Grinder
RRID:SCR_000168
Type: Tool

Proper Citation

Grinder (RRID:SCR_000168)

Resource Information

URL: http://sourceforge.net/projects/biogrinder/

Proper Citation: Grinder (RRID:SCR_000168)

Description: An open-source bioinformatic tool to create simulated omic shotgun and amplicon sequence libraries for all main sequencing platforms. The tool is available through multiple interfaces like GUI, CLI and API. It is useful for simulating clinical or environmental microbial communities and complements the use of in vitro mock communities.

Abbreviations: Grinder

Resource Type: software resource

Defining Citation: PMID:22434876, DOI:10.1093/nar/gks251

Keywords: simulation, amplicon, shotgun, genomic sequencing, clinical, metagenomic, transcriptomic and metatranscriptomic

Availability: Public, Open Source

Resource Name: Grinder

Resource ID: SCR_000168

Alternate IDs: OMICS_01508

 Alternate URLs: https://sources.debian.org/src/grinder/

Record Creation Time: 20220129T080200+0000
Ratings and Alerts

No rating or validation information has been found for Grinder.

No alerts have been found for Grinder.

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 FDI Lab - SciCrunch.org.

