

Resource Summary Report

Generated by ASWG on Aug 3, 2026

Plotly

RRID:SCR_013991

Type: Tool

Proper Citation

Plotly (RRID:SCR_013991)

Resource Information

URL: <https://plot.ly>

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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: 20260801T190456+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 441 mentions in open access literature.

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

Rraku E, et al. (2026) Developing Del2Phen: A Novel Phenotype Description Tool for Chromosome Deletions. Human mutation, 2026, 5239482.

Lateef MO, et al. (2026) Diversity, recombination and misclassification in the family Geminiviridae: Insight from bioinformatics analysis. PloS one, 21(1), e0338481.

Meghdadi B, et al. (2026) Digital twins for in vivo metabolic flux estimations in patients with brain cancer. Cell metabolism, 38(1), 228.

Hennion N, et al. (2025) Identification of early genes in the pathophysiology of fibrotic interstitial lung disease in a new model of pulmonary fibrosis. Cellular and molecular life sciences : CMLS, 82(1), 115.

Brenman-Suttner DB, et al. (2025) Exploring the genetics of social behaviour in C. calcarata. Scientific reports, 15(1), 5580.

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

Li Y, et al. (2025) Potential links between human bloodstream infection by Salmonella enterica serovar Typhimurium and international transmission to Colombia. PLoS neglected tropical diseases, 19(1), e0012801.

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

Siam MHB, et al. (2025) CAFE: an integrated web app for high-dimensional analysis and visualization in spectral flow cytometry. Bioinformatics (Oxford, England), 41(6).

Dong J, et al. (2025) MTD: A cloud-based omics database and interactive platform for Myceliophthora thermophila. Synthetic and systems biotechnology, 10(3), 783.

Chen SH, et al. (2025) Unveiling the immune microenvironment of complex tissues and tumors in transcriptomics through a deconvolution approach. BMC cancer, 25(Suppl 1), 733.

Mumme HL, et al. (2025) Identification of leukemia-enriched signature through the development of a comprehensive pediatric single-cell atlas. *Nature communications*, 16(1), 4114.

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).

Sadok N, et al. (2025) Performing the HINTS-exam using a mixed-reality head-mounted display in patients with acute vestibular syndrome: a feasibility study. *Frontiers in neurology*, 16, 1576959.

Chuah A, et al. (2025) EDAmame: interactive exploratory data analyses with explainable models. *Bioinformatics (Oxford, England)*, 41(6).

Eichwald T, et al. (2025) OPIOID-EXPRESSING B CELLS SILENCE TUMOR-INFILTRATING NOCICEPTOR NEURONS. *Research square*.

Litvin U, et al. (2025) Viro3D: a comprehensive database of virus protein structure predictions. *Molecular systems biology*, 21(11), 1599.

Parrondo-Pizarro R, et al. (2025) Enhancing molecular property prediction through data integration and consistency assessment. *Journal of cheminformatics*, 17(1), 163.

Krull NK, et al. (2025) IDBac: an open-access web platform and compendium for the identification of bacteria by MALDI-TOF mass spectrometry. *bioRxiv : the preprint server for biology*.

Sobsey CA, et al. (2025) Characterization of 53 Multiplexed Targeted Proteomics Assays for Verification Studies in Cancer Cell Lines. *Journal of proteome research*, 24(2), 459.