# Snakemake

**RRID:** SCR_003475  
**Type:** Tool

## Proper Citation

Snakemake (RRID:SCR_003475)

## Resource Information

**URL:** [https://bitbucket.org/johanneskoester/snakemake/wiki/](https://bitbucket.org/johanneskoester/snakemake/wiki/)

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**Description:** A Python based language and execution environment for make-like workflows. The system supports the use of automatically inferred multiple named wildcards (or variables) in input and output filenames.

**Resource Type:** Resource, software resource

**References:** [PMID:22908215](https://www.ncbi.nlm.nih.gov/pubmed/22908215)

**Keywords:** python, workflow, bio.tools

**Parent Organization:** Google Code

**Availability:** MIT License

**Website Status:** Last checked up

**Abbreviations:** Snakemake

**Resource Name:** Snakemake

**Resource ID:** SCR_003475

**Alternate IDs:** OMICS_02299, biotools:snakemake

**Alternate URLs:** [https://bio.tools/snakemake](https://bio.tools/snakemake)
Ratings and Alerts

No rating or validation information has been found for Snakemake.

No alerts have been found for Snakemake.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 82 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.


Hendrickx APA, et al. (2021) -like genome architecture among carbapenemase-producing and the Netherlands. Microbial genomics, 7(5).


Istace B, et al. (2021) Sequencing and Chromosome-Scale Assembly of Plant Genomes, as a Use Case. Biology, 10(8).


