

# Resource Summary Report

Generated by [kravitz2](#) on Sep 4, 2026

## PAIPline

RRID:SCR\_024151

Type: Tool

### Proper Citation

PAIPline (RRID:SCR\_024151)

### Resource Information

**URL:** <https://gitlab.com/andreas.andrusch/pipeline>

**Proper Citation:** PAIPline (RRID:SCR\_024151)

**Description:** Software Python program to search for pathogen nucleic acid sequences in NGS datasets.Used for pathogen identification in metagenomic and clinical next generation sequencing samples.

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

**Defining Citation:** [PMID:30423069](#)

**Keywords:** Python program, pathogen nucleic acid sequences in NGS datasets search, pathogen identification, metagenomic and clinical next generation sequencing samples,

**Availability:** Free, Available for download, Freely available,

**Resource Name:** PAIPline

**Resource ID:** SCR\_024151

**Old URLs:** <https://sources.debian.org/src/pipeline/>

**License:** GNU GENERAL PUBLIC LICENSE

**Record Creation Time:** 20230824T050212+0000

**Record Last Update:** 20260903T235920+0000

### Ratings and Alerts

No rating or validation information has been found for PAIPline.

No alerts have been found for PAIPLine.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.