

Resource Summary Report

Generated by [NIF](#) on May 4, 2025

[tidyr](#)

RRID:SCR_017102

Type: Tool

Proper Citation

tidyr (RRID:SCR_017102)

Resource Information

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

Proper Citation: tidyr (RRID:SCR_017102)

Description: Software R package for easily tidy data with spread and gather functions. Used for simplifying process of creating tidy data.

Resource Type: software resource, data management software, software application

Keywords: tidy, data, spread, gather, function

Funding:

Availability: Free, Available for download, Freely available

Resource Name: tidyr

Resource ID: SCR_017102

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

License: MIT

Record Creation Time: 20220129T080333+0000

Record Last Update: 20250503T060659+0000

Ratings and Alerts

No rating or validation information has been found for tidyr.

No alerts have been found for tidyr.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 39 mentions in open access literature.

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

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.

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

Loh CA, et al. (2024) High-fidelity, large-scale targeted profiling of microsatellites. Genome research, 34(7), 1008.

Abebe JS, et al. (2024) Nanopore guided annotation of transcriptome architectures. mSystems, 9(7), e0050524.

Klooster IT, et al. (2024) A systematic review on eHealth technology personalization approaches. iScience, 27(9), 110771.

Popovi? L, et al. (2024) Profiling of ERBB receptors and downstream pathways reveals selectivity and hidden properties of ERBB4 antagonists. iScience, 27(2), 108839.

Borgmästars E, et al. (2024) Multi-omics profiling to identify early plasma biomarkers in pre-diagnostic pancreatic ductal adenocarcinoma: a nested case-control study. Translational oncology, 48, 102059.

Sui L, et al. (2024) Multi-proteomics and interactome dataset of tick-borne encephalitis virus infected host cells. Scientific data, 11(1), 1280.

Golov AK, et al. (2024) A genome-wide nucleosome-resolution map of promoter-centered interactions in human cells corroborates the enhancer-promoter looping model. eLife, 12.

Thomas RC, et al. (2024) Second generation lethality in RNAseH2a knockout zebrafish. Nucleic acids research, 52(18), 11014.

Rissi DV, et al. (2024) Comparative Genomics of Different Lifestyle Fungi in Helotiales (Leotiomyces) Reveals Temperature and Ecosystem Adaptations. Journal of fungi (Basel,

Switzerland), 10(12).

Duarte VdS, et al. (2024) Host DNA depletion methods and genome-centric metagenomics of bovine hindmilk microbiome. *mSphere*, 9(1), e0047023.

Abebe JS, et al. (2024) Nanopore Guided Annotation of Transcriptome Architectures. *bioRxiv* : the preprint server for biology.

Popovi? L, et al. (2024) Protocol for identifying properties of ERBB receptor antagonists using the barcoded ERBBprofiler assay. *STAR protocols*, 5(2), 102987.

Kerstens M, et al. (2024) PLETHORA transcription factors promote early embryo development through induction of meristematic potential. *Development (Cambridge, England)*, 151(12).

Frey N, et al. (2024) Endolysosomal TRPML1 channel regulates cancer cell migration by altering intracellular trafficking of E-cadherin and β 1-integrin. *The Journal of biological chemistry*, 300(1), 105581.

Diaz GB, et al. (2024) Maximum velocity and leg-specific ground reaction force production change with radius during flat curve sprinting. *The Journal of experimental biology*, 227(4).

Hauptfeld E, et al. (2024) Integrating taxonomic signals from MAGs and contigs improves read annotation and taxonomic profiling of metagenomes. *Nature communications*, 15(1), 3373.

Rietz M, et al. (2024) Facilitating ambulatory heart rate variability analysis using accelerometry-based classifications of body position and self-reported sleep. *Physiological measurement*, 45(5).

Haufe Y, et al. (2024) Symmetrical Bispyridinium Compounds Act as Open Channel Blockers of Cation-Selective Ion Channels. *ACS pharmacology & translational science*, 7(3), 771.