# SSAKE

**RRID:** SCR_010753  
**Type:** Tool

## Proper Citation

SSAKE (RRID:SCR_010753)

## Resource Information

### URL:

http://www.bcgsc.ca/platform/bioinfo/software/ssake

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SSAKE (RRID:SCR_010753)

### Description:

Software designed to help leverage the information from short sequences reads by stringently clustering them into contigs that can be used to characterize novel sequencing targets.

### Abbreviations:

SSAKE

### Resource Type:

software resource

### Defining Citation:

DOI:10.1093/bioinformatics/btl629

### Keywords:

bio.tools

### Resource Name:

SSAKE

### Resource ID:

SCR_010753

### Alternate IDs:

biotools:ssake, OMICS_00033

### Alternate URLs:


### Record Creation Time:

20220129T080300+0000

### Record Last Update:

20240424T182902+0000

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### Ratings and Alerts

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No rating or validation information has been found for SSAKE.

No alerts have been found for SSAKE.

Data and Source Information

**Source:** SciCrunch Registry

Usage and Citation Metrics

We found 11 mentions in open access literature.

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


Li E, et al. (2016) Isolation and characterization of a bacteriophage phiEap-2 infecting multidrug resistant Enterobacter aerogenes. Scientific reports, 6, 28338.


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