**Trinity**

RRID:SCR_013048  
Type: Tool

**Proper Citation**

Trinity (RRID:SCR_013048)

**Resource Information**

**URL:** [http://trinityrnaseq.sourceforge.net/](http://trinityrnaseq.sourceforge.net/)

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

**Resource Name:** Trinity

**Proper Citation:** Trinity (RRID:SCR_013048)

**Resource Type:** Resource, software resource

**Resource ID:** SCR_013048

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

**Website Status:** Last checked up

**Alternate IDs:** OMICS_01327

**Abbreviations:** Trinity

**Mentions Count:** 4074

**Ratings and Alerts**

No rating or validation information has been found for Trinity.

No alerts have been found for Trinity.
Usage and Citation Metrics

We found 4074 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch Infrastructure](#).


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).

Samsulrizal NH, et al. (2020) transcriptome dataset of accession MS007. Data in brief, 28, 104811.


