

Resource Summary Report

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[ggvenn](#)

RRID:SCR_025300

Type: Tool

Proper Citation

ggvenn (RRID:SCR_025300)

Resource Information

URL: <https://CRAN.R-project.org/package=ggvenn>

Proper Citation: ggvenn (RRID:SCR_025300)

Description: Software R package to draw venn diagram by ggplot2.

Resource Type: source code, software resource, software toolkit

Keywords: draw venn diagram,

Availability: Free, Available for download, Freely available

Resource Name: ggvenn

Resource ID: SCR_025300

License: MIT license

Record Creation Time: 20240502T053246+0000

Record Last Update: 20260728T043722+0000

Ratings and Alerts

No rating or validation information has been found for ggvenn.

No alerts have been found for ggvenn.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 18 mentions in open access literature.

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

Musillo C, et al. (2025) Maternal stressors disrupt mouse placental proteome and fetal brain development in a sex-specific fashion through inflammation and oxidative stress. *Molecular psychiatry*, 30(11), 5072.

Liu Y, et al. (2025) Uncovering potential biomarkers of endometriosis: transcriptomic and single-cell analysis. *Frontiers in medicine*, 12, 1528434.

Pessina P, et al. (2025) Selective RNA sequestration in biomolecular condensates directs cell fate transitions. *bioRxiv : the preprint server for biology*.

Jensen A, et al. (2025) Inflammation and response to bacterial infection as potential drivers of equine odontoclastic tooth resorption and hypercementosis: A proteomics insight. *Equine veterinary journal*, 57(4), 977.

Hailati J, et al. (2025) Targeting PDCD4 in cancer and atrial fibrillation: mechanistic insights from integrated multi-omics and single-cell analysis. *Frontiers in oncology*, 15, 1593815.

Zeng XX, et al. (2025) Identification of biomarkers associated with proliferation and differentiation of mesenchymal stem cells in pulmonary adenocarcinoma and establishment of prognostic models. *Hereditas*, 162(1), 118.

Peng G, et al. (2025) Identification of genes related to the effects of bicalutamide and zoladex on benign prostatic hyperplasia. *Scientific reports*, 15(1), 32901.

Ordóñez A, et al. (2025) Evaluating agar-plating and dilution-to-extinction isolation methods for generating oak-associated microbial culture collections. *ISME communications*, 5(1), ycaf019.

Qian D, et al. (2025) Identification and validation of microglia-associated genes in ischemic stroke using single-cell and bulk RNA-seq. *Molecular brain*, 18(1), 91.

Wu J, et al. (2025) Prognostic value assessment and in vitro validation of mitochondria-ferroptosis-related genes in multiple myeloma. *Scientific reports*, 15(1), 41548.

Sun M, et al. (2025) Transcriptome combined with Mendelian randomization to identify key genes related to polyamine metabolism in childhood obesity and elucidate their molecular regulatory mechanisms. *Scientific reports*, 15(1), 17799.

Zhang J, et al. (2025) Identification and validation of prognostic genes associated with m6A-regulated programmed cell death in acute lymphoblastic leukemia. *Scientific reports*, 15(1), 38248.

Deffner M, et al. (2024) Chemokine-mediated cell migration into the central nervous system in progressive multifocal leukoencephalopathy. *Cell reports. Medicine*, 5(7), 101622.

Wang L, et al. (2024) Targeting the HSP47-collagen axis inhibits brain metastasis by reversing M2 microglial polarization and restoring anti-tumor immunity. *Cell reports. Medicine*, 5(5), 101533.

Cheung A, et al. (2024) Anti-EGFR Antibody-Drug Conjugate Carrying an Inhibitor Targeting CDK Restricts Triple-Negative Breast Cancer Growth. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 30(15), 3298.

Kerstens M, et al. (2024) PLETHORA transcription factors promote early embryo development through induction of meristematic potential. *Development (Cambridge, England)*, 151(12).

Wang N, et al. (2024) TFE3 and TP53 were novel diagnostic biomarkers related to mitochondrial autophagy in chronic rhinosinusitis with nasal polyps. *Frontiers in genetics*, 15, 1423778.

Lehle JD, et al. (2024) An in vitro approach reveals molecular mechanisms underlying endocrine disruptor-induced epimutagenesis. *eLife*, 13.