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: 3737

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 3737 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [scicrunch](https://scicrunch).


Takeda S, et al. (2019) Comparative transcriptome analysis of galls from four different host plants suggests the molecular mechanism of gall development. PloS one, 14(10), e0223686.


