# **Resource Summary Report**

Generated by NIF on Apr 17, 2025

# **SAT-Assembler**

RRID:SCR\_012108

Type: Tool

### **Proper Citation**

SAT-Assembler (RRID:SCR\_012108)

#### **Resource Information**

**URL:** http://sourceforge.net/projects/sat-assembler/

Proper Citation: SAT-Assembler (RRID:SCR\_012108)

Description: A targeted gene assembly software program which aims to recover gene

families of particular interest to biologists.

Resource Type: software resource

**Defining Citation: PMID:25122209** 

**Keywords:** standalone software, python

**Funding:** 

**Availability:** Creative Commons License

Resource Name: SAT-Assembler

Resource ID: SCR\_012108

Alternate IDs: OMICS\_05397

**Record Creation Time:** 20220129T080308+0000

Record Last Update: 20250410T070229+0000

## Ratings and Alerts

No rating or validation information has been found for SAT-Assembler.

No alerts have been found for SAT-Assembler.

### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 2 mentions in open access literature.

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

Guo J, et al. (2019) Review, Evaluation, and Directions for Gene-Targeted Assembly for Ecological Analyses of Metagenomes. Frontiers in genetics, 10, 957.

Zhang Y, et al. (2014) A scalable and accurate targeted gene assembly tool (SAT-Assembler) for next-generation sequencing data. PLoS computational biology, 10(8), e1003737.