

# Resource Summary Report

Generated by [RRID](#) 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](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

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.