Motawei MI, et al. (2025) Exploring molecular variation and combining ability of local and exotic bread wheat genotypes under well-watered and drought conditions. *PeerJ*, 13, e18994.