# **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on May 10, 2025

### qrqc

RRID:SCR\_006867

Type: Tool

### **Proper Citation**

qrqc (RRID:SCR\_006867)

#### Resource Information

URL: http://bioconductor.org/packages/2.8/bioc/html/qrqc.html

**Proper Citation:** qrqc (RRID:SCR\_006867)

**Description:** Software R package to quickly scan reads and gather statistics on base and quality frequencies, read length, k-mers by position, and frequent sequences. Produces graphical output of statistics for use in quality control pipelines, and an optional HTML quality report. S4 SequenceSummary objects allow specific tests and functionality to be written around the data collected.

Abbreviations: qrqc

**Synonyms:** quick read quality control, Quick Read Quality Control

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

**Keywords:** Quickly scan reads, read length, k-mers, position, frequent sequences, quality control pipeline, HTML quality report, bio.tools

Funding:

Availability: Free, Available for download, Freely available

Resource Name: qrqc

Resource ID: SCR\_006867

Alternate IDs: biotools:qrqc, OMICS\_01071

Alternate URLs: https://github.com/vsbuffalo/qrqc, https://bio.tools/qrqc

License: GNU General Public License, version 2

**Record Creation Time:** 20220129T080238+0000

**Record Last Update:** 20250509T055819+0000

### **Ratings and Alerts**

No rating or validation information has been found for qrqc.

No alerts have been found for qrqc.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Rosenberg MW, et al. (2022) Comprehensive molecular characterization of a rare case of Philadelphia chromosome-positive acute myeloid leukemia. Cold Spring Harbor molecular case studies, 8(6).

Pereira FL, et al. (2016) Evaluating the efficacy of the new Ion PGM Hi-Q Sequencing Kit applied to bacterial genomes. Genomics, 107(5), 189.