

Resource Summary Report

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Modern Applied Statistics with S

RRID:SCR_019125

Type: Tool

Proper Citation

Modern Applied Statistics with S (RRID:SCR_019125)

Resource Information

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

Proper Citation: Modern Applied Statistics with S (RRID:SCR_019125)

Description: Software R package to support functions and datasets for Venables and Ripley MASS. Used for statistical and graphical analysis of data.

Abbreviations: MASS

Resource Type: software resource, software toolkit

Keywords: Statistical analysis, graphical analysis, data analysis, Venables MASS, Ripley MASS, function support

Availability: Free, Freely available

Resource Name: Modern Applied Statistics with S

Resource ID: SCR_019125

License: GPL v3

Record Creation Time: 20220129T080343+0000

Record Last Update: 20260904T000510+0000

Ratings and Alerts

No rating or validation information has been found for Modern Applied Statistics with S.

No alerts have been found for Modern Applied Statistics with S.

Data and Source Information

Usage and Citation Metrics

We found 29 mentions in open access literature.

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

Komrokji SR, et al. (2026) Geospatial Analysis of Accredited Lung Cancer Screening Facilities in Florida Reveals Suboptimal Alignment with High-Risk Populations. *Cancer research communications*, 6(2), 294.

Bellman S, et al. (2026) Predictors of Fatal Outcomes among Pediatric Patients Hospitalized for Rocky Mountain Spotted Fever, Sonora, Mexico, 2004-20241. *Emerging infectious diseases*, 32(2), 180.

Huie JM, et al. (2026) Limb kinematics and morphology improve salamander climbing performance. *The Journal of experimental biology*, 229(11).

Walker AB, et al. (2026) The Neuroanatomy of the Hawaiian Bobtail Squid Juvenile Bacterial Light Organ. *bioRxiv : the preprint server for biology*.

Bourguet P, et al. (2025) Major alleles of CDCA7 shape CG methylation in *Arabidopsis thaliana*. *Nature plants*, 11(12), 2511.

Ospina OE, et al. (2025) spatialGE Is a User-Friendly Web Application That Facilitates Spatial Transcriptomics Data Analysis. *Cancer research*, 85(5), 848.

Do TTN, et al. (2025) Multi-transcriptomics predicts clinical outcome in systemically untreated breast cancer patients with extensive follow-up. *Breast cancer research : BCR*, 27(1), 133.

Caksa S, et al. (2025) Elevated Transglutaminase-2 in SOX10-Deficient Melanoma Promotes Tumor Onset and Decreases Intratumoral CD4+ T Cells. *Cancer research*.

Tan A, et al. (2024) A multi-peak performance landscape for scale biting in an adaptive radiation of pupfishes. *The Journal of experimental biology*, 227(16).

Lubben N, et al. (2024) LRRK2 kinase inhibition reverses G2019S mutation-dependent effects on tau pathology progression. *Translational neurodegeneration*, 13(1), 13.

Chuhma N, et al. (2024) Regional heterogeneity in the membrane properties of mouse striatal neurons. *Frontiers in cellular neuroscience*, 18, 1412897.

Lempidakis E, et al. (2024) Turbulence causes kinematic and behavioural adjustments in a flapping flier. *Journal of the Royal Society, Interface*, 21(212), 20230591.

Vatsa N, et al. (2024) Network analysis of α -synuclein pathology progression reveals p21-activated kinases as regulators of vulnerability. *bioRxiv : the preprint server for biology*.

Koike K, et al. (2023) A quantitative sequencing method using synthetic internal standards including functional and phylogenetic marker genes. *Environmental microbiology reports*, 15(6), 497.

Sutton GJ, et al. (2023) Determining energy expenditure in a large seabird using accelerometry. *The Journal of experimental biology*, 226(23).

Whyte BA, et al. (2023) The role of body size and cuticular hydrocarbons in the desiccation resistance of invasive Argentine ants (*Linepithema humile*). *The Journal of experimental biology*, 226(16).

Weeks K, et al. (2023) Characterization of microbiomic and geochemical compositions across the photosynthetic fringe. *Frontiers in microbiology*, 14, 1176606.

Shahbazi M, et al. (2023) Comprehensive association analysis of speech recognition thresholds after cisplatin-based chemotherapy in survivors of adult-onset cancer. *Cancer medicine*, 12(3), 2999.

Salimi A, et al. (2023) InterOpt: Improved gene expression quantification in qPCR experiments using weighted aggregation of reference genes. *iScience*, 26(10), 107945.

Marabita F, et al. (2022) Multiomics and digital monitoring during lifestyle changes reveal independent dimensions of human biology and health. *Cell systems*, 13(3), 241.