

Resource Summary Report

Generated by [Kravitz](#) 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

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 [Kravitz](#) .

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).