

Resource Summary Report

Generated by [NIF](#) on Aug 28, 2026

seaborn

RRID:SCR_018132

Type: Tool

Proper Citation

seaborn (RRID:SCR_018132)

Resource Information

URL: <https://seaborn.pydata.org/>

Proper Citation: seaborn (RRID:SCR_018132)

Description: Software Python tool as data visualization library based on matplotlib. Provides interface for drawing attractive and informative statistical graphics. Statistical data visualization using matplotlib.

Resource Type: data processing software, data visualization software, software application, software library, software resource, software toolkit

Keywords: Data visualization library, statistical graphic, statistical data visualization

Availability: Free, Available for download, Freely available

Resource Name: seaborn

Resource ID: SCR_018132

Alternate URLs: <https://github.com/mwaskom/seaborn/tree/v0.10.0>

License: BSD 3-Clause "New" or "Revised" License

Record Creation Time: 20220129T080338+0000

Record Last Update: 20260821T194125+0000

Ratings and Alerts

No rating or validation information has been found for seaborn.

No alerts have been found for seaborn.

Data and Source Information

Usage and Citation Metrics

We found 479 mentions in open access literature.

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

Arun S, et al. (2026) Recommended Cardiometabolic Screening Guidelines for Unhoused Adults: A Street Medicine Needs Assessment. Clinics and practice, 16(4).

Joshi A, et al. (2026) A multi-agent large language model framework for intelligent vendor evaluation and risk-aware procurement decisions. Scientific reports, 16(1).

Alam MA, et al. (2026) Association between Diagnostic History and Cancer Incidence within 5 Years: A Real-world Observational Analysis. Cancer research communications, 6(5), 1083.

Brima Y, et al. (2026) An empirical study of machine learning robustness and scalability for imbalanced tabular clinical data in emergency and critical care. Scientific reports, 16(1).

Tan Z, et al. (2026) Dissecting the genetic architecture of seed-related traits in Brassica napus by integrating multi-omics analysis and VIS-NIR hyperspectral imaging. Genome biology, 27(1).

Sochodolsky K, et al. (2026) BDNF Regulates Pituitary Stem Cell Engagement towards precursor state. bioRxiv : the preprint server for biology.

Wan M, et al. (2026) A meta learning and task adaptive approach for drug target affinity prediction. Nature communications, 17(1).

Shi Y, et al. (2026) Genomic, Clinical, and Spatial Predictors of Durable Response to BRAF/MEK Inhibition in BRAF-Mutant Melanoma. bioRxiv : the preprint server for biology.

Min J, et al. (2026) Transarterial embolization for chronic wrist pain: a preliminary study of feasibility, safety, and pain reduction. CVIR endovascular, 9(1).

Wang F, et al. (2026) Artificial intelligence-enabled non-invasive cataract diagnosis and grading system using anterior segment images. Chinese medical journal, 139(12), 1888.

Li H, et al. (2026) LeGenD: High-throughput N-glycan profiling using explainable AI and lectin profiling. The Journal of biological chemistry, 302(7), 113234.

Garralda E, et al. (2026) Broad Utility of Ultrasensitive Analysis of ctDNA Dynamics across Solid Tumors Treated with Immunotherapy. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(2), 333.

Elbahnsi A, et al. (2026) Mapping the inhibition landscape of P-glycoprotein via conformational ensemble docking. Scientific reports, 16(1).

Salem M, et al. (2026) Genetic diversity and mobile genetic element associated multidrug resistance in *Salmonella enterica* from broiler chickens in Egypt. *Scientific reports*, 16(1).

??ktem A, et al. (2026) Adaptive responses of genome-reduced *Bacillus subtilis* during enzyme secretion in a breathing vessel culture. *Microbial cell factories*, 25(1).

Daneshnia F, et al. (2026) Profound cell wall remodeling in *Candida parapsilosis* during systemic infection confers simultaneous tolerance to echinocandins and host immunity. *Microbiology spectrum*, 14(4), e0304325.

Liu D, et al. (2026) Predominant Merkel Cell Polyomavirus DNA Detection in Essential Thrombocythemia within Myeloproliferative Neoplasms. *Cancer research communications*, 6(4), 742.

Jiang J, et al. (2026) A CNN-transformer dual-branch network with structure-aware loss for high-resolution edge detection. *Scientific reports*, 16(1).

Zeevi D, et al. (2026) Integrating Multimodal Neuroimaging and Physical-Health Markers for Autism Spectrum Disorder in the ABCD Study. *Journal of integrative neuroscience*, 25(2), 48212.

Wang QS, et al. (2026) Platform-dependent effects of genetic variants on plasma APOL1. *iScience*, 29(3), 114953.