

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

SciPipe

RRID:SCR_017086

Type: Tool

Proper Citation

SciPipe (RRID:SCR_017086)

Resource Information

URL: <http://scipipe.org>

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Description: Software tool for writing scientific workflows of shell commands that depend on each other in Go programming language. Workflow library for agile development of complex and dynamic bioinformatics pipelines. Used in bioinformatics and cheminformatics, but works for any problem involving pipelines of commandline applications.

Resource Type: data processing software, software application, software library, software resource, software toolkit, source code, workflow software

Defining Citation: [DOI:10.1101/380808](https://doi.org/10.1101/380808)

Keywords: writing, scientific, workflow, shell, command, line, application, processing, software, Go

Funding: European Union's Horizon 2020 research and innovation programme 654241; National Bioinformatics Infrastructure Sweden (NBIS) ; Swedish e-Science Research Centre (SeRC) ; Swedish strategic research programme eSENCE

Availability: Free, Available for download, Freely available

Resource Name: SciPipe

Resource ID: SCR_017086

Alternate IDs: 10.5281/zenodo.1157941

Alternate URLs: <https://github.com/scipipe/scipipe/>, <https://pharmb.io/tool/scipipe/>

License: MIT

Record Creation Time: 20220129T080333+0000

Record Last Update: 20260905T132818+0000

Ratings and Alerts

No rating or validation information has been found for SciPipe.

No alerts have been found for SciPipe.

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 [Kravitz](#) .

Lampa S, et al. (2019) SciPipe: A workflow library for agile development of complex and dynamic bioinformatics pipelines. GigaScience, 8(5).