

Resource Summary Report

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

Bambus

RRID:SCR_005068

Type: Tool

Proper Citation

Bambus (RRID:SCR_005068)

Resource Information

URL: <http://sourceforge.net/apps/mediawiki/amos/index.php?title=Bambus2>

Proper Citation: Bambus (RRID:SCR_005068)

Description: Software for scaffolding to address some of the challenges encountered when analyzing metagenomes. Scaffolding represents the task of ordering and orienting contigs by incorporating additional information about their relative placement along the genome. While most other scaffolders are closely tied to a specific assembly program, Bambus accepts the output from most current assemblers and provides the user with great flexibility in choosing the scaffolding parameters. In particular, Bambus is able to accept contig linking data other than specified by mate-pairs. Such sources of information include alignment to a reference genome (Bambus can directly use the output of MUMmer), physical mapping data, or information about gene synteny.

Abbreviations: Bambus

Synonyms: Bambus 2, Bambus 2.0

Resource Type: software resource

Defining Citation: [PMID:21926123](#)

Keywords: scaffolding

Availability: Open unspecified license

Resource Name: Bambus

Resource ID: SCR_005068

Alternate IDs: OMICS_01432

Alternate URLs: <http://sourceforge.net/apps/mediawiki/amos/index.php?title=Bambus>

Record Creation Time: 20220129T080228+0000

Record Last Update: 20260905T132529+0000

Ratings and Alerts

No rating or validation information has been found for Bambus.

No alerts have been found for Bambus.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.