

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

## MHAP

RRID:SCR\_024113

Type: Tool

### Proper Citation

MHAP (RRID:SCR\_024113)

### Resource Information

**URL:** <https://mhap.readthedocs.io/en/stable/>

**Proper Citation:** MHAP (RRID:SCR\_024113)

**Description:** Software tool as reference implementation of probabilistic sequence overlapping algorithm. Used to detect overlaps between noisy long-read sequence data.

**Synonyms:** MinHash Alignment Process, mhap

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

**Defining Citation:** [PMID:26006009](#)

**Keywords:** detect overlaps between noisy long-read sequence data, probabilistic sequence overlapping algorithm,

**Availability:** Free, Available for download, Freely available,

**Resource Name:** MHAP

**Resource ID:** SCR\_024113

**Alternate IDs:** OMICS\_13515

**Alternate URLs:** <https://sources.debian.org/src/mhap/>, <https://github.com/marbl/MHAP>

**License:** Apache-2.0 license

**Record Creation Time:** 20230824T050212+0000

**Record Last Update:** 20260905T133029+0000

### Ratings and Alerts

No rating or validation information has been found for MHAP.

No alerts have been found for MHAP.

---

## Data and Source Information

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

---

## Usage and Citation Metrics

We have not found any literature mentions for this resource.