Resource Summary Report

Generated by Kravitz on Sep 10, 2026

MAP

RRID:SCR_013216
Type: Tool

Proper Citation

MAP (RRID:SCR_013216)

Resource Information

URL: http://bioinfo.ctb.pku.edu.cn/MAP/

Proper Citation: MAP (RRID:SCR_013216)

Description: This resource is out of service. Documented on February 23,2021. Software for de novo metagenomic assembly program for shotgun DNA reads., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Synonyms: Metagenomic Assembly Program

Resource Type: software resource

Defining Citation: PMID:22495746

Keywords: bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: MAP

Resource ID: SCR_013216

Alternate IDs: OMICS_01424, biotools:MAP

Alternate URLs: https://bio.tools/MAP

License: GNU General Public License

Record Creation Time: 20220129T080314+0000

Record Last Update: 20260905T132734+0000

Ratings and Alerts

No rating or validation information has been found for MAP.

No alerts have been found for MAP.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at Kravitz .

Kayani MUR, et al. (2021) Genome-resolved metagenomics using environmental and clinical samples. Briefings in bioinformatics, 22(5).