

# Resource Summary Report

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

## simhtsd

RRID:SCR\_006822

Type: Tool

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### Proper Citation

simhtsd (RRID:SCR\_006822)

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### Resource Information

**URL:** <http://sourceforge.net/projects/simhtsd/>

**Proper Citation:** simhtsd (RRID:SCR\_006822)

**Description:** Software that given a reference sequence, will create a large set of short nucleotide reads, simulating the output from today's high-throughput DNA sequencers, such as the Illumina Genome Analyzer II.

**Abbreviations:** simhtsd

**Synonyms:** Simulate High-Throughput Sequencing Data

**Resource Type:** software resource

**Keywords:** command-line, perl

**Availability:** GNU General Public License, v2, v3

**Resource Name:** simhtsd

**Resource ID:** SCR\_006822

**Alternate IDs:** OMICS\_00256

**Record Creation Time:** 20220129T080238+0000

**Record Last Update:** 20260905T132602+0000

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### Ratings and Alerts

No rating or validation information has been found for simhtsd.

No alerts have been found for simhtsd.

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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.