

Resource Summary Report

Generated by [NIF](#) on Apr 20, 2025

HaTSPiL

RRID:SCR_017059

Type: Tool

Proper Citation

HaTSPiL (RRID:SCR_017059)

Resource Information

URL: <https://github.com/dodomorandi/hatspil>

Proper Citation: HaTSPiL (RRID:SCR_017059)

Description: Software Python tool for high throughput sequencing analysis, focused on high reliability, modularity and customisability.

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

Keywords: next, generation, sequencing, pipeline, metadata, analysis, mutation, barcoding, customisability, modularity, bio.tools

Funding:

Availability: Free, Available for download, Freely available

Resource Name: HaTSPiL

Resource ID: SCR_017059

Alternate IDs: biotools:HatSPiL

Alternate URLs: <https://bio.tools/HaTSPiL>

License: MIT License

Record Creation Time: 20220129T080333+0000

Record Last Update: 20250420T014824+0000

Ratings and Alerts

No rating or validation information has been found for HaTSPiL.

No alerts have been found for HaTSPiL.

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 [NIF](#).

Morandi E, et al. (2019) HaTSPiL: A modular pipeline for high-throughput sequencing data analysis. PloS one, 14(10), e0222512.