# **Resource Summary Report**

Generated by NIF on May 15, 2025

# Ray

RRID:SCR 001916

Type: Tool

### **Proper Citation**

Ray (RRID:SCR\_001916)

#### Resource Information

URL: http://sourceforge.net/projects/denovoassembler/files/

**Proper Citation:** Ray (RRID:SCR\_001916)

**Description:** Software that assembles reads obtained with new sequencing technologies

(Illumina, 454, SOLiD) using MPI 2.2.

**Abbreviations:** Ray

Synonyms: Ray - a de novo assembler using MPI 2.2, Ray - Parallel genome assemblies

for parallel DNA sequencing

**Resource Type:** software resource

**Defining Citation:** PMID:20958248, DOI:10.1089/cmb.2009.0238

**Keywords:** mpi, bio.tools

**Funding:** 

Resource Name: Ray

Resource ID: SCR\_001916

Alternate IDs: OMICS\_00027, biotools:ray

Alternate URLs: https://bio.tools/ray, https://sources.debian.org/src/ray/

**Record Creation Time:** 20220129T080210+0000

Record Last Update: 20250420T014045+0000

## **Ratings and Alerts**

No rating or validation information has been found for Ray.

No alerts have been found for Ray.

### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.

Arikan M, et al. (2024) gNOMO2: a comprehensive and modular pipeline for integrated multiomics analyses of microbiomes. GigaScience, 13.