

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](#) on Apr 23, 2025

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 resource, software toolkit

Keywords: Consistent wrappers, common string operations

Funding:

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: 20250422T060304+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

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 12 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

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

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

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.

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

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.

Wang K, et al. (2024) Protocol for fast clonal family inference and analysis from large-scale B cell receptor repertoire sequencing data. *STAR protocols*, 5(2), 102969.

Yan S, et al. (2024) A CC-NB-ARC-LRR Gene Regulates Bract Morphology in Cotton. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 11(44), e2406111.

Sharker Y, et al. (2024) Pairwise Accelerated Failure Time Regression Models for Infectious Disease Transmission in Close-Contact Groups With External Sources of Infection. *Statistics in medicine*, 43(27), 5138.

Lehle JD, et al. (2024) An in vitro approach reveals molecular mechanisms underlying endocrine disruptor-induced epimutagenesis. *eLife*, 13.

Kraus F, et al. (2023) PARK15/FBXO7 is dispensable for PINK1/Parkin mitophagy in

iNeurons and HeLa cell systems. EMBO reports, 24(8), e56399.

van der Vegt I, et al. (2023) A multi-modal panel dataset to understand the psychological impact of the pandemic. Scientific data, 10(1), 537.