## **Resource Summary Report**

Generated by NIF on May 17, 2025

# **MCODE**

RRID:SCR 015828

Type: Tool

### **Proper Citation**

MCODE (RRID:SCR\_015828)

#### Resource Information

**URL:** http://apps.cytoscape.org/apps/mcode

**Proper Citation:** MCODE (RRID:SCR\_015828)

**Description:** Software that clusters a given network based on topology to find densely connected regions. It can be used to find protein complexes and parts of pathways in proteinprotein interaction networks or protein families in protein similarity networks.

**Resource Type:** software application, data analysis software, data processing software, algorithm resource, image analysis software, software resource

Defining Citation: PMID:12525261, PMID:25570063

**Keywords:** cluster, clustering, graph analysis, network, network topology, interconnectivity

Funding: Canadian Institutes of Health Research (CIHR); Ontario Research and Development Challenge Fund;

MDS-Sciex

Availability: Free, Available for download

Resource Name: MCODE

Resource ID: SCR\_015828

Alternate URLs: http://baderlab.org/Software/MCODE

**Record Creation Time:** 20220129T080327+0000

Record Last Update: 20250517T060227+0000

### **Ratings and Alerts**

No rating or validation information has been found for MCODE.

No alerts have been found for MCODE.

### **Data and Source Information**

Source: SciCrunch Registry

### **Usage and Citation Metrics**

We found 1036 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.

Ren F, et al. (2025) A small-molecule TNIK inhibitor targets fibrosis in preclinical and clinical models. Nature biotechnology, 43(1), 63.

Yuan HJ, et al. (2025) Exploration and verification of the therapeutic mechanism of shenfu injection in sepsis-induced myocardial injury. PloS one, 20(1), e0317738.

Ren J, et al. (2025) Active Ingredients and Potential Mechanism of Additive Sishen Decoction in Treating Rheumatoid Arthritis with Network Pharmacology and Molecular Dynamics Simulation and Experimental Verification. Drug design, development and therapy, 19, 405.

Zhuang Y, et al. (2025) Core microbe Bifidobacterium in the hindgut of calves improves the growth phenotype of young hosts by regulating microbial functions and host metabolism. Microbiome, 13(1), 13.

Yang J, et al. (2025) Perilipin-2 mediates ferroptosis in oligodendrocyte progenitor cells and myelin injury after ischemic stroke. Neural regeneration research, 20(7), 2015.

Li XY, et al. (2025) Tryptophan metabolism-related gene CYP1B1 serves as a shared biomarker for both Parkinson's disease and insomnia. Scientific reports, 15(1), 1362.

Zhang L, et al. (2025) Comprehensive analysis pinpoints CCNA2 as a prognostic and immunological biomarker in non-small cell lung cancer. BMC pulmonary medicine, 25(1), 14.

Zhao N, et al. (2025) Identification of critical endoplasmic reticulum stress-related genes in advanced atherosclerotic plaque. Scientific reports, 15(1), 2107.

George S, et al. (2025) Feasibility of structuring electronic health record data to facilitate real-world data research: ICAREdata methods applied to multicenter cancer clinical trials. Cancer, 131(1), e35528.

Li J, et al. (2025) Knockdown of CENPF induces cell cycle arrest and inhibits epithelial?mesenchymal transition progression in glioma. Oncology letters, 29(1), 61.

Zhu HR, et al. (2025) Double-negative T cells with a distinct transcriptomic profile are abundant in the peripheral blood of patients with breast cancer. Breast cancer research and treatment, 209(1), 103.

Ma Y, et al. (2025) Exploring the comorbidity mechanisms between atherosclerosis and hashimoto's thyroiditis based on microarray and single-cell sequencing analysis. Scientific reports, 15(1), 1792.

Linares-Pineda TM, et al. (2025) Longitudinal DNA methylation profiles in saliva of offspring from mothers with gestational diabetes: associations with early childhood growth patterns. Cardiovascular diabetology, 24(1), 15.

Zhang X, et al. (2025) MARCHF8-mediated ubiquitination via TGFBI regulates NF-?B dependent inflammatory responses and ECM degradation in intervertebral disc degeneration. PloS one, 20(1), e0314021.

Long H, et al. (2025) Proteomic Characterization of Liver Cancer Cells Treated with Clinical Targeted Drugs for Hepatocellular Carcinoma. Biomedicines, 13(1).

Hong Y, et al. (2024) Identification of molecular subtypes and diagnostic model in clear cell renal cell carcinoma based on collagen-related genes may predict the response of immunotherapy. Frontiers in pharmacology, 15, 1325447.

Zhao M, et al. (2024) Il1r2 and Tnfrsf12a in transcranial magnetic stimulation effect of ischemic stroke via bioinformatics analysis. Medicine, 103(4), e36109.

Xu D, et al. (2024) Integrative bioinformatics analysis of miRNA and mRNA expression profiles and identification of associated miRNA-mRNA network in intracranial aneurysms. Non-coding RNA research, 9(2), 471.

Ge X, et al. (2024) Roles of pyroptosis and immune infiltration in aortic dissection. Frontiers in molecular biosciences, 11, 1277818.

Chen L, et al. (2024) Identification and validation of mutual hub genes in idiopathic pulmonary fibrosis and rheumatoid arthritis-associated usual interstitial pneumonia. Heliyon, 10(7), e28088.