

Resource Summary Report

Generated by RRID on Aug 2, 2026

dynamicTreeCut

RRID:SCR_024280

Type: Tool

Proper Citation

dynamicTreeCut (RRID:SCR_024280)

Resource Information

URL: <https://cran.r-project.org/web/packages/dynamicTreeCut/index.html>

Proper Citation: dynamicTreeCut (RRID:SCR_024280)

Description: Software R package contains methods for detection of clusters in hierarchical clustering dendrograms.

Synonyms: r-cran-dynamictreecut, dynamicTreeCut: Methods for Detection of Clusters in Hierarchical Clustering Dendrograms

Resource Type: software resource, software toolkit

Keywords: clusters detection, hierarchical clustering dendrograms,

Availability: Free, Available for download, Freely available,

Resource Name: dynamicTreeCut

Resource ID: SCR_024280

Alternate IDs: OMICS_23803

Alternate URLs: <https://sources.debian.org/src/r-cran-dynamictreecut/>

License: GPL-2 | GPL-3 [expanded from: GPL (? 2)]

Record Creation Time: 20230830T050217+0000

Record Last Update: 20260801T190856+0000

Ratings and Alerts

No rating or validation information has been found for dynamicTreeCut.

No alerts have been found for dynamicTreeCut.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Tlemsani C, et al. (2024) Sarcoma_CellminerCDB: A tool to interrogate the genomic and functional characteristics of a comprehensive collection of sarcoma cell lines. iScience, 27(6), 109781.

Li C, et al. (2024) scRank infers drug-responsive cell types from untreated scRNA-seq data using a target-perturbed gene regulatory network. Cell reports. Medicine, 5(6), 101568.

Nouri N, et al. (2024) GENIX enables comparative network analysis of single-cell RNA sequencing to reveal signatures of therapeutic interventions. Cell reports methods, 4(6), 100794.

Finlay CM, et al. (2023) T helper 2 cells control monocyte to tissue-resident macrophage differentiation during nematode infection of the pleural cavity. Immunity, 56(5), 1064.

Funnell T, et al. (2019) Integrated structural variation and point mutation signatures in cancer genomes using correlated topic models. PLoS computational biology, 15(2), e1006799.