# **Resource Summary Report**

Generated by NIF on Apr 19, 2025

## **QcReads**

RRID:SCR\_013002

Type: Tool

### **Proper Citation**

QcReads (RRID:SCR\_013002)

#### **Resource Information**

URL: http://sourceforge.net/projects/qcreads/

**Proper Citation:** QcReads (RRID:SCR\_013002)

**Description:** Provides an efficient tool for trimming adapter sequences and low quality sequences, in raw reads generated by the high throughput sequencing platforms.

Abbreviations: QcReads

Resource Type: software resource

**Funding:** 

Resource Name: QcReads

Resource ID: SCR\_013002

Alternate IDs: OMICS\_01088

**Record Creation Time:** 20220129T080313+0000

Record Last Update: 20250410T070323+0000

## Ratings and Alerts

No rating or validation information has been found for QcReads.

No alerts have been found for QcReads.

### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 1 mentions in open access literature.

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

Ma YF, et al. (2019) Population Genomics Analysis Revealed Origin and High-altitude Adaptation of Tibetan Pigs. Scientific reports, 9(1), 11463.