

Resource Summary Report

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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: 20250416T063653+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.

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.

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

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

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.