

Resource Summary Report

Generated by [RRID](#) on Aug 15, 2026

[circlize](#)

RRID:SCR_002141

Type: Tool

Proper Citation

circlize (RRID:SCR_002141)

Resource Information

URL: <http://cran.r-project.org/web/packages/circlize/>

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Description: Software package that implements and enhances circular visualization in R. Due to natural born feature of R to draw statistical graphics, this package can provide more general and flexible way to visualize huge information in circular style.

Synonyms: circlize: Circular Visualization in R

Resource Type: software application, standalone software, software resource

Defining Citation: [PMID:24930139](#)

Keywords: circular visualization, draw statistical graphics, circular style graphics, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: circlize

Resource ID: SCR_002141

Alternate IDs: OMICS_04658, biotools:circlize

Alternate URLs: <https://github.com/jokergoo/circlize>, <https://bio.tools/circlize>

License: GNU General Public License, v2, v3

Record Creation Time: 20220129T080211+0000

Record Last Update: 20260815T042427+0000

Ratings and Alerts

No rating or validation information has been found for circlize.

No alerts have been found for circlize.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1289 mentions in open access literature.

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

Zhao C, et al. (2026) Progranulin promotes glioma progression via interaction with cathepsin D and serves as a diagnostic and prognostic biomarker. *Discover oncology*, 17(1), 336.

Avolio M, et al. (2026) The Molecular and Functional Landscape of Resistance to FOLFIRI Chemotherapy in Metastatic Colorectal Cancer. *Cancer discovery*, 16(2), 270.

Schulz A, et al. (2026) LAMA5 links extracellular matrix organization to a candidate WNT-associated endothelial signaling niche during human chondrogenesis. *iScience*, 29(8), 116687.

Jin K, et al. (2026) Bibliometric and visualized analysis of the therapeutic application of suprachoroidal space from 2000 to 2024. *International journal of surgery (London, England)*, 112(1), 1635.

Wang JY, et al. (2026) *Fusobacterium nucleatum* manipulates host autophagy to promote its intracellular survival and treatment resistance in nasopharyngeal carcinoma. *Molecular cancer*, 25(1).

Sun J, et al. (2026) B cell imprinting in children impairs antibodies to the haemagglutinin stalk. *Nature*, 653(8114), 528.

Zhang Z, et al. (2026) Systematic identification of cell-cell interactions associated with the severity of patients with Alzheimer's disease. *Journal of Alzheimer's disease : JAD*, 111(3), 1156.

Venkatesan S, et al. (2026) Cell type-agnostic transcriptomic signatures enable uniform comparisons of neural maturation. *PLoS biology*, 24(4), e3003757.

Ward MH, et al. (2026) Metabolites from plasma-like medium fuel nitrogen metabolism and influence proliferation in *Leptospira interrogans*. *bioRxiv : the preprint server for biology*.

Zhong J, et al. (2026) Transcriptomic and proteomic profiling of pulmonary mucormycosis reveals a failed activation of host immune response. *Frontiers in immunology*, 17, 1732782.

de Gennaro L, et al. (2026) Evolution of rDNA-Linked Segmental Duplications as Lineage-Specific Mosaics in Great Apes. *Genes*, 17(2).

Selvatici S, et al. (2026) Genomic identification and complete mitochondrial recovery of a Late Holocene porcupine (*Erethizon dorsatum*) mummy from Yukon permafrost. *Scientific reports*, 16(1).

Peng SL, et al. (2026) CS2164 combined with radiation suppressed tumor growth via ??-H2AX persistence and ROS accumulation. *BMC cancer*, 26(1).

Bellomo SE, et al. (2026) Genome-wide biomarker analysis across the full spectrum of HER2-expressing breast cancers to reveal a clonal chromosome 17 imbalance defining unfavourable HER2-low disease. *Biomarker research*, 14(1).

Krey K, et al. (2026) The ISG Atlas: a loss-of-function analysis characterizes antiviral properties of interferon stimulated genes. *Nature communications*, 17(1).

Gonzalez Mateu M, et al. (2026) Functional and ecological drivers of bacterial interactions in glacier-fed stream biofilms. *mSystems*, 11(6), e0037026.

Jain R, et al. (2026) Insulin signaling engages divergent transcriptional mechanisms in neural and metabolic tissues. *BMC molecular and cell biology*, 27(1).

Chen T, et al. (2026) Multi-Omics Data Reveal Estrogen-Driven Dysregulation and Stromal-Epithelial Signaling Alterations in Endometrial Polyps. *FASEB journal : official publication of the Federation of American Societies for Experimental Biology*, 40(5), e71645.

Diaz M, et al. (2026) Novel driver gene MDC1 confers homologous recombination repair deficiency and genomic instability in chemoresistant relapsing ovarian cancer. *Journal of translational medicine*, 24(1).

Nunes de Almeida F, et al. (2026) A forward genetic screen identifies Sirtuin1 as a driver of neuroendocrine prostate cancer. *The Journal of experimental medicine*, 223(7).