

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

Scaffold builder

RRID:SCR_000556

Type: Tool

Proper Citation

Scaffold builder (RRID:SCR_000556)

Resource Information

URL: http://edwards.sdsu.edu/scaffold_builder/

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Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 6,2023. Tool designed to generate scaffolds (super contigs of sequences joined by N-bases) using the homology provided by a closely related reference sequence. Scaffold_builder is an advanced wrapper for Nucmer, written in Python that resolves several situations that may arise when mapping contigs to the reference genome.

Abbreviations: scaffold_builder

Resource Type: analysis service resource, data analysis service, production service resource, service resource

Defining Citation: [PMID:24267787](#)

Keywords: scaffolding

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Scaffold builder

Resource ID: SCR_000556

Alternate IDs: OMICS_00046

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260905T132423+0000

Ratings and Alerts

No rating or validation information has been found for Scaffold builder.

No alerts have been found for Scaffold builder.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.