## **Resource Summary Report**

Generated by NIF on Apr 16, 2025

# <u>CLIIQ</u>

RRID:SCR\_009972 Type: Tool

**Proper Citation** 

CLIIQ (RRID:SCR\_009972)

#### **Resource Information**

URL: http://cliiq.sourceforge.net/Home

Proper Citation: CLIIQ (RRID:SCR\_009972)

**Description:** An algorithm to simultaneously identify and quantify expressed isoforms based on RNA-Seq data from multiple sample(s) in a population.

Abbreviations: CLIIQ

Resource Type: software resource

Funding:

Resource Name: CLIIQ

Resource ID: SCR\_009972

Alternate IDs: OMICS\_01272

Record Creation Time: 20220129T080256+0000

Record Last Update: 20250410T065937+0000

#### **Ratings and Alerts**

No rating or validation information has been found for CLIIQ.

No alerts have been found for CLIIQ.

### Data and Source Information

Source: <u>SciCrunch Registry</u>

## Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

Bernard E, et al. (2015) A convex formulation for joint RNA isoform detection and quantification from multiple RNA-seq samples. BMC bioinformatics, 16, 262.