

Resource Summary Report

Generated by [NIF](#) on Aug 26, 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: 20260821T043137+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 [NIF](#) .

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.

Menozi 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.