Resource Summary Report

Generated by <u>NIF</u> on May 25, 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: 20250525T032609+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 <u>NIF</u>.

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.