

Resource Summary Report

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GeneVenn

RRID:SCR_012117

Type: Tool

Proper Citation

GeneVenn (RRID:SCR_012117)

Resource Information

URL: <http://genevenn.sourceforge.net/>

Proper Citation: GeneVenn (RRID:SCR_012117)

Description: A web application creating Venn diagrams from two or three gene lists.

Resource Type: analysis service resource, service resource, data analysis service, production service resource

Defining Citation: [PMID:17597932](#)

Keywords: web app

Resource Name: GeneVenn

Resource ID: SCR_012117

Alternate IDs: OMICS_05568

Record Creation Time: 20220129T080308+0000

Record Last Update: 20260728T043402+0000

Ratings and Alerts

No rating or validation information has been found for GeneVenn.

No alerts have been found for GeneVenn.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 101 mentions in open access literature.

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

Mayyas A, et al. (2025) Deciphering the Anti-Diabetic Potential of Gymnema Sylvestre Using Integrated Computer-Aided Drug Design and Network Pharmacology. *Journal of cellular and molecular medicine*, 29(1), e70349.

Nagarajan V, et al. (2025) SCassist: An AI Based Workflow Assistant for Single-Cell Analysis. *bioRxiv : the preprint server for biology*.

Liu B, et al. (2025) Single-Cell and Spatial Transcriptomics Identified Fatty Acid-Binding Proteins Controlling Endothelial Glycolytic and Arterial Programming in Pulmonary Hypertension. *Arteriosclerosis, thrombosis, and vascular biology*, 45(7), 1145.

Jinesh G, et al. (2025) CEBPB, C19MC, and Defective Autophagy Drive Novel Podosomal Belt to Macropinocytosis Transition, Lipid Accumulation, and HBV A-to-I RNA-editing. *Research square*.

Ortabozkoyun H, et al. (2024) Members of an array of zinc-finger proteins specify distinct Hox chromatin boundaries. *Molecular cell*, 84(18), 3406.

Balestrini PA, et al. (2024) Transcription factor-based transdifferentiation of human embryonic to trophoblast stem cells. *Development (Cambridge, England)*, 151(17).

Sukhija N, et al. (2024) Genome-wide selection signatures address trait specific candidate genes in cattle indigenous to arid regions of India. *Animal biotechnology*, 35(1), 2290521.

Szakats S, et al. (2024) Identification of novel microRNAs in the embryonic mouse brain using deep sequencing. *Molecular and cellular biochemistry*, 479(2), 297.

Caballero-Valderrama MR, et al. (2024) Early Myocardial Strain Reduction and miR-122-5p Elevation Associated with Interstitial Fibrosis in Anthracycline-Induced Cardiotoxicity. *Biomedicines*, 13(1).

Jinesh GG, et al. (2024) C19MC drives nucleolar invasion of mitochondria and meiotic nuclear division in human cancers. *iScience*, 27(11), 111132.

Vaikakkara Chithran A, et al. (2023) Adult expression of Semaphorins and Plexins is essential for motor neuron survival. *Scientific reports*, 13(1), 5894.

Jones C, et al. (2023) HNF4 α Acts as Upstream Functional Regulator of Intestinal Wnt3 and Paneth Cell Fate. *Cellular and molecular gastroenterology and hepatology*, 15(3), 593.

Tomaž Š, et al. (2023) A mini-TGA protein modulates gene expression through heterogeneous association with transcription factors. *Plant physiology*, 191(3), 1934.

Jaime OG, et al. (2023) SIX1+PAX3+ identify a progenitor for myogenic lineage commitment from hPSCs. *Development (Cambridge, England)*, 150(14).

Kain BN, et al. (2023) Hematopoietic stem and progenitor cells confer cross-protective trained immunity in mouse models. *iScience*, 26(9), 107596.

Cowell LM, et al. (2023) Regulation of gene expression downstream of a novel Fgf/Erk pathway during *Xenopus* development. *PloS one*, 18(10), e0286040.

Zhang Y, et al. (2023) MicroRNA-142-3p promotes renal cell carcinoma progression by targeting RhoBTB3 to regulate HIF-1 signaling and GGT/GSH pathways. *Scientific reports*, 13(1), 5935.

Maddali P, et al. (2023) Induction of pro-inflammatory genes by fibronectin DAMPs in three fibroblast cell lines: Role of TAK1 and MAP kinases. *PloS one*, 18(5), e0286390.

Hussein M, et al. (2023) Integrated Transcriptomic and Metabolomic Mapping Reveals the Mechanism of Action of Ceftazidime/Avibactam against Pan-Drug-Resistant *Klebsiella pneumoniae*. *ACS infectious diseases*, 9(12), 2409.

Cibichakravarthy B, et al. (2022) Comparative Proteomics of *Coxiella* like Endosymbionts (CLEs) in the Symbiotic Organs of *Rhipicephalus sanguineus* Ticks. *Microbiology spectrum*, 10(1), e0167321.