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.
Abbreviations: Trinity
Resource Type: software resource
Defining Citation: DOI:10.1038/nbt.1883
Keywords: bio.tools
Resource Name: Trinity
Resource ID: SCR_013048
Alternate IDs: biotools:trinity, OMICS_01327
Record Creation Time: 20220129T080314+0000
Record Last Update: 20240424T182930+0000

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 7958 mentions in open access literature.

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

Frese AN, et al. (2024) Quantitative proteome dynamics across embryogenesis in a model chordate. iScience, 27(4), 109355.

Niu X, et al. (2024) The genome assembly and annotation of the white-lipped tree pit viper Trimeresurus albolabris. GigaByte (Hong Kong, China), 2024, gigabyte106.


Yang Z, et al. (2024) Two horizontally acquired bacterial genes steer the exceptionally efficient and flexible nitrogenous waste cycling in whiteflies. Science advances, 10(5), eadi3105.


Buller-Peralta I, et al. (2024) Comprehensive allostatic load risk index is associated with increased frontal and left parietal white matter hyperintensities in mid-life cognitively healthy adults. Scientific reports, 14(1), 573.


Wang T, et al. (2024) Transcriptome-Wide Identification of Cytochrome P450s in Tea Black Tussock Moth (Dasychira baibarana) and Candidate Genes Involved in Type-II Sex Pheromone Biosynthesis. Insects, 15(2).