# **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

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

**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: 20250420T014800+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.