SortMeRNA
RRID:SCR_014402
Type: Tool

Proper Citation
SortMeRNA (RRID:SCR_014402)

Resource Information

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

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.

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

Keywords: sequence analysis software, filter, map, otu picking, ngs, sort, rna, rrna

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

Availability: Freely available

Website Status: Last checked up

Mentions Count: 153

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

No alerts have been found for SortMeRNA.

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Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 153 mentions in open access literature.

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


Li F, et al. (2019) Comparative metagenomic and metatranscriptomic analyses reveal the
breed effect on the rumen microbiome and its associations with feed efficiency in beef cattle. Microbiome, 7(1), 6.


