

# Resource Summary Report

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

## PPR-Meta

RRID:SCR\_016915

Type: Tool

### Proper Citation

PPR-Meta (RRID:SCR\_016915)

### Resource Information

**URL:** [http://cqb.pku.edu.cn/ZhuLab/PPR\\_Meta](http://cqb.pku.edu.cn/ZhuLab/PPR_Meta)

**Proper Citation:** PPR-Meta (RRID:SCR\_016915)

**Description:** Software tool to identify metagenomic sequences of phages, chromosomes or plasmids. Used for identifying phages and plasmids from metagenomic fragments using deep learning., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

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

**Keywords:** identify, metagenomic, sequence, phage, chromosome, plasmid, fragment, machine, learning

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** PPR-Meta

**Resource ID:** SCR\_016915

**Alternate URLs:** <https://github.com/zhenchengfang/PPR-Meta>

**License:** GNU General Public License v3.0

**Record Creation Time:** 20220129T080332+0000

**Record Last Update:** 20260905T132815+0000

### Ratings and Alerts

No rating or validation information has been found for PPR-Meta.

No alerts have been found for PPR-Meta.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Feng T, et al. (2024) MOBFinder: a tool for mobilization typing of plasmid metagenomic fragments based on a language model. GigaScience, 13.

Marquet M, et al. (2022) What the Phage: a scalable workflow for the identification and analysis of phage sequences. GigaScience, 11.

Pradier L, et al. (2021) PlasForest: a homology-based random forest classifier for plasmid detection in genomic datasets. BMC bioinformatics, 22(1), 349.

Fang Z, et al. (2019) PPR-Meta: a tool for identifying phages and plasmids from metagenomic fragments using deep learning. GigaScience, 8(6).