

Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

[Plantagora](#)

RRID:SCR_001227

Type: Tool

Proper Citation

Plantagora (RRID:SCR_001227)

Resource Information

URL: <http://www.plantagora.org/>

Proper Citation: Plantagora (RRID:SCR_001227)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on August 18,2025. A web-based plant genome assembly simulation platform whose resources include out of the box scripts for analyzing assembly data, an on-demand web graphing tool to model your experiment, and a downloadable database with metrics and parameters from over 3,000 simulated genome assemblies.

Abbreviations: Plantagora

Synonyms: Plantagora - a resource for plant genome assembly

Resource Type: data or information resource, data set, software resource

Defining Citation: [PMID:22174807](#)

Keywords: next-generation sequencing, genome assembly, sequencing, genome, 454, illumina, simulate

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Plantagora

Resource ID: SCR_001227

Alternate IDs: OMICS_02116

Record Creation Time: 20220129T080206+0000

Record Last Update: 20260905T133036+0000

Ratings and Alerts

No rating or validation information has been found for Plantagora.

No alerts have been found for Plantagora.

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 [RRID](#) .

Ramazzotti M, et al. (2015) riboFrame: An Improved Method for Microbial Taxonomy Profiling from Non-Targeted Metagenomics. Frontiers in genetics, 6, 329.