Resource Summary Report

Generated by kravitz2 on Aug 1, 2026

AmpliconTagger

RRID:SCR_019112
Type: Tool

Proper Citation

AmpliconTagger (RRID:SCR_019112)

Resource Information

URL: https://jtremblay.github.io/amplicontagger.html

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Description: Software tool as rRNA marker gene amplicon pipeline coded in python framework that enables fine tuning and integration of virtually any potential rRNA gene amplicon bioinformatic procedure. Designed to work within HPC environment, supporting complex network of job dependencies with smart restart mechanism in case of job failure or parameter modifications.

Resource Type: software resource, software application, workflow software, data processing software

Defining Citation: PMID:31816087

Keywords: High Performance Computing, HPC environment, rRNA gene amplicons, rRNA marker, gene amplicon pipeline, bio.tools

Availability: Free, Freely available

Resource Name: AmpliconTagger

Resource ID: SCR_019112

Alternate IDs: SCR_019113, biotools:amplicontagger

Alternate URLs: https://bitbucket.org/jtremblay514/nrc_pipeline_public/src/master/, https://jtremblay.github.io/amplicontagger_guide.html, https://bio.tolols/amplicontagger

License: GNU GPL

Record Creation Time: 20220129T080343+0000

Record Last Update: 20260801T042824+0000

Ratings and Alerts

No rating or validation information has been found for AmpliconTagger.

No alerts have been found for AmpliconTagger.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at kravitz2 .

Ducret J, et al. (2025) Role of biofilm during groundwater biofiltration of manganese. Scientific reports, 15(1), 41330.

Middleton H, et al. (2024) Rhizospheric miRNAs affect the plant microbiota. ISME communications, 4(1), ycae120.

Schreiber L, et al. (2021) In situ microcosms deployed at the coast of British Columbia (Canada) to study dilbit weathering and associated microbial communities under marine conditions. FEMS microbiology ecology, 97(7).