## **Resource Summary Report**

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# **IVA**

RRID:SCR\_024044 Type: Tool

**Proper Citation** 

IVA (RRID:SCR\_024044)

#### **Resource Information**

URL: https://github.com/sanger-pathogens/iva

Proper Citation: IVA (RRID:SCR\_024044)

**Description:** Software tool as de novo assembler designed to assemble virus genomes that have no repeat sequences, using Illumina read pairs sequenced from mixed populations at extremely high and variable depth.

**Synonyms:** Iterative Virus Assembler - de novo virus assembler of Illumina paired reads., iva

**Resource Type:** data analysis software, software application, software resource, data processing software

Defining Citation: PMID:25725497

**Keywords:** de novo assembler, assemble virus genomes, no repeat sequences, Illumina reads, pairs sequenced from mixed populations,

#### Funding:

Availability: Free, Available for download, Freely available,

**Resource Name:** IVA

Resource ID: SCR\_024044

Alternate IDs: OMICS\_08188

Alternate URLs: https://sources.debian.org/src/iva/, http://sanger-pathogens.github.io/iva/

License: GNU GENERAL PUBLIC LICENSE

**Record Creation Time:** 20230824T050211+0000

Record Last Update: 20250426T060956+0000

### **Ratings and Alerts**

No rating or validation information has been found for IVA.

No alerts have been found for IVA.

Data and Source Information

Source: <u>SciCrunch Registry</u>

**Usage and Citation Metrics** 

We have not found any literature mentions for this resource.