

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

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: 20260821T194143+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 259 mentions in open access literature.

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

Fujiwara K, et al. (2026) Evolutionary adaptation of bacterial proteomes to translation-impeding sequences. The EMBO journal, 45(6), 1957.

Dong C, et al. (2026) Metabolic and inflammatory burden modifies muscle cardiovascular association across aging cohorts within an intrinsic capacity framework. iScience, 29(5), 115477.

Solgama JP, et al. (2026) Pharmacoepidemiology of Oxycodone in the USA: An Observational Study of ARCOS, Medicaid, and Medicare Drug Databases. Pharmacology research & perspectives, 14(1), e70210.

Oberg AL, et al. (2026) Pancreatic Cancer Detection Consortium Biomarker Bakeoff: A Phase II Blinded Biomarker Validation and Panel Discovery Study. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(12), 2438.

Rodriguez Y, et al. (2026) The Ovine Brain as a Model for Human Neurodevelopment: Immunohistochemical Profiling of Brain Maturation Markers in Preterm Lambs. The Journal of comparative neurology, 534(3), e70149.

McDonald TO, et al. (2026) BESTDR Enables Bayesian Quantification of Mechanism-Specific Drug Responses. Cancer research, 86(5), 1286.

Yao J, et al. (2026) Combining plasma biomarkers, clinical parameters, and neuroimaging features for differential diagnosis of Parkinson's disease and atypical parkinsonian syndromes: a multidimensional modeling approach. Frontiers in aging neuroscience, 18, 1727812.

Khodadad CLM, et al. (2026) Microbial community characterization of multi-crop growouts in the XROOTS aeroponic-hydroponic system on the International Space Station. Frontiers in microbiomes, 5, 1779816.

Nahar S, et al. (2026) Co-mutations of CTNNB1 and PTEN drive aggressive tumor progression in endometrial cancer. Disease models & mechanisms, 19(7).

Andersen L, et al. (2026) Cell-Free DNA-Derived Immune Cell Ratios Uncover Cancer-Associated Systemic Changes. Cancer research communications, 6(4), 811.

Zambrano-Harvey AR, et al. (2026) Cancer incidence in the Pijao indigenous population of colombia using administrative health records, 1996-2022. *Scientific reports*, 16(1).

You Y, et al. (2026) Dynamic single-cell transcriptomics reveals Isamp-guided neural network formation in male *S. japonicum* driving female reproduction. *Nature communications*, 17(1), 1602.

Huang Z, et al. (2026) Single-dose psilocybin promotes cell-type-specific changes of neurons in the orbitofrontal cortex. *Neurotherapeutics : the journal of the American Society for Experimental NeuroTherapeutics*, 23(1), e00841.

Thygesen VFDF, et al. (2026) The genomic history of *Streptococcus mutans* from the Mesolithic until modern times. *Genome biology*, 27(1).

Coppola MR, et al. (2026) Zebrafish as a model for Catel-Manzke syndrome-identification and characterization of the zebrafish TGDS ortholog. *The FEBS journal*, 293(8), 2248.

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.

Zhao Z, et al. (2026) CCR5+ CD8+ T Cells Are Associated with Poor Response to PD-1 Blockade Therapy. *International journal of molecular sciences*, 27(11).

Ma L, et al. (2026) Immunogenicity assessment and epitope mapping of the ASFV proteome by profiling serum antibodies with ASFV antigen phage libraries. *Communications biology*, 9(1).

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.

Rigby RE, et al. (2026) Single-cell analysis of signalling and transcriptional responses to type I interferons. *EMBO reports*, 27(9), 2491.