Resource Summary Report

Generated by dkNET on May 19, 2025

clustergrammer

RRID:SCR_015681

Type: Tool

Proper Citation

clustergrammer (RRID:SCR_015681)

Resource Information

URL: http://amp.pharm.mssm.edu/clustergrammer/

Proper Citation: clustergrammer (RRID:SCR_015681)

Description: Clustergrammer is a web-based tool for visualizing and analyzing high-dimensional data as interactive and shareable hierarchically clustered heatmaps. Clustergrammer enables intuitive exploration of high-dimensional data and has several optional biology-specific features.

Resource Type: software tool, data visualization tool

Defining Citation: DOI:10.1038/sdata.2017.151

Keywords: bio.tools

Funding:

Resource Name: clustergrammer

Resource ID: SCR 015681

Alternate IDs: biotools:clustergrammer

Alternate URLs: https://bio.tools/clustergrammer

Record Creation Time: 20220129T080327+0000

Record Last Update: 20250516T054109+0000

Ratings and Alerts

No rating or validation information has been found for clustergrammer.

No alerts have been found for clustergrammer.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 34 mentions in open access literature.

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

Ibe M, et al. (2024) Familial schwannomatosis carrying LZTR1 variant p.R340X with brain tumor: A case report. Molecular genetics and metabolism reports, 40, 101107.

Lee BH, et al. (2024) High-dimensional profiling of regulatory T cells in psoriasis reveals an impaired skin-trafficking property. EBioMedicine, 100, 104985.

Qin S, et al. (2024) Shade responses and resistant mechanisms in Spatholobus suberectus. Heliyon, 10(6), e28077.

Ryan D, et al. (2024) An expanded transcriptome atlas for Bacteroides thetaiotaomicron reveals a small RNA that modulates tetracycline sensitivity. Nature microbiology, 9(4), 1130.

Lee YJ, et al. (2024) A mutation in the low-complexity domain of splicing factor hnRNPA1 linked to amyotrophic lateral sclerosis disrupts distinct neuronal RNA splicing networks. Genes & development, 38(1-2), 11.

Cathomas F, et al. (2024) Circulating myeloid-derived MMP8 in stress susceptibility and depression. Nature, 626(8001), 1108.

Thompson RC, et al. (2023) Molecular states during acute COVID-19 reveal distinct etiologies of long-term sequelae. Nature medicine, 29(1), 236.

Russo S, et al. (2023) Peripheral immune-derived matrix metalloproteinase promotes stress susceptibility. Research square.

Vázquez-Blomquist D, et al. (2023) Gene expression profiling unveils the temporal dynamics of CIGB-300-regulated transcriptome in AML cell lines. BMC genomics, 24(1), 373.

Velletri T, et al. (2022) Single cell-derived spheroids capture the self-renewing subpopulations of metastatic ovarian cancer. Cell death and differentiation, 29(3), 614.

Keller A, et al. (2022) miRNATissueAtlas2: an update to the human miRNA tissue atlas. Nucleic acids research, 50(D1), D211.

Kim KB, et al. (2022) KIX domain determines a selective tumor-promoting role for EP300 and its vulnerability in small cell lung cancer. Science advances, 8(7), eabl4618.

Du R, et al. (2021) Inhibition of ERAD synergizes with FTS to eradicate pancreatic cancer cells. BMC cancer, 21(1), 237.

Geanon D, et al. (2021) A streamlined whole blood CyTOF workflow defines a circulating immune cell signature of COVID-19. Cytometry. Part A: the journal of the International Society for Analytical Cytology, 99(5), 446.

Hodaei M, et al. (2021) Anticancer drug discovery from Iranian Chrysanthemum cultivars through system pharmacology exploration and experimental validation. Scientific reports, 11(1), 11767.

Ahn JH, et al. (2021) TNF? enhances trovafloxacin-induced in vitro hepatotoxicity by inhibiting protective autophagy. Toxicology letters, 342, 73.

Wen H, et al. (2021) Incorporating Differential Gene Expression Analysis with Predictive Biomarkers to Identify Novel Therapeutic Drugs for Fuchs Endothelial Corneal Dystrophy. Journal of ophthalmology, 2021, 5580595.

Xiong S, et al. (2021) Stem cell transplantation rescued a primary open-angle glaucoma mouse model. eLife, 10.

Dong N, et al. (2021) Ion channel profiling of the Lymnaea stagnalis ganglia via transcriptome analysis. BMC genomics, 22(1), 18.

Bisserier M, et al. (2021) Combination Therapy with STAT3 Inhibitor Enhances SERCA2a-Induced BMPR2 Expression and Inhibits Pulmonary Arterial Hypertension. International journal of molecular sciences, 22(17).