Resource Summary Report

Generated by NIF on Apr 26, 2025

HiLive

RRID:SCR_016134

Type: Tool

Proper Citation

HiLive (RRID:SCR_016134)

Resource Information

URL: https://gitlab.com/SimonHTausch/HiLive

Proper Citation: HiLive (RRID:SCR_016134)

Description: Software tool for performing read mapping that maps Illumina HiSeq sequencer read alignments when they are produced. Used in Next Generation Sequencing in time critical, clinical applications.

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

Defining Citation: PMID:27794555

Keywords: perform, read, mapping, sequence, alignment, analysis, Illumina, time, critical, clinical, application

Funding: the German Federal Ministry of Health IIA5-2512-FSB-725

Availability: Free, Available for download

Resource Name: HiLive

Resource ID: SCR_016134

Alternate IDs: OMICS_13393

Alternate URLs: https://sources.debian.org/src/hilive/

Old URLs: https://sourceforge.net/projects/hilive/

License: BSD License

Record Creation Time: 20220129T080329+0000

Record Last Update: 20250426T060527+0000

Ratings and Alerts

No rating or validation information has been found for HiLive.

No alerts have been found for HiLive.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.