

Resource Summary Report

Generated by [NIF](#) on Aug 30, 2026

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

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

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: 20260829T182555+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.