

Resource Summary Report

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jVenn

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

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

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: 20260804T043620+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 248 mentions in open access literature.

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

Li X, et al. (2026) Coinfection with fowl adenovirus serotypes 1 and 4 (FAdV-1 and -4) enhances FAdV-4 replication through FAdV-1-mediated upregulation of HSPA2 expression. *Virulence*, 17(1), 2605751.

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.

He W, et al. (2025) SKSR1 identified as key virulence factor in *Cryptosporidium* by genetic crossing. *Nature communications*, 16(1), 4694.

Xiao Y, et al. (2025) Causal Association Between Lung Cancer and Rash: A Bidirectional Mendelian Randomization Study. *Clinical Medicine Insights. Oncology*, 19, 11795549251341559.

Guo Y, et al. (2025) The mechanisms of environmental pollutant acetyl tributyl citrate induced oral squamous cell carcinoma using network toxicology, molecular docking and molecular dynamics simulation. *International journal of surgery (London, England)*, 111(11), 7873.

Özkan H, et al. (2025) Expression Patterns of miR-29b and miR-148a in Colostrum of Awassi Sheep With Higher Immunoglobulin G Levels. *Veterinary medicine and science*, 11(3), e70335.

Zhou L, et al. (2025) Integrating machine learning and genetic evidence to uncover novel gene biomarkers for colorectal cancer diagnosis. *Discover oncology*, 16(1), 675.

Li G, et al. (2025) The mechanism of Shancigu and its monomer in the development of colorectal cancer based on network pharmacology. *Scientific reports*, 15(1), 23186.

Man Y, et al. (2025) Molecular identification of novel duck associated Chapparvovirus in ducks, first report from China. *Poultry science*, 104(5), 104984.

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

He J, et al. (2025) Investigation of the relationship between chronic hepatitis B and tuberculosis using bioinformatics and systems biology approaches. *Frontiers in medicine*, 12, 1519216.

Mao K, et al. (2025) A cross-sectional and bioinformatics-based analysis: perirenal fat thickness as a superior predictor of kidney stone disease. *Lipids in health and disease*, 24(1), 269.

Han GP, et al. (2025) Transcriptomic analysis of the breast muscle at different ages during the grower phase in broilers. *Poultry science*, 104(12), 105968.

Lin T, et al. (2025) Identification of RNA processing factor LSM5 as a new adverse biomarker in nasopharyngeal carcinoma. *Scientific reports*, 15(1), 9901.

Guo J, et al. (2025) Network Pharmacology Analysis and In Vitro Validation of the Active Ingredients and Potential Mechanisms of *Gynostemma Pentaphyllum* Against Esophageal Cancer. *Combinatorial chemistry & high throughput screening*, 28(3), 500.

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.

Wang Y, et al. (2025) Molecular mechanism underlying the protective effects of ischemic preconditioning in total knee arthroplasty. *Chinese journal of traumatology = Zhonghua chuang shang za zhi*, 28(4), 257.

Wu Y, et al. (2025) Inactivation of β -1,3-glucan synthase-like 5 confers broad-spectrum resistance to *Plasmodiophora brassicae* pathotypes in cruciferous plants. *Nature genetics*, 57(9), 2302.

Stetsenko V, et al. (2025) Human CD4⁺ T cells recognize *Mycobacterium tuberculosis*-infected macrophages amid broader responses. *The Journal of experimental medicine*, 222(12).

Hsieh LT, et al. (2025) The *Mycobacterium ulcerans* toxin mycolactone causes destructive Sec61-dependent loss of the endothelial glycocalyx and vessel basement membrane to drive skin necrosis. *eLife*, 12.