

Resource Summary Report

Generated by ASWG on Aug 3, 2026

dplyr

RRID:SCR_016708

Type: Tool

Proper Citation

dplyr (RRID:SCR_016708)

Resource Information

URL: <https://cran.r-project.org/web/packages/dplyr/index.html>

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Description: Software tool for working with data frame like objects, both in memory and out of memory. It is a grammar of data manipulation, providing a consistent set of verbs that help you solve the most common data manipulation challenges: mutate, select, filter, summarise, arrange.

Resource Type: data analysis software, software resource, data processing software, software application

Keywords: data, frame, object, in and of memory, grammar, manipulation, set, verb, mutate, select, filter, summarise, arrange

Availability: Open Source, Available for download

Resource Name: dplyr

Resource ID: SCR_016708

Alternate URLs: <https://github.com/tidyverse/dplyr>, <https://dplyr.tidyverse.org/>

License: MIT

Record Creation Time: 20220129T080332+0000

Record Last Update: 20260803T040722+0000

Ratings and Alerts

No rating or validation information has been found for dplyr.

No alerts have been found for dplyr.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 200 mentions in open access literature.

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

Tomasovic LM, et al. (2026) Mapping the Geographic Distribution of Lung Transplant Centers and Smoking-Associated Diseases Across the United States. Transplantation proceedings.

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

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

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.

Zhou D, et al. (2025) Essential Regulation of Spermatogonial Stem Cell Fate Decisions and Male Fertility by APBB1 via Interaction with KAT5 and GDF15 in Humans and Mice. Research (Washington, D.C.), 8, 0647.

Mohamad-Ramshan R, et al. (2025) Nucleotide level mapping of uracils in murine heavy chain switch regions shows correlation between uracilation and positions of switch junctions created during class-switch recombination. NAR molecular medicine, 2(1), ugaf003.

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

Sharma S, et al. (2025) Taurine from tumour niche drives glycolysis to promote leukaemogenesis. Nature, 644(8075), 263.

O'Connell TM, et al. (2025) Pathway Volcano: an interactive tool for pathway guided visualization of differential expression data. Bioinformatics (Oxford, England), 41(7).

Duan B, et al. (2025) Widespread epistasis shapes RNA polymerase II active site function and evolution. Nature communications, 16(1), 7993.

Uehara T, et al. (2025) Biotin methyl ester enhances cargo release in RUSH system and enables rapid biotinylation with TurboID. Communications biology, 8(1), 1767.

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),

Mondragon-Estrada E, et al. (2025) Protocol to analyze deep-learning-predicted functional scores for noncoding de novo variants and their correlation with complex brain traits. STAR protocols, 6(2), 103738.

Nath R, et al. (2025) Lineage-Specific Class-A GPCR Dynamics Reflect Diverse Chemosensory Adaptations in Lophotrochozoa. Molecular biology and evolution, 42(3).

Song Z, et al. (2025) CTG-Initiated Cryptic Peptide Translation Up- and Downstream of a Canonical ATG Start Codon Is Enhanced by TLR Stimulation and Induces Tumor Regression in Mice. Cancer immunology research, 13(8), 1246.

Frey HC, et al. (2025) A membrane lipid signature unravels the dynamic landscape of group 1 innate lymphoid cells across the health-disease continuum. iScience, 28(3), 112043.

Fraga-González G, et al. (2025) Affording reusable data: recommendations for researchers from a data-intensive project. Scientific data, 12(1), 258.

Wang G, et al. (2025) Development and external validation of a model for post-endoscopic retrograde cholangiopancreatography pancreatitis. iScience, 28(6), 112570.

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).

Huo W, et al. (2025) Integrated Transcriptomics and Metabolomics Provide Insight into Degeneration-Related Molecular Mechanisms of Morchella importuna During Repeated Subculturing. Journal of fungi (Basel, Switzerland), 11(6).