Resource Summary Report

Generated by NIF on Apr 30, 2025

PoolHap

RRID:SCR_012129

Type: Tool

Proper Citation

PoolHap (RRID:SCR_012129)

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: standalone software, software resource, software application

Defining Citation: PMID:21264334

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

Funding:

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: 20250430T055801+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.