Trinity
RRID:SCR_013048
Type: Tool

Proper Citation

Trinity (RRID:SCR_013048)

Resource Information

URL: http://trinityrnaseq.sourceforge.net/

Proper Citation: Trinity (RRID:SCR_013048)

Description: Software for the efficient and robust de novo reconstruction of transcriptomes from RNA-seq data.

Resource Type: Resource, software resource

Keywords: bio.tools

Parent Organization: Broad Institute, Hebrew University of Jerusalem; Jerusalem; Israel, SourceForge

Website Status: Last checked up

Abbreviations: Trinity

Resource Name: Trinity

Resource ID: SCR_013048

Alternate IDs: OMICS_01327, biotools:trinity

Alternate URLs: https://bio.tools/trinity

Ratings and Alerts

No rating or validation information has been found for Trinity.
No alerts have been found for Trinity.

Data and Source Information

**Source:** SciCrunch Registry

Usage and Citation Metrics

We found 5042 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](https://fdi.lab/scicrunch.org).


Nor Muhammad NA, et al. (2021) Data on RNA-seq analysis of the cocoa pod borer pest(Snellen) (Lepidoptera: Gracillariidae). Data in brief, 34, 106638.


Chen MS, et al. (2020) De novo genome assembly and Hi-C analysis reveal an association between chromatin architecture alterations and sex differentiation in the woody plant Jatropha curcas. GigaScience, 9(2).


Li RT, et al. (2020) A moth odorant receptor highly expressed in the ovipositor is involved in
detecting host-plant volatiles. eLife, 9.


Zhao H, et al. (2020) Mycoparasitism illuminated by genome and transcriptome sequencing of, an important biocontrol fungus of the plant pathogen. Microbial genomics.
