Resource Summary Report

Generated by RRID on Apr 28, 2025

SHORTY

RRID:SCR_002048

Type: Tool

Proper Citation

SHORTY (RRID:SCR_002048)

Resource Information

URL: http://www.cs.sunysb.edu/~skiena/shorty/

Proper Citation: SHORTY (RRID:SCR_002048)

Description: Software for targeted de novo assembly of microreads with mate pair

information and sequencing errors.

Resource Type: data processing software, sequence analysis software, software resource,

data analysis software, software application

Defining Citation: PMID:19208115

Keywords: sequencing, dna, de novo, microreads, assembler, bio.tools

Funding:

Availability: Available for download, Free

Resource Name: SHORTY

Resource ID: SCR_002048

Alternate IDs: biotools:shorty, OMICS_00030

Alternate URLs: https://bio.tools/shorty

Record Creation Time: 20220129T080211+0000

Record Last Update: 20250428T052919+0000

Ratings and Alerts

No rating or validation information has been found for SHORTY.

No alerts have been found for SHORTY.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID.

Kobeni S, et al. (2020) The Dynamic Changes of African Elephant Milk Composition over Lactation. Animals: an open access journal from MDPI, 10(6).

El-Metwally S, et al. (2013) Next-generation sequence assembly: four stages of data processing and computational challenges. PLoS computational biology, 9(12), e1003345.

Hossain MS, et al. (2009) Crystallizing short-read assemblies around seeds. BMC bioinformatics, 10 Suppl 1(Suppl 1), S16.