

Resource Summary Report

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HiLive

RRID:SCR_016134

Type: Tool

Proper Citation

HiLive (RRID:SCR_016134)

Resource Information

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

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

Defining Citation: [PMID:27794555](https://pubmed.ncbi.nlm.nih.gov/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: 20260912T195835+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.