

# Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 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:** sequence analysis software, data analysis software, software resource, software application, data processing software

**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:** 20260728T043530+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 [nidm-terms](#) .

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