# **Resource Summary Report**

Generated by NIF on May 13, 2025

# **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

**Funding:** 

Resource Name: GramCluster

Resource ID: SCR\_000272

Alternate IDs: OMICS\_01956

**Record Creation Time:** 20220129T080200+0000

**Record Last Update:** 20250420T013938+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.