## **Resource Summary Report**

Generated by NIF on May 4, 2025

# tidyverse

RRID:SCR\_019186

Type: Tool

## **Proper Citation**

tidyverse (RRID:SCR\_019186)

#### **Resource Information**

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

**Proper Citation:** tidyverse (RRID:SCR\_019186)

**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, software application, software resource, data processing software, software toolkit

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

Funding:

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: 20250503T060837+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 145 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.

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

Monteiro-Cardoso VF, et al. (2025) The bile acid chenodeoxycholic acid associates with reduced stroke in humans and mice. Journal of lipid research, 66(1), 100712.

Puffet AS, et al. (2025) Validation of the Emotionally Congruent and Incongruent Face-Body Static Set (ECIFBSS). Behavior research methods, 57(1), 41.

Hoyer MJ, et al. (2024) Combinatorial selective ER-phagy remodels the ER during neurogenesis. Nature cell biology, 26(3), 378.

Ghezzi H, et al. (2024) PUPpy: a primer design pipeline for substrain-level microbial detection and absolute quantification. mSphere, 9(7), e0036024.

Sweeney CJ, et al. (2024) Circulating Tumor DNA Assessment for Treatment Monitoring Adds Value to PSA in Metastatic Castration-Resistant Prostate Cancer. Clinical cancer research: an official journal of the American Association for Cancer Research, 30(18), 4115.

Stankovic S, et al. (2024) Genetic links between ovarian ageing, cancer risk and de novo mutation rates. Nature, 633(8030), 608.

Nowak M, et al. (2024) Single-cell Al-based detection and prognostic and predictive value of DNA mismatch repair deficiency in colorectal cancer. Cell reports. Medicine, 5(9), 101727.

Pan ZF, et al. (2024) Protocol for analysis of plasma proteomes from patients with hepatocellular carcinoma receiving combination therapy. STAR protocols, 5(4), 103308.

Emonet A, et al. (2024) Polyploid genome assembly of Cardamine chenopodiifolia. GigaByte (Hong Kong, China), 2024, gigabyte145.

Poch T, et al. (2024) Intergenic risk variant rs56258221 skews the fate of naive CD4+ T cells via miR4464-BACH2 interplay in primary sclerosing cholangitis. Cell reports. Medicine, 5(7), 101620.

Go S, et al. (2024) Tissue-resident natural killer cells support survival in pancreatic cancer through promotion of cDC1-CD8 T activity. eLife, 13.

Köster T, et al. (2024) At-RS31 orchestrates hierarchical cross-regulation of splicing factors and integrates alternative splicing with TOR-ABA pathways. bioRxiv: the preprint server for biology.

Thomas P, et al. (2024) High affinity mAb infusion can enhance maximum affinity maturation during HIV Env immunization. iScience, 27(4), 109495.

Mondal AK, et al. (2024) Selective transcriptomic dysregulation of metabolic pathways in liver and retina by short- and long-term dietary hyperglycemia. iScience, 27(2), 108979.

Perkins MA, et al. (2024) Protocol for programing a degree-to-careers dashboard in R using Posit and Shiny. STAR protocols, 5(1), 102902.

Kawashima S, et al. (2024) Association of gut microbiome with COPD in Japanese male residents: the SESSA study. ERJ open research, 10(1).

Diego JG, et al. (2024) Breakthrough infections by SARS-CoV-2 variants boost cross-reactive hybrid immune responses in mRNA-vaccinated Golden Syrian hamsters. PLoS pathogens, 20(1), e1011805.

Liu D, et al. (2024) Protocol for analyzing plasma cell-free DNA fragment end motifs from ultra-low-pass whole-genome sequencing. STAR protocols, 5(4), 103357.

Ponkilainen V, et al. (2024) Regional variation in low-value musculoskeletal surgery: a nationwide study from the Finnish Care Register. Acta orthopaedica, 95, 553.