

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 18, 2026

GramCluster

RRID:SCR_000272

Type: Tool

Proper Citation

GramCluster (RRID:SCR_000272)

Resource Information

URL: <http://bioinfo.unl.edu/gramcluster.php>

Proper Citation: GramCluster (RRID:SCR_000272)

Description: Software implementing a fast and accurate progressive clustering algorithm that relies on a grammar-based sequence distance and is particularly useful in clustering large datasets.

Abbreviations: GramCluster

Resource Type: software resource

Defining Citation: [PMID:21167044](#)

Keywords: 16s sequence, cluster

Availability: Free, Available for download, Freely available

Resource Name: GramCluster

Resource ID: SCR_000272

Alternate IDs: OMICS_01956

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260912T195506+0000

Ratings and Alerts

No rating or validation information has been found for GramCluster.

No alerts have been found for GramCluster.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.