Resource Summary Report

Generated by dkNET on Apr 29, 2025

MBCluster.Seq

RRID:SCR_005079

Type: Tool

Proper Citation

MBCluster.Seq (RRID:SCR_005079)

Resource Information

URL: http://cran.r-project.org/web/packages/MBCluster.Seq/index.html

Proper Citation: MBCluster.Seq (RRID:SCR_005079)

Description: Software to cluster genes based on Poisson or Negative-Binomial model for

RNA-Seq or other digital gene expression (DGE) data.

Abbreviations: MBCluster.Seq

Synonyms: MBCluster.Seq: Model-Based Clustering for RNA-seq Data

Resource Type: software resource

Defining Citation: PMID:24191069

Keywords: bio.tools

Funding:

Availability: GNU General Public License, >/=v3

Resource Name: MBCluster.Seq

Resource ID: SCR_005079

Alternate IDs: OMICS 01417, biotools:mbcluster.seq

Alternate URLs: https://bio.tools/mbcluster.seq

Record Creation Time: 20220129T080228+0000

Record Last Update: 20250420T014243+0000

Ratings and Alerts

No rating or validation information has been found for MBCluster.Seq.

No alerts have been found for MBCluster.Seq.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Thorpe P, et al. (2014) Genomic characterisation of the effector complement of the potato cyst nematode Globodera pallida. BMC genomics, 15(1), 923.