

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

riboPicker

RRID:SCR_000360

Type: Tool

Proper Citation

riboPicker (RRID:SCR_000360)

Resource Information

URL: <http://ribopicker.sourceforge.net/>

Proper Citation: riboPicker (RRID:SCR_000360)

Description: Software to automatically identify and efficiently remove rRNA-like sequences from metatranscriptomic and metagenomic datasets.

Resource Type: software resource

Defining Citation: [PMID:22155869](#)

Keywords: standalone software, perl, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: riboPicker

Resource ID: SCR_000360

Alternate IDs: OMICS_02618, biotools:ribopicker

Alternate URLs: <https://bio.tools/ribopicker>

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260905T132417+0000

Ratings and Alerts

No rating or validation information has been found for riboPicker.

No alerts have been found for riboPicker.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [Kravitz](#) .

Kulkarni KS, et al. (2016) De novo Transcriptome Sequencing to Dissect Candidate Genes Associated with Pearl Millet-Downy Mildew (*Sclerospora graminicola* Sacc.) Interaction. *Frontiers in plant science*, 7, 847.

Wang K, et al. (2016) Clam focal and systemic immune responses to QPX infection revealed by RNA-seq technology. *BMC genomics*, 17, 146.