

Resource Summary Report

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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: data management software, software application, software resource

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: 20250330T061630+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 [FDI Lab - SciCrunch.org](#).

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.

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.

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

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).

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.

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).

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.