

Resource Summary Report

Generated by T1D on Sep 5, 2026

PoolHap

RRID:SCR_012129

Type: Tool

Proper Citation

PoolHap (RRID:SCR_012129)

Resource Information

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

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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](https://pubmed.ncbi.nlm.nih.gov/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: 20260904T000449+0000

Ratings and Alerts

No rating or validation information has been found for PoolHap.

No alerts have been found for PoolHap.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.