

# Resource Summary Report

Generated by [ASWG](#) on Aug 4, 2026

## SPHINX

RRID:SCR\_000534

Type: Tool

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### Proper Citation

SPHINX (RRID:SCR\_000534)

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### Resource Information

**URL:** <http://metagenomics.atc.tcs.com/SPHINX/>

**Proper Citation:** SPHINX (RRID:SCR\_000534)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 1, 2023. Hybrid binning tool that achieves high binning efficiency by utilizing both "compositional" and "similarity" features of the query sequence during the binning process. SPHINX can analyze sequences in metagenomic data sets as rapidly as composition based approaches, but nevertheless has the accuracy and specificity of similarity based algorithms., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

**Abbreviations:** SPHINX

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

**Defining Citation:** [PMID:21030462](#)

**Keywords:** metagenome, taxonomy, sequence, binning

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** SPHINX

**Resource ID:** SCR\_000534

**Alternate IDs:** OMICS\_01466

**Record Creation Time:** 20220129T080202+0000

**Record Last Update:** 20260804T043104+0000

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### Ratings and Alerts

No rating or validation information has been found for SPHINX.

No alerts have been found for SPHINX.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.