

Resource Summary Report

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NanoAmpli-Seq

RRID:SCR_016710

Type: Tool

Proper Citation

NanoAmpli-Seq (RRID:SCR_016710)

Resource Information

URL: <https://github.com/umerijaz/nanopore>

Proper Citation: NanoAmpli-Seq (RRID:SCR_016710)

Description: Software for a workflow for amplicon sequencing from mixed microbial communities on the nanopore sequencing platform. Used for full-length SSU rRNA gene sequencing.

Resource Type: data analysis software, data processing software, sequence analysis software, software application, software resource

Defining Citation: [DOI:10.1101/244517](https://doi.org/10.1101/244517)

Keywords: amplicon, sequencing, 16S rRNA, mixed, microbial, communities, nanopore, platform, gene

Funding: EPSRC EP/M016811/1;
NERC NE/L011956/1

Availability: Free, Available for download, Freely available

Resource Name: NanoAmpli-Seq

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Alternate URLs: <https://www.protocols.io/view/nanoampli-seq-bioinformatics-workflow-u25eyg6>

Record Creation Time: 20220129T080332+0000

Record Last Update: 20260905T132813+0000

Ratings and Alerts

No rating or validation information has been found for NanoAmpli-Seq.

No alerts have been found for NanoAmpli-Seq.

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 [Kravitz](#) .

Calus ST, et al. (2018) NanoAmpli-Seq: a workflow for amplicon sequencing for mixed microbial communities on the nanopore sequencing platform. GigaScience, 7(12).