Resource Summary Report

Generated by NIF on May 27, 2025

Plotly

RRID:SCR_013991
Type: Tool

Proper Citation

Plotly (RRID:SCR_013991)

Resource Information

URL: https://plot.ly

Proper Citation: Plotly (RRID:SCR_013991)

Description: Web application to collaboratively create interactive graphs and share them with others. JavaScript toolbox plotly.js enables users with little to no web development skills to make scientific charts.

Resource Type: collaboration tool, software resource

Keywords: collaboration, interactive, tool, graph, plot, chart

Funding: Siemens ;
RealVentures ;
Rho ;
Silicon Valley Bank ;
CNRC-NRC ;
MHS Capital

Availability: Restricted

Resource Name: Plotly

Resource ID: SCR_013991

Alternate URLs: https://plot.ly/product/enterprise/, https://github.com/plotly

License URLs: https://plot.ly/terms-of-service/, https://plot.ly/privacy/

Record Creation Time: 20220129T080318+0000

Record Last Update: 20250527T055340+0000

Ratings and Alerts

No rating or validation information has been found for Plotly.

No alerts have been found for Plotly.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 346 mentions in open access literature.

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

G?sior ?, et al. (2025) Proteomics analysis in rats reveals convergent mechanisms between major depressive disorder and dietary zinc deficiency. Pharmacological reports : PR, 77(1), 145.

Blanco E, et al. (2025) Dominant negative variants in ITPR3 impair T cell Ca2+ dynamics causing combined immunodeficiency. The Journal of experimental medicine, 222(1).

Niraula D, et al. (2025) Intricacies of human-AI interaction in dynamic decision-making for precision oncology. Nature communications, 16(1), 1138.

Saito MA, et al. (2025) Detection of Iron Protein Supercomplexes in Pseudomonas aeruginosa by Native Metalloproteomics. bioRxiv : the preprint server for biology.

Owen CM, et al. (2025) Artificial intelligence driven clustering of blood pressure profiles reveals frailty in orthostatic hypertension. Experimental physiology, 110(2), 230.

Lin V, et al. (2025) TCR3d 2.0: expanding the T cell receptor structure database with new structures, tools and interactions. Nucleic acids research, 53(D1), D604.

Hu ZC, et al. (2025) Evolution of a SHOOTMERISTEMLESS transcription factor binding site promotes fruit shape determination. Nature plants, 11(1), 23.

Romaszko-Wojtowicz A, et al. (2025) Impact of seasonal biometeorological conditions and particulate matter on asthma and COPD hospital admissions. Scientific reports, 15(1), 450.

Zhuo L, et al. (2025) MAPbrain: a multi-omics atlas of the primate brain. Nucleic acids

research, 53(D1), D1055.

Reiter MA, et al. (2024) Dashing Growth Curves: a web application for rapid and interactive analysis of microbial growth curves. BMC bioinformatics, 25(1), 67.

König S, et al. (2024) Observations from the Proteomics Bench. Proteomes, 12(1).

Xiang Q, et al. (2024) The geriatric nutrition risk index is longitudinally associated with incident Sarcopenia: evidence from a 5-year prospective cohort. Aging clinical and experimental research, 36(1), 52.

Felbinger N, et al. (2024) Proscan: a structure-based proline design web server. Nucleic acids research, 52(W1), W280.

Sung PY, et al. (2024) Recruitment of multi-segment genomic RNAs by Bluetongue virus requires a preformed RNA network. Nucleic acids research, 52(14), 8500.

Tsota M, et al. (2024) Investigation of Antihypertensive Properties of Chios Mastic via Monitoring microRNA-21 Expression Levels in the Plasma of Well-Controlled Hypertensive Patients. Non-coding RNA, 10(3).

Gerayeli FV, et al. (2024) Single-cell sequencing reveals cellular landscape alterations in the airway mucosa of patients with pulmonary long COVID. The European respiratory journal, 64(5).

Choi J, et al. (2024) Stemformatics data portal enables transcriptional benchmarking of lab-derived myeloid cells. Stem cell reports, 19(6), 922.

Boutouchent N, et al. (2024) Urogenital colonization and pathogenicity of E. Coli in the vaginal microbiota during pregnancy. Scientific reports, 14(1), 25523.

Scheufen Tieghi R, et al. (2024) A Novel Machine Learning Model and a Web Portal for Predicting the Human Skin Sensitization Effects of Chemical Agents. Toxics, 12(11).

Petrovskiy DV, et al. (2024) Extended range proteomic analysis of blood plasma from schizophrenia patients. Frontiers in molecular biosciences, 11, 1483933.