Resource Summary Report

Generated by T1D on Sep 17, 2026

stats

RRID:SCR_025968
Type: Tool

Proper Citation

stats (RRID:SCR_025968)

Resource Information

URL: https://stat.ethz.ch/R-manual/R-devel/library/stats/html/00Index.html

Proper Citation: stats (RRID:SCR_025968)

Description: Software package contains functions for statistical calculations and random number generation.

Resource Type: software application, software resource

Keywords: functions, statistical calculations, random number generation,

Availability: Free, Freely available

Resource Name: stats

Resource ID: SCR_025968

Record Creation Time: 20241024T053301+0000

Record Last Update: 20260912T200454+0000

Ratings and Alerts

No rating or validation information has been found for stats.

No alerts have been found for stats.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 30 mentions in open access literature.

Listed below are recent publications. The full list is available at T1D .

Komrokji SR, et al. (2026) Geospatial Analysis of Accredited Lung Cancer Screening Facilities in Florida Reveals Suboptimal Alignment with High-Risk Populations. Cancer research communications, 6(2), 294.

Yogo A, et al. (2026) Spatial Transcriptomics Reveals Location-Specific Tumor Cell Subtypes and Signaling within Multifocal Small Intestinal Neuroendocrine Tumors. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(10), 2098.

Altendorf L, et al. (2026) Molecular changes during AT/RT progression associated with epithelial-mesenchymal transition and extracellular matrix changes. Acta neuropathologica, 152(1).

De Marco L, et al. (2026) NOUS-209 Off-the-shelf Immunotherapy Has the Potential to Hit Primary and Metachronous Colorectal and Urothelial Cancers in Lynch Syndrome. Molecular cancer therapeutics, 25(4), 650.

Zhang Y, et al. (2026) Co-occurrence of transcriptionally distinct persister cell states underpins neoadjuvant therapy resistance in triple?negative breast cancer. Genome medicine, 18(1).

Gurjao C, et al. (2026) Germline Predisposition to Oncogenic Alkylating Damage in Colorectal Cancer. Cancer epidemiology, biomarkers & prevention : a publication of the American Association for Cancer Research, cosponsored by the American Society of Preventive Oncology, 35(3), 420.

Lorenzo MCC, et al. (2026) Single-Cell Morphometrics From Hematoxylin and Eosin-Stained Images Reveals Subtype-Specific Features in Chronic Lymphocytic Leukemia. Cytometry. Part A : the journal of the International Society for Analytical Cytology, 109(1), 62.

Tabilas C, et al. (2026) Single-cell gene expression and TCR profiling reveal age-related differences in recent thymic emigrants. iScience, 29(5), 115582.

Wild C, et al. (2026) Restoring structural complexity in temperate forests increases bat and bird diversity. Current biology : CB, 36(11), 2840.

Licona Angeles J, et al. (2025) Community Composition of Epibiont Hydroids of the Naturalized Alien Macroalga Acanthophora spicifera in Pichilingue, Mexico. Biology, 14(1).

Alves da Quinta D, et al. (2025) Genetic Ancestry, Intrinsic Tumor Subtypes, and Breast Cancer Survival in Latin American Women. Cancer research communications, 5(7), 1070.

Kisakol B, et al. (2025) High-Resolution Spatial Proteomics Characterises Colorectal Cancer Consensus Molecular Subtypes. bioRxiv : the preprint server for biology.

Bensberg M, et al. (2025) Exposing the DNA methylation-responsive compartment of the leukaemic genome in T-ALL cell lines support its potential as a novel therapeutic target in

T-ALL. Clinical epigenetics, 17(1), 114.

Mier-Quesada Z, et al. (2025) Micro and macro structural brain plastic changes induced by sexual experience in male rats. PloS one, 20(10), e0334959.

van Velthoven CTJ, et al. (2025) Transcriptomic and spatial organization of telencephalic GABAergic neurons. Nature, 647(8088), 143.

Shafer AM, et al. (2025) Immunocompetent Murine Models Recapitulate the Heterogeneous Tumor-Immune Microenvironment of Human Liposarcoma. Clinical cancer research : an official journal of the American Association for Cancer Research, 31(23), 5078.

García-Ruiz S, et al. (2025) Splicing accuracy varies across human introns, tissues, age and disease. Nature communications, 16(1), 1068.

Levé M, et al. (2025) Metabolomics and metagenomics in mice reveal the role of the gut microbiota in tryptophan metabolism. iScience, 28(11), 113751.

Menozzi E, et al. (2025) Exploring the relationship between GBA1 host genotype and gut microbiome in the GBA1 L444P/WT mouse model: implications for Parkinson's disease pathogenesis. Frontiers in neuroscience, 19, 1546203.

Gemünd I, et al. (2025) Characterization of human CMV-specific CD8+ T cells using multi-layer single-cell omics. Cell reports methods, 5(7), 101085.