Resource Summary Report

Generated by RRID on Apr 28, 2025

<u>Libra</u>

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, sequence analysis software, software resource, data analysis software, software application, data analytics software

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: 20250428T054009+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: <u>SciCrunch Registry</u>

Usage and Citation Metrics

We have not found any literature mentions for this resource.