Grinder
RRID:SCR_000168
Type: Tool

Proper Citation
Grinder (RRID:SCR_000168)

Resource Information

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

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.

Resource Name: Grinder

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Resource Type: Resource, software resource

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

Resource ID: SCR_000168

Parent Organization: SourceForge

References: PMID:22434876

Availability: Public, Open Source

Website Status: Last checked up

Alternate IDs: OMICS_01508

Abbreviations: Grinder
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 Infrastructure.

