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

Resource ID: SCR_014402

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.

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**Data and Source Information**

**Source:** [SciCrunch Registry](https://SciCrunch.org)

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**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](https://FDI Lab - SciCrunch.org).


Pascual A, et al. (2020) Effect of dietary supplementation with yeast cell wall extracts on


