SGA
RRID:SCR_001982
Type: Tool

Proper Citation

SGA (RRID:SCR_001982)

Resource Information

URL: https://github.com/jts/sga

Proper Citation: SGA (RRID:SCR_001982)

Description: Software package that functions as a de novo genome assembler based on the concept of string graphs. It is designed as a modular set of programs used to assemble large genomes from high coverage short read data.

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

References: DOI:10.1101/gr.126953.111

Keywords: string, graph assembler, de novo assembly

Availability: Open source

Website Status: Last checked up

Resource Name: SGA

Resource ID: SCR_001982

Alternate IDs: OMICS_00028

Ratings and Alerts

No rating or validation information has been found for SGA.
No alerts have been found for SGA.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 12 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](http://FDI Lab - SciCrunch.org).


