

# Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 2026

## GramCluster

RRID:SCR\_000272

Type: Tool

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### Proper Citation

GramCluster (RRID:SCR\_000272)

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### 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:** 20260808T185714+0000

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### Ratings and Alerts

No rating or validation information has been found for GramCluster.

No alerts have been found for GramCluster.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.