

# Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

## PoolHap

RRID:SCR\_012129

Type: Tool

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### Proper Citation

PoolHap (RRID:SCR\_012129)

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### Resource Information

**URL:** <https://github.com/Gregor-Mendel-Institute/poolhap>

**Proper Citation:** PoolHap (RRID:SCR\_012129)

**Description:** Software tool for inferring haplotypes from pooled sequencing. Enables to infer strain numbers and haplotype frequencies in silico from sequences of pooled samples.

**Synonyms:** Inferring Haplotype frequencies from Pooled sequencing, poolhap2, PoolHap2

**Resource Type:** software application, software resource, standalone software

**Defining Citation:** [PMID:21264334](#)

**Keywords:** inferring haplotypes, pooled sequencing, haplotype frequencies, infer strain numbers, pooled samples sequences, bio.tools

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

**Resource Name:** PoolHap

**Resource ID:** SCR\_012129

**Alternate IDs:** biotools:poolhap, OMICS\_05832

**Alternate URLs:** <https://bio.tools/poolhap>

**License:** MIT License

**Record Creation Time:** 20220129T080308+0000

**Record Last Update:** 20260905T133254+0000

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### Ratings and Alerts

No rating or validation information has been found for PoolHap.

No alerts have been found for PoolHap.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.