Resource Summary Report

Generated by NIF on Apr 27, 2025

Libra

RRID:SCR 016608

Type: Tool

Proper Citation

Libra (RRID:SCR_016608)

Resource Information

URL: https://github.com/iychoi/libra

Proper Citation: Libra (RRID:SCR_016608)

Description: Hadoop based tool for massive comparative metagenomics analysis. Compute the similarity between metagenomic samples.

Resource Type: data processing software, data analysis software, sequence analysis software, software resource, data analytics software, software application

Keywords: gene, distance, matrix, computation, k-mer-based, sequence, comparison, Hadoop, metagenomic, sample, bio.tools

Funding: NSF 1640775

Availability: Free, Available for download, Freely available

Resource Name: Libra

Resource ID: SCR 016608

Alternate IDs: biotools:Libra_k-mer

Alternate URLs: https://bio.tools/Libra_k-mer

License: Apache License

Record Creation Time: 20220129T080331+0000

Record Last Update: 20250426T060557+0000

Ratings and Alerts

No rating or validation information has been found for Libra.

No alerts have been found for Libra.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.