SortMeRNA
RRID:SCR_014402
Type: Tool

Proper Citation
SortMeRNA (RRID:SCR_014402)

Resource Information

URL: http://bioinfo.lifl.fr/RNA/sortmerna/

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Description: Sequence analysis software for filtering, mapping and OTU-picking NGS reads. SortMeRNA takes as input a file of reads (fasta or fastq format) and one or multiple rRNA database file(s), and sorts apart rRNA and rejected reads into two files specified by the user.

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

References: PMID:23071270, DOI:10.1093/bioinformatics/bts611

Keywords: sequence analysis software, filter, map, otu picking, ngs, sort, rna, rRNA, bio.tools

Availability: Freely available

Website Status: Last checked up

Resource Name: SortMeRNA

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Alternate IDs: biotools:sortmerna

Alternate URLs: https://bio.tools/sortmerna

Ratings and Alerts
No rating or validation information has been found for SortMeRNA.

No alerts have been found for SortMeRNA.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 225 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch.org.


Sillen WMA, et al. (2020) Nanoparticle treatment of maize analyzed through the metatranscriptome: compromised nitrogen cycling, possible phytopathogen selection, and plant hormesis. Microbiome, 8(1), 127.