

# Resource Summary Report

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## [fastqz](#)

RRID:SCR\_001006

Type: Tool

### Proper Citation

fastqz (RRID:SCR\_001006)

### Resource Information

**URL:** <http://www.mattmahoney.net/dc/fastqz/>

**Proper Citation:** fastqz (RRID:SCR\_001006)

**Description:** Source code used to compress FASTQ files. FASTQ is DNA sequencing machine output.

**Resource Type:** software application, data management software, software resource, source code

**Defining Citation:** [PMID:23533605](#)

**Keywords:** compress, source code, fastq output, fastq dna, fastq sequencing, fastq compress, fastq compressor, fastq files, bio.tools

**Availability:** Free, Available for download, Freely available

**Resource Name:** fastqz

**Resource ID:** SCR\_001006

**Alternate IDs:** OMICS\_00956, biotools:fastqz

**Alternate URLs:** <https://bio.tools/fastqz>

**License:** BSD 2 clause license

**Record Creation Time:** 20220129T080205+0000

**Record Last Update:** 20260728T043106+0000

### Ratings and Alerts

No rating or validation information has been found for fastqz.

No alerts have been found for fastqz.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.