

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

stringr

RRID:SCR_022813

Type: Tool

Proper Citation

stringr (RRID:SCR_022813)

Resource Information

URL: <https://stringr.tidyverse.org/>

Proper Citation: stringr (RRID:SCR_022813)

Description: Software package to provide set of functions designed to make working with strings easy. Consistent wrappers for common string operations.

Resource Type: software toolkit, software resource

Keywords: Consistent wrappers, common string operations

Availability: Free, Available for download, Freely available

Resource Name: stringr

Resource ID: SCR_022813

Alternate URLs: <https://CRAN.R-project.org/package=stringr>

License: GPL v2

Record Creation Time: 20221005T050138+0000

Record Last Update: 20260729T044547+0000

Ratings and Alerts

No rating or validation information has been found for stringr.

No alerts have been found for stringr.

Data and Source Information

Usage and Citation Metrics

We found 26 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

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.

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.

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

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

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.

Kremer LS, et al. (2025) The bottleneck for maternal transmission of mtDNA is linked to purifying selection by autophagy. *Science advances*, 11(46), eaea4660.

Li S, et al. (2025) Exosomes originating from neural stem cells undergoing necroptosis participate in cellular communication by inducing TSC2 upregulation of recipient cells following spinal cord injury. *Neural regeneration research*, 20(11), 3273.

Ellegaard KM, et al. (2025) SARS-CoV-2 sequencing artifacts associated with targeted PCR enrichment and read mapping. *PloS one*, 20(10), e0334009.

Vandana JJ, et al. (2025) ChemPerturb-seq screen identifies a small molecule cocktail enhancing human beta cell survival after subcutaneous transplantation. *Cell stem cell*, 32(8), 1299.

Tenedini FM, et al. (2025) Inflammatory cytokine *upd3* induces axon length-dependent synapse removal by glia. *Proceedings of the National Academy of Sciences of the United States of America*, 122(21), e2422752122.

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

Michel L, et al. (2025) Unraveling variability in young children's brain responses to varied sensory stimulations. *iScience*, 28(11), 113773.

Jayasinghe RG, et al. (2025) Single-cell transcriptomic profiling reveals diversity in human iNKT cells across hematologic tissues. *Cell reports*, 44(5), 115587.

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

Bortel P, et al. (2024) Systematic Optimization of Automated Phosphopeptide Enrichment for High-Sensitivity Phosphoproteomics. *Molecular & cellular proteomics : MCP*, 23(5), 100754.

Muse VP, et al. (2024) Protocol for EHR laboratory data preprocessing and seasonal adjustment using R and RStudio. *STAR protocols*, 5(1), 102912.

Kerzel T, et al. (2024) VisualZoneR: A computational protocol to identify compartmental zones from single-cell spatial transcriptomics using R. *STAR protocols*, 5(3), 103196.

Aribi HB, et al. (2024) NeuroVar: an open-source tool for the visualization of gene expression and variation data for biomarkers of neurological diseases. *GigaByte (Hong Kong, China)*, 2024, gigabyte143.

Navarro J, et al. (2024) Social mood during the Covid-19 vaccination process in Spain. A sentiment analysis of tweets and social network leaders. *Heliyon*, 10(1), e23958.