Resource Summary Report

Generated by dkNET on May 19, 2025

smashpp

RRID:SCR 018307

Type: Tool

Proper Citation

smashpp (RRID:SCR_018307)

Resource Information

URL: https://github.com/smortezah/smashpp

Proper Citation: smashpp (RRID:SCR_018307)

Description: Software tool to find and visualize rearrangements in DNA sequences.

Synonyms: Smash++

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

Keywords: Find sequence rearrangement, visualize sequence rearrangement, DNA, DNA sequence, DNA sequence rearrangement, bio.tools

Funding:

Availability: Free, Available for download, Freely available

Resource Name: smashpp

Resource ID: SCR_018307

Alternate IDs: biotools:smashpp, BioTools:smashpp

Alternate URLs: https://bio.tools/smashpp, https://bio.tools/smashpp,

https://bio.tools/smashpp

License: GNU General Public License v3.0

Record Creation Time: 20220129T080339+0000

Record Last Update: 20250519T204027+0000

Ratings and Alerts

No rating or validation information has been found for smashpp.

No alerts have been found for smashpp.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>dkNET</u>.

Hosseini M, et al. (2020) Smash++: an alignment-free and memory-efficient tool to find genomic rearrangements. GigaScience, 9(5).