

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 17, 2026

[rworldmap](#)

RRID:SCR_023394

Type: Tool

Proper Citation

rworldmap (RRID:SCR_023394)

Resource Information

URL: <https://cran.r-project.org/web/packages/rworldmap/>

Proper Citation: rworldmap (RRID:SCR_023394)

Description: Software R package for mapping country data. Enables mapping of country level and gridded user datasets by facilitating joining to modern world maps and offering visualisation options. Country borders are derived from Natural Earth data v 1.4.0.

Resource Type: software resource, software toolkit

Keywords: mapping country data, joining to modern world maps, modern world maps, modern world maps visualisation,

Availability: Free, Available for download, Freely available

Resource Name: rworldmap

Resource ID: SCR_023394

License: GPL v3

Record Creation Time: 20230321T180026+0000

Record Last Update: 20260912T200304+0000

Ratings and Alerts

No rating or validation information has been found for rworldmap.

No alerts have been found for rworldmap.

Data and Source Information

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Delalay G, et al. (2025) The use of scenario tree models in support of animal health surveillance: A scoping review. Preventive veterinary medicine, 234, 106371.

Islam MS, et al. (2024) COVID-19 contact tracing and quarantine policies in the Indo-Pacific Region: A mixed-methods study of experiences of public health professionals. PLOS global public health, 4(5), e0003121.

Jmel H, et al. (2024) Genetic diversity of variants involved in drug response among Tunisian and Italian populations toward personalized medicine. Scientific reports, 14(1), 5842.

Lu L, et al. (2024) West Nile virus spread in Europe: Phylogeographic pattern analysis and key drivers. PLoS pathogens, 20(1), e1011880.

Raman G, et al. (2023) Population Structure and Genetic Diversity Analyses Provide New Insight into the Endemic Species *Aster spathulifolius* Maxim. and Its Evolutionary History. Plants (Basel, Switzerland), 13(1).

Chu R, et al. (2022) Plastome-based phylogeny and biogeography of *Lactuca L.* (Asteraceae) support revised lettuce gene pool categories. Frontiers in plant science, 13, 978417.