

# Resource Summary Report

Generated by [kravitz2](#) on Jul 30, 2026

## MetaCluster-TA

RRID:SCR\_004599

Type: Tool

### Proper Citation

MetaCluster-TA (RRID:SCR\_004599)

### Resource Information

**URL:** <http://i.cs.hku.hk/~alse/MetaCluster/>

**Proper Citation:** MetaCluster-TA (RRID:SCR\_004599)

**Description:** A software for binning and annotating short paired-end reads.

**Abbreviations:** MetaCluster-TA

**Resource Type:** software resource

**Defining Citation:** [PMID:24564377](#)

**Keywords:** binning, annotation, metagenomics

**Resource Name:** MetaCluster-TA

**Resource ID:** SCR\_004599

**Alternate IDs:** OMICS\_01473

**Record Creation Time:** 20220129T080225+0000

**Record Last Update:** 20260725T190555+0000

### Ratings and Alerts

No rating or validation information has been found for MetaCluster-TA.

No alerts have been found for MetaCluster-TA.

### 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 [kravitz2](#) .

Malla MA, et al. (2018) Exploring the Human Microbiome: The Potential Future Role of Next-Generation Sequencing in Disease Diagnosis and Treatment. *Frontiers in immunology*, 9, 2868.

Alvarenga DO, et al. (2017) A Metagenomic Approach to Cyanobacterial Genomics. *Frontiers in microbiology*, 8, 809.

Escobar-Zepeda A, et al. (2015) The Road to Metagenomics: From Microbiology to DNA Sequencing Technologies and Bioinformatics. *Frontiers in genetics*, 6, 348.

Kumar S, et al. (2015) Metagenomics: Retrospect and Prospects in High Throughput Age. *Biotechnology research international*, 2015, 121735.