

Resource Summary Report

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maps

RRID:SCR_019296

Type: Tool

Proper Citation

maps (RRID:SCR_019296)

Resource Information

URL: <https://CRAN.R-project.org/package=maps>

Proper Citation: maps (RRID:SCR_019296)

Description: Software R package to draw geographical maps.

Resource Type: data processing software, data visualization software, software application, software resource, software toolkit

Keywords: Maps display, geographical maps drawing

Availability: Free, Available for download, Freely available

Resource Name: maps

Resource ID: SCR_019296

License: GPL-2

Record Creation Time: 20220129T080344+0000

Record Last Update: 20260829T182621+0000

Ratings and Alerts

No rating or validation information has been found for maps.

No alerts have been found for maps.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 34 mentions in open access literature.

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

Hazell Pickering S, et al. (2026) Multimodal epigenetic and enhancer network remodeling shape the transcriptional landscape of human beige adipocytes. *Communications biology*, 9(1), 191.

Wang SY, et al. (2026) Anti-granulocyte macrophage colony-stimulating factor autoantibodies: A rising cause of infectious diseases. *PLoS pathogens*, 22(4), e1014116.

Mori S, et al. (2025) Biodiversity, Phylogeny, and Evolutionary Dynamics of Gall Midges on Japanese Beech Trees. *Ecology and evolution*, 15(7), e71621.

Guo Y, et al. (2025) A haplotype-based evolutionary history of barley domestication. *Nature*, 647(8090), 680.

Berger C, et al. (2025) In situ and in vitro cryo-EM reveal structures of mycobacterial encapsulin assembly intermediates. *Communications biology*, 8(1), 245.

Sadler MC, et al. (2025) Genetic determinants of zinc homeostasis and its role in cardiometabolic diseases. *PLoS genetics*, 21(12), e1011928.

Glemain B, et al. (2024) Estimating SARS-CoV-2 infection probabilities with serological data and a Bayesian mixture model. *Scientific reports*, 14(1), 9503.

Zhao D, et al. (2024) Members of the class Candidatus Ordosarchaeia imply an alternative evolutionary scenario from methanogens to haloarchaea. *The ISME journal*, 18(1).

Wright DJ, et al. (2024) An international estimate of the prevalence of differing visual imagery abilities. *Frontiers in psychology*, 15, 1454107.

Letcher B, et al. (2024) Role for gene conversion in the evolution of cell-surface antigens of the malaria parasite *Plasmodium falciparum*. *PLoS biology*, 22(3), e3002507.

Wiley LK, et al. (2024) Building a vertically integrated genomic learning health system: The biobank at the Colorado Center for Personalized Medicine. *American journal of human genetics*, 111(1), 11.

Speth DR, et al. (2024) Genetic potential for aerobic respiration and denitrification in globally distributed respiratory endosymbionts. *Nature communications*, 15(1), 9682.

Abhari N, et al. (2024) Measuring genetic diversity across populations. *PLoS computational biology*, 20(12), e1012651.

Stormo BM, et al. (2024) Intrinsically disordered sequences can tune fungal growth and the cell cycle for specific temperatures. *Current biology : CB*, 34(16), 3722.

Saha I, et al. (2023) The AAA+ chaperone VCP disaggregates Tau fibrils and generates aggregate seeds in a cellular system. *Nature communications*, 14(1), 560.

Tamaki I, et al. (2023) Phylogenetic, population structure, and population demographic analyses reveal that *Vicia sepium* in Japan is native and not introduced. *Scientific reports*, 13(1), 20746.

Kusmec A, et al. (2023) A genetic tradeoff for tolerance to moderate and severe heat stress in US hybrid maize. *PLoS genetics*, 19(7), e1010799.

Chong F, et al. (2023) High concentrations of floating neustonic life in the plastic-rich North Pacific Garbage Patch. *PLoS biology*, 21(5), e3001646.

Wu D, et al. (2022) Dual genome-wide coding and lncRNA screens in neural induction of induced pluripotent stem cells. *Cell genomics*, 2(11).

Lenton TM, et al. (2022) Resilience of countries to COVID-19 correlated with trust. *Scientific reports*, 12(1), 75.