

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/>

Proper Citation: circlize (RRID:SCR_002141)

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.

Makhmut A, et al. (2026) Spatial proteomics of ovarian cancer precursors delineates early disease changes and drug targets. *Molecular systems biology*, 22(1), 7.

Luo H, et al. (2026) EasyAmplicon 2: Expanding PacBio and Nanopore Long Amplicon Sequencing Analysis Pipeline for Microbiome. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 13(1), e12447.

Zhu Y, et al. (2026) Single-cell RNA sequencing unveils enhanced antitumor immunity in colorectal cancer with PD-1 blockade and LINC00673 deletion. *Biomarker research*, 14(1), 16.

Schoof M, et al. (2026) Investigation of a global mouse methylome atlas reveals subtype-specific copy number alterations in pediatric cancer models. *Nature genetics*, 58(1), 143.

Shi J, et al. (2026) Nap1L4a Cooperates with Scl/Klf1 to Recruit H2A.Z in Mediating Interactions Among Cis-Regulatory Elements and Transcription Required for Primitive Erythropoiesis in Zebrafish. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 13(12), e13762.

Perdios C, et al. (2026) RhCMV expands CCR5+ memory T cells and promotes SIV reservoir seeding in the gut mucosa. *JCI insight*, 11(1).

Singh P, et al. (2026) Loss of KDM6A-mediated genomic instability and metabolic reprogramming regulates response to therapeutic perturbations in bladder cancer. *Nature communications*, 17(1), 1382.

Backenstose NJC, et al. (2026) Delayed transcriptional response of *Daphnia pulex* to thermal stress. *G3 (Bethesda, Md.)*, 16(3).

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

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).

Chan AHP, et al. (2026) Developing a Single-Cell Spatial Transcriptomics Workflow for In Vivo Evaluation of Implanted Biomaterials. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 13(24), e13242.

Leong MK, et al. (2026) From fish to invertebrates: multi-marker eDNA metabarcoding for monitoring wetland biodiversity and non-indigenous species in Macao SAR China. *Scientific reports*, 16(1).

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

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.

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.

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