

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: software application, software resource, data management software

Keywords: tidy, data, spread, gather, function

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: 20260803T040727+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

Usage and Citation Metrics

We found 62 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Allen AM, et al. (2026) Differential neuronal survival defines a novel axis of sexual dimorphism in the Drosophila brain. *Cell genomics*, 101125.

Meghdadi B, et al. (2026) Digital twins for in vivo metabolic flux estimations in patients with brain cancer. *Cell metabolism*, 38(1), 228.

Davar D, et al. (2025) Combined Targeting of PD-1 and TIM-3 in Patients with Locally Advanced or Metastatic Non-Small Cell Lung Cancer: AMBER Part 2B. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 31(16), 3443.

Christensen JFMM, et al. (2025) Implementing physician-led medication reviews for patients with diabetes and severe mental disorder: A randomized controlled trial. *British journal of clinical pharmacology*, 91(9), 2543.

Sudhindar PD, et al. (2025) Urinary renal epithelial cells can be used for NPHP1 phenotyping and a personalized therapeutic strategy. *Journal of cell science*, 138(20).

Puffet AS, et al. (2025) The role of cognitive load in automatic integration of emotional information from face and body. *Scientific reports*, 15(1), 28184.

Davar D, et al. (2025) Combined Targeting of PD-1 and TIM-3 in Patients with Locally Advanced or Metastatic Melanoma: AMBER Cohorts 1c, 1e, and 2A. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 31(16), 3433.

Gonzalez-Lozano MA, et al. (2025) EndoMAP.v1 charts the structural landscape of human early endosome complexes. *Nature*, 643(8070), 252.

Sümer AP, et al. (2025) Earliest modern human genomes constrain timing of Neanderthal admixture. *Nature*, 638(8051), 711.

Hekimoğlu O, et al. (2025) Genome-wide insights into genetic diversity of endemic and non-endemic *Ixodes ricinus* populations. *Scientific reports*, 15(1), 34199.

Butic AB, et al. (2025) PD-1 regulates CD4⁺ T cell-mediated CD8⁺ T cell responses in the brain to balance viral control and neuroinflammation. *bioRxiv : the preprint server for biology*.

Landi D, et al. (2025) A Checkpoint Reversal Receptor Mediates Bipartite Activation and Enhances CAR T-cell Function. *Cancer research communications*, 5(3), 527.

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

Rokes AB, et al. (2025) History shapes regulatory and evolutionary responses to tigecycline in two reference strains of *Acinetobacter baumannii*. *Microbiology (Reading, England)*, 171(6).

Oliver-Domínguez I, et al. (2025) Memory recognition elicits autonomic-like responses in crayfish. *The Journal of experimental biology*, 228(12).

Burchert HH, et al. (2025) Oxygen dissociation curve inflection point during incremental exercise: a trigger for the Bohr effect. *Pflügers Archiv : European journal of physiology*, 477(8), 1075.

Tenger-Trolander A, et al. (2025) Genomic resources for the scuttle fly *Megaselia abdita*: a model organism for comparative developmental studies in flies. *Development (Cambridge, England)*, 152(22).

Fuhrmann E, et al. (2025) Extended poly(A) tails are a shared feature of herpesvirus mRNAs. *bioRxiv : the preprint server for biology*.

Duarte VDS, et al. (2025) Longitudinal study of the udder microbiome using genome-centric metagenomics uncovers pathogen-driven adaptation and succession. *NPJ biofilms and microbiomes*, 11(1), 227.

Allen AM, et al. (2025) A high-resolution atlas of the brain predicts lineage and birth order underlying neuronal identity. *Cell genomics*, 101103.