**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.


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.

Warr A, et al. (2020) An improved pig reference genome sequence to enable pig genetics
and genomics research. GigaScience, 9(6).


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