

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

simhtsd

RRID:SCR_006822

Type: Tool

Proper Citation

simhtsd (RRID:SCR_006822)

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: 20260725T190632+0000

Ratings and Alerts

No rating or validation information has been found for simhtsd.

No alerts have been found for simhtsd.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.