

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 analysis software, data processing 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: 20260905T132748+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 2608 mentions in open access literature.

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

Jing M, et al. (2026) Osteoblast-Derived Osteomodulin Suppresses Bone Loss in Osteoporosis by Inhibiting Osteoclastogenesis. FASEB journal : official publication of the Federation of American Societies for Experimental Biology, 40(10), e71934.

Sanks J, et al. (2026) Predicting ecology and hearing sensitivities in Parapontoporia-An extinct long-snouted dolphin. Anatomical record (Hoboken, N.J. : 2007), 309(4), 853.

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.

Huang Q, et al. (2026) Mesenchymal Stem Cell-Derived Extracellular Vesicles Modulate the Course of Peritoneal Inflammation Through Metabolic and Epigenetic Regulation. Advanced science (Weinheim, Baden-Wurttemberg, Germany), 13(4), e08645.

Jain I, et al. (2026) Effect of 2D and 3D ECM and Biomechanical Cues on Human iPSC-Derived Liver Progenitor Cell Differentiation. Advanced healthcare materials, 15(4), e01370.

Gao C, et al. (2026) Enhance the annotation of long noncoding RNAs in the duck genome using multi-tissue transcriptome data. Poultry science, 105(2), 106239.

Luka? NB, et al. (2026) INSL3 Expression in Leydig Cells is a Biomarker for Immunocastration in Boars: Transcriptional Evidence. Andrology, 14(3), 782.

Stahl S, et al. (2026) Aging of the adaptive immune system affects the gut microbiome and systemic levels of vitamin B6. Microbiome, 14(1).

Shi JR, et al. (2026) An ERR?-ZEB1 transcriptional signature predicts survival in triple-negative breast cancers. Molecular medicine (Cambridge, Mass.), 32(1).

Kleinberger M, et al. (2026) Peripheral blood DNA methylation profile and immune checkpoint inhibitor response in cancer patients. ESMO open, 11(5), 106940.

Bruse J, et al. (2026) Hepatic FGF21 is not required for fasting metabolism but guides protein appetite post energy depletion. EMBO reports, 27(12), 3189.

Morelli G, et al. (2026) Traumatic life experiences during critical periods lead to diverse developmental trajectories. Cell reports. Medicine, 7(6), 102798.

Jenickova E, et al. (2026) The faecal bulk heterogeneity: implications for homogenisation and spot-sampling strategies for metabolomic investigations. Metabolomics : Official journal of the Metabolomic Society, 22(3).

Rodríguez-Correa E, et al. (2026) A kinetics-based model of haematopoiesis reveals extrinsic regulation of skewed lineage output from stem cells. *Nature cell biology*, 28(6), 1127.

Bruno V, et al. (2026) High dose of psilocybin induces acute behavioral changes without inducing conditioned place preference in Sprague-Dawley rats. *Journal of psychopharmacology (Oxford, England)*, 40(5), 863.

Porter D, et al. (2026) N1M²U-modified mRNA vaccines break the mold in fish by enhancing innate immune activation. *Molecular therapy. Nucleic acids*, 37(1), 102862.

Fan Y, et al. (2026) Diel-driven variations of leaf dark respiration and metabolite levels in C3 and C4 grasses. *Plant physiology*, 201(2).

Huang S, et al. (2026) Potassium-regulated sucrose and trehalose-6-phosphate metabolism underpins root adaptation to drought in sweet potato (*Ipomoea batatas* (L.) Lam): an integrated transcriptomic and metabolomic study. *BMC plant biology*, 26(1).

Gomez-Escudero J, et al. (2026) Tumour endothelial cell reprogramming orchestrates angiocrine signalling to drive chemoresistance in breast cancer. *Angiogenesis*, 29(3).

Bermejo-Haro MY, et al. (2026) Coordination Between Treg Cells and Bifidobacterium in the Immune-Bacterial Network of Human Colostrum. *Journal of immunology research*, 2026(1), e8677501.