

# Resource Summary Report

Generated by [NIF](#) on Aug 10, 2026

## CongrPE

RRID:SCR\_013190

Type: Tool

### Proper Citation

CongrPE (RRID:SCR\_013190)

### Resource Information

**URL:** <http://sourceforge.net/projects/congrpe/>

**Proper Citation:** CongrPE (RRID:SCR\_013190)

**Description:** A de novo assembly algorithm for Next-Generation Sequencing technology.

**Abbreviations:** CongrPE

**Resource Type:** software resource

**Resource Name:** CongrPE

**Resource ID:** SCR\_013190

**Alternate IDs:** OMICS\_00011

**Record Creation Time:** 20220129T080314+0000

**Record Last Update:** 20260808T190020+0000

### Ratings and Alerts

No rating or validation information has been found for CongrPE.

No alerts have been found for CongrPE.

### 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](#) .

Thangam M, et al. (2015) CRCDA--Comprehensive resources for cancer NGS data analysis. Database : the journal of biological databases and curation, 2015.