Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 8, 2025

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

Funding:

Resource Name: MetaCluster-TA

Resource ID: SCR_004599

Alternate IDs: OMICS_01473

Record Creation Time: 20220129T080225+0000

Record Last Update: 20250214T183019+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 FDI Lab - SciCrunch.org.

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.

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

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