Resource Summary Report

Generated by <u>NIF</u> on Apr 26, 2025

<u>jVenn</u>

RRID:SCR_016343 Type: Tool

Proper Citation

jVenn (RRID:SCR_016343)

Resource Information

URL: http://jvenn.toulouse.inra.fr/app/example.html

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Description: JavaScript plug-in software application for web environments to analyze data. It is an interactive Venn diagram viewer.Used for comparing lists with Venn Diagrams. It handles up to six input lists and presents results using classical or Edwards-Venn layouts. User interactions can be controlled and customized.

Resource Type: data processing software, data analysis software, data visualization software, software resource, software application

Keywords: analyze, data, Venn, diagram, visualisation, independent, platform

Funding:

Availability: Free, Available for download, Freely available

Resource Name: jVenn

Resource ID: SCR_016343

Alternate URLs: http://bioinfo.genotoul.fr/jvenn

License: GNU GPL

Record Creation Time: 20220129T080330+0000

Record Last Update: 20250426T060538+0000

Ratings and Alerts

No rating or validation information has been found for jVenn.

No alerts have been found for jVenn.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 214 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>NIF</u>.

Škandík M, et al. (2025) Age-associated microglial transcriptome leads to diminished immunogenicity and dysregulation of MCT4 and P2RY12/P2RY13 related functions. Cell death discovery, 11(1), 16.

Cao W, et al. (2025) Prognostic values of intracellular cell-related genes in esophageal cancer and their regulatory mechanisms. BMC cancer, 25(1), 105.

Dong Z, et al. (2025) A KSHV-targeted small molecule efficiently blocks SARS-CoV-2 infection via inhibiting expression of EGFR and Cyclin A2. Emerging microbes & infections, 14(1), 2440490.

Yang Z, et al. (2025) Study on the In Vitro and In Vivo Antioxidant Activity and Potential Mechanism of Polygonum viviparum L. Antioxidants (Basel, Switzerland), 14(1).

Wang C, et al. (2024) Integrated chemical analysis, metabolic profiling, network pharmacology, molecular docking and toxicity prediction to reveal the active ingredients and their safety of raw and prepared rhubarbs in the treatment of gastric ulcers. Frontiers in pharmacology, 15, 1481091.

Zhu BT, et al. (2024) Estrogen: the forgotten player in metaflammation. Frontiers in pharmacology, 15, 1478819.

Wang C, et al. (2024) Molecular mechanisms regulating glucose metabolism in quinoa (Chenopodium quinoa Willd.) seeds under drought stress. BMC plant biology, 24(1), 796.

Wang Y, et al. (2024) Natural hydrogen gas and engineered microalgae prevent acute lung injury in sepsis. Materials today. Bio, 28, 101247.

Li C, et al. (2024) Genetic relationship analysis and core collection construction of Eucalyptus grandis from Dongmen improved variety base: the largest eucalypt germplasm resource in China. BMC plant biology, 24(1), 1240.

Zheng L, et al. (2024) Expression of overall survival-EMT-immune cell infiltration genes predict the prognosis of glioma. Non-coding RNA research, 9(2), 407.

Shen R, et al. (2024) Alisol A, the Eye-Entering Ingredient of Alisma orientale, Relieves Macular Edema Through TNF-? as Revealed by UPLC-Triple-TOF/MS, Network Pharmacology, and Zebrafish Verification. Drug design, development and therapy, 18, 3361.

Chen N, et al. (2024) Exploring the mechanism of agarwood moxa smoke in treating sleep disorders based on GC-MS and network pharmacology. Frontiers in medicine, 11, 1400334.

Sakarin S, et al. (2024) Proteomic analysis of the serum in dogs with pulmonary hypertension secondary to myxomatous mitral valve disease: the preliminary study. Frontiers in veterinary science, 11, 1327453.

Monti M, et al. (2024) Novel cellular systems unveil mucosal melanoma initiating cells and a role for PI3K/Akt/mTOR pathway in mucosal melanoma fitness. Journal of translational medicine, 22(1), 35.

Zhong Z, et al. (2024) Exploring shared molecular signatures and regulatory mechanisms in nonalcoholic steatohepatitis and inflammatory bowel disease using integrative bioinformatics analysis. Scientific reports, 14(1), 12085.

Zhu B, et al. (2024) Identification of Key Genes Related to Skin Burns Based on Bioinformatics Analysis. Journal of burn care & research : official publication of the American Burn Association, 45(5), 1183.

Li D, et al. (2024) Comprehensive bioinformatics analysis and systems biology approaches to identify the interplay between COVID-19 and pericarditis. Frontiers in immunology, 15, 1264856.

Xie X, et al. (2024) Genome-wide characterization and metabolite profiling of Cyathus olla: insights into the biosynthesis of medicinal compounds. BMC genomics, 25(1), 618.

Wang X, et al. (2024) Hypermethylation of CDKN2A CpG island drives resistance to PRC2 inhibitors in SWI/SNF loss-of-function tumors. Cell death & disease, 15(11), 794.

Shu X, et al. (2024) Screening of pathologically significant diagnostic biomarkers in tears of thyroid eye disease based on bioinformatic analysis and machine learning. Frontiers in cell and developmental biology, 12, 1486170.