

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

netClass

RRID:SCR_005672

Type: Tool

Proper Citation

netClass (RRID:SCR_005672)

Resource Information

URL: <http://sourceforge.net/projects/netclassr/>

Proper Citation: netClass (RRID:SCR_005672)

Description: An R package for network-based feature (gene) selection for biomarkers discovery via integrating biological information. The package adapts the following 5 algorithms for classifying and predicting gene expression data using prior knowledge: # average gene expression of pathway (aep); # pathway activities classification (PAC); # Hub network classification (hubc); # filter via top ranked genes (FrSVM); # network smoothed t-statistic (stSVM).

Abbreviations: netClass

Resource Type: software resource

Defining Citation: [PMID:24443376](#)

Keywords: protein-protein interaction network, biomarker discovery, classification, microarray, gene expression profile, protein-protein interaction, gene expression, gene, biomarker

Availability: Free, Public

Resource Name: netClass

Resource ID: SCR_005672

Alternate IDs: OMICS_02241

Record Creation Time: 20220129T080231+0000

Record Last Update: 20260725T190608+0000

Ratings and Alerts

No rating or validation information has been found for netClass.

No alerts have been found for netClass.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.