Resource Summary Report

Generated by NIF on May 28, 2025

Google Charts

RRID:SCR_014010
Type: Tool

Proper Citation

Google Charts (RRID:SCR_014010)

Resource Information

URL: https://developers.google.com/chart/

Proper Citation: Google Charts (RRID:SCR_014010)

Description: A web application which allows users to create data visualizations using JavaScript that can be embedded in a web page. Users can load some Google Chart libraries, list the data to be charted, select options to customize their chart, and create a chart object with a personally chosen id. Then, in the web page, users create a

with that id to display the Google Chart. Charts are customizable and interactive, and since they are rendered using HTML5/SVG technology, charts are cross-browser compatibility and cross platform portability to iPhones, iPads and Android.
Resource Type: web application, software resource

Keywords: web application, data visualization, chart, interactive

Funding:

Availability: Free, Public

Resource Name: Google Charts

Resource ID: SCR_014010

License URLs: https://developers.google.com/site-terms, https://www.google.com/intl/en/policies/privacy/

Record Creation Time: 20220129T080318+0000

Record Last Update: 20250528T061136+0000

Ratings and Alerts

No rating or validation information has been found for Google Charts.

No alerts have been found for Google Charts.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 27 mentions in open access literature.

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

Paley S, et al. (2024) The Comparative Genome Dashboard. bioRxiv : the preprint server for biology.

Hauptman N, et al. (2024) AmiCa: Atlas of miRNA-gene correlations in cancer. Computational and structural biotechnology journal, 23, 2277.

Paley S, et al. (2024) The Comparative Genome Dashboard. Frontiers in microbiology, 15, 1447632.

Mahajan P, et al. (2024) IDSL.GOA: gene ontology analysis for interpreting metabolomic datasets. Scientific reports, 14(1), 1299.

Mahajan P, et al. (2023) IDSL.GOA: Gene Ontology Analysis for Metabolomics. bioRxiv : the preprint server for biology.

Seol D, et al. (2022) Microbial Identification Using rRNA Operon Region: Database and Tool for Metataxonomics with Long-Read Sequence. Microbiology spectrum, 10(2),

e0201721.

, et al. (2022) 3DGenBench: a web-server to benchmark computational models for 3D Genomics. Nucleic acids research, 50(W1), W4.

Rios RA, et al. (2021) Country transition index based on hierarchical clustering to predict next COVID-19 waves. Scientific reports, 11(1), 15271.

Mandai S, et al. (2021) Burden of kidney disease on the discrepancy between reasons for hospital admission and death: An observational cohort study. PloS one, 16(11), e0258846.

Joshi M, et al. (2021) Preparing Infographics for Post-publication Promotion of Research on Social Media. Journal of Korean medical science, 36(5), e41.

Kajiya-Kanegae H, et al. (2021) OryzaGenome2.1: Database of Diverse Genotypes in Wild Oryza Species. Rice (New York, N.Y.), 14(1), 24.

Yang N, et al. (2021) Distinct Retrotransposon Evolution Profile in the Genome of Rabbit (Oryctolagus cuniculus). Genome biology and evolution, 13(8).

Ravichandran J, et al. (2021) NeurotoxKb 1.0: Compilation, curation and exploration of a knowledgebase of environmental neurotoxicants specific to mammals. Chemosphere, 278, 130387.

Schwengers O, et al. (2020) ASA3P: An automatic and scalable pipeline for the assembly, annotation and higher-level analysis of closely related bacterial isolates. PLoS computational biology, 16(3), e1007134.

Mochão H, et al. (2020) KiMoSys 2.0: an upgraded database for submitting, storing and accessing experimental data for kinetic modeling. Database : the journal of biological databases and curation, 2020.

Florez H, et al. (2020) Online dashboard and data analysis approach for assessing COVID-19 case and death data. F1000Research, 9, 570.

Abriata LA, et al. (2020) Building blocks for commodity augmented reality-based molecular visualization and modeling in web browsers. PeerJ. Computer science, 6, e260.

Jevšinek Skok D, et al. (2019) The integrative knowledge base for miRNA-mRNA expression in colorectal cancer. Scientific reports, 9(1), 18065.

Togninalli M, et al. (2018) The AraGWAS Catalog: a curated and standardized Arabidopsis thaliana GWAS catalog. Nucleic acids research, 46(D1), D1150.

Sugimoto N, et al. (2018) Genome-wide analysis of the spatiotemporal regulation of firing and dormant replication origins in human cells. Nucleic acids research, 46(13), 6683.