

Resource Summary Report

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[circIncRNAnet](#)

RRID:SCR_015794

Type: Tool

Proper Citation

circIncRNAnet (RRID:SCR_015794)

Resource Information

URL: <http://app.cgu.edu.tw/circInc/>

Proper Citation: circIncRNAnet (RRID:SCR_015794)

Description: Web application for mapping functional networks of long or circular forms of non-coding RNAs. It supports the uploading and processing of user-defined NGS-based gene expression matrix data.

Resource Type: data or information resource, data processing resource, database, software resource, web application

Keywords: mapping, long form, circular form, non-coding rna, rna mapping, regulatory rna, ncRNA, bio.tools, FASEB list

Availability: Freely available, Free, Available for download, Runs on Linux, Demo available, Tutorial available

Resource Name: circIncRNAnet

Resource ID: SCR_015794

Alternate IDs: biotools:circIncrnanet

Alternate URLs: <https://github.com/smw1414/circIncRNAnet>,
<https://bio.tools/circIncrnanet>

Record Creation Time: 20220129T080327+0000

Record Last Update: 20260821T194027+0000

Ratings and Alerts

No rating or validation information has been found for circIncRNAnet.

No alerts have been found for circIncRNA.net.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 33 mentions in open access literature.

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

Jihen S, et al. (2026) Filter bank CSP with Riemannian weighting for disability-centric motor imagery brain computer interface. *Brain informatics*, 13(1).

Wang B, et al. (2023) A systematic review and meta-analysis of the association between HOTAIR polymorphisms and susceptibility to breast cancer. *Archives of medical science : AMS*, 19(1), 128.

Tang LH, et al. (2023) LINC01268 promotes epithelial-mesenchymal transition, invasion and metastasis of gastric cancer via the PI3K/Akt signaling pathway and targeting MARCKS. *World journal of gastrointestinal oncology*, 15(8), 1366.

Hu F, et al. (2023) Circular RNAs: implications of signaling pathways and bioinformatics in human cancer. *Cancer biology & medicine*, 20(2), 104.

Liu Y, et al. (2022) Biogenesis, functions, and clinical implications of circular RNAs in non-small cell lung cancer. *Molecular therapy. Nucleic acids*, 27, 50.

Xie X, et al. (2021) LncRNA CDKN2B-AS1 stabilized by IGF2BP3 drives the malignancy of renal clear cell carcinoma through epigenetically activating NUF2 transcription. *Cell death & disease*, 12(2), 201.

Tang X, et al. (2021) Review on circular RNAs and new insights into their roles in cancer. *Computational and structural biotechnology journal*, 19, 910.

Cheng D, et al. (2021) Cancer-related circular RNA: diverse biological functions. *Cancer cell international*, 21(1), 11.

Zaiou M, et al. (2020) circRNAs Signature as Potential Diagnostic and Prognostic Biomarker for Diabetes Mellitus and Related Cardiovascular Complications. *Cells*, 9(3).

Gao L, et al. (2020) Identification of long noncoding RNA RP11-89K21.1 and RP11-357H14.17 as prognostic signature of endometrial carcinoma via integrated bioinformatics analysis. *Cancer cell international*, 20, 268.

Yao L, et al. (2020) Decreased expression of the long non-coding RNA HOXD-AS2 promotes gastric cancer progression by targeting HOXD8 and activating PI3K/Akt signaling pathway. *World journal of gastrointestinal oncology*, 12(11), 1237.

Xu J, et al. (2020) LncRNA BBOX1-AS1 upregulates HOXC6 expression through miR-361-3p and HuR to drive cervical cancer progression. *Cell proliferation*, 53(7), e12823.

Zhang G, et al. (2020) LINC01419 facilitates hepatocellular carcinoma growth and metastasis through targeting EZH2-regulated RECK. *Aging*, 12(11), 11071.

Huan L, et al. (2020) Hypoxia induced LUCAT1/PTBP1 axis modulates cancer cell viability and chemotherapy response. *Molecular cancer*, 19(1), 11.

Liang X, et al. (2020) FOXD2-AS1 Predicts Dismal Prognosis for Oral Squamous Cell Carcinoma and Regulates Cell Proliferation. *Cell transplantation*, 29, 963689720964411.

Xu L, et al. (2020) LncRNA SNHG11 facilitates tumor metastasis by interacting with and stabilizing HIF-1 α . *Oncogene*, 39(46), 7005.

Ye P, et al. (2020) H3K27ac-activated LINC00519 promotes lung squamous cell carcinoma progression by targeting miR-450b-5p/miR-515-5p/YAP1 axis. *Cell proliferation*, 53(5), e12797.

Zhang Y, et al. (2020) Identification of Potential Prognostic Long Non-Coding RNA Biomarkers for Predicting Recurrence in Patients with Cervical Cancer. *Cancer management and research*, 12, 719.

Wang Y, et al. (2020) High Cancer Susceptibility Candidate 8 Expression Is Associated With Poor Prognosis of Pancreatic Adenocarcinoma: Validated Analysis Based on Four Cancer Databases. *Frontiers in cell and developmental biology*, 8, 392.

Lin YZ, et al. (2020) Bioinformatics Analysis of the Expression of Key Long Intergenic Non-Protein Coding RNA Genes in Bladder Cancer. *Medical science monitor : international medical journal of experimental and clinical research*, 26, e920504.