

Resource Summary Report

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tidyverse

RRID:SCR_019186

Type: Tool

Proper Citation

tidyverse (RRID:SCR_019186)

Resource Information

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

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Description: Software tool as collection of R packages designed for data science which share underlying design philosophy, grammar, and data structures. Packages work in harmony because they share common data representations and API design. Used in everyday data analyses.

Synonyms: tidyverse 1.3.0

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

Keywords: data science, R package, data analysis, common data representation

Availability: Free, Available for download, Freely available

Resource Name: tidyverse

Resource ID: SCR_019186

Alternate URLs: <https://www.tidyverse.org/>, <https://github.com/tidyverse/tidyverse>

License: GPL v3

Record Creation Time: 20220129T080343+0000

Record Last Update: 20260804T043711+0000

Ratings and Alerts

No rating or validation information has been found for tidyverse.

No alerts have been found for tidyverse.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 209 mentions in open access literature.

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

Yeki S, et al. (2026) Sex-Specific doublesex Regulation Targeting the Color-Patterning Gene *h* Underlies the Evolution of Wing Sexual Dimorphism in the Harlequin Ladybug *Harmonia axyridis*. *Evolution & development*, 28(1), e70028.

Dunn LN, et al. (2026) Altered hepatic metabolism in Down syndrome. *Cell reports*, 45(1), 116835.

Madsen RR, et al. (2025) Oncogenic PIK3CA corrupts growth factor signaling specificity. *Molecular systems biology*, 21(2), 126.

Li J, et al. (2025) Epidemiological status, development trends, and risk factors of disability-adjusted life years due to diabetic kidney disease: A systematic analysis of Global Burden of Disease Study 2021. *Chinese medical journal*, 138(5), 568.

Rodrigues JA, et al. (2025) Evidence of diversification and geographic separation in a lineage of *Campylobacter jejuni* co-circulating in cattle and humans. *Microbial genomics*, 11(10).

Mayer M, et al. (2025) Deciphering the cellular tumor microenvironment landscape in salivary gland carcinomas using multiplexed imaging mass cytometry. *Journal of experimental & clinical cancer research : CR*, 44(1), 288.

Frigard R, et al. (2025) Daily activity rhythms, sleep and pregnancy are fundamentally related in the Pacific beetle mimic cockroach, *Diploptera punctata*. *The Journal of experimental biology*, 228(15).

Köster T, et al. (2025) At-RS31 orchestrates hierarchical cross-regulation of splicing factors and integrates alternative splicing with TOR-ABA pathways. *The New phytologist*, 247(2), 738.

Strömbäck C, et al. (2025) Positive anticipated affective reactions increase pro-environmental behavior. *iScience*, 28(5), 112389.

Wang T, et al. (2025) Predictive and therapeutic value of the ferroptosis gene *CISD1* in non-small cell lung cancer. *Oncology letters*, 29(5), 235.

Rackoll T, et al. (2025) Patient perspective and research prioritization in rehabilitation after stroke - results from an online survey in Germany. *BMC health services research*, 25(1), 1564.

Olweny G, et al. (2025) Protocol for identifying Mycobacterium tuberculosis infection status through airway microbiome profiling. STAR protocols, 6(1), 103574.

Chiang AC, et al. (2025) Real-World Validity of Tissue-Agnostic Circulating Tumor DNA Response Monitoring in Lung Cancers Treated With Chemotherapy, Immunotherapy, or Targeted Agents. JTO clinical and research reports, 6(9), 100829.

Stefanatos R, et al. (2025) Developmental mitochondrial complex I activity determines lifespan. EMBO reports, 26(8), 1957.

Lynn HM, et al. (2025) Sequential co-assembly reduces computational resources and errors in metagenome-assembled genomes. Cell reports methods, 5(3), 101005.

Bhattacharya D, et al. (2025) Zebrafish model of palmitic acid induced MAFLD recapitulates pathways conserved in mice and humans. Scientific reports, 15(1), 33343.

Ma H, et al. (2025) Temporal trends and disparities in stroke and diabetes mellitus comorbidity-related mortality from 1999 to 2023. Scientific reports, 15(1), 43967.

Li S, et al. (2025) Exosomes originating from neural stem cells undergoing necroptosis participate in cellular communication by inducing TSC2 upregulation of recipient cells following spinal cord injury. Neural regeneration research, 20(11), 3273.

Marín-Garzón NA, et al. (2025) Detection and functional assessment of structural variants using whole-genome re-sequencing data in Nellore cattle. Scientific reports, 15(1), 30364.

Woo MS, et al. (2025) Glia inflammation and cell death pathways drive disease progression in preclinical and early AD. EMBO molecular medicine, 17(11), 3064.