Trinity
RRID:SCR_013048
Type: Tool

Proper Citation

Trinity (RRID:SCR_013048)

Resource Information

URL: 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](https://www.fdicluster.org/SciCrunch).


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).


Roy D, et al. (2020) Rhodopsin and melanopsin coexist in mammalian sperm cells and activate different signaling pathways for thermotaxis. Scientific reports, 10(1), 112.


Konopová B, et al. (2020) Transcriptome of pleuropodia from locust embryos supports that these organs produce enzymes enabling the larva to hatch. Frontiers in zoology, 17, 4.
