

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, software resource, source code, data management software

Defining Citation: [PMID:23533605](#)

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

Funding:

Availability: Free, Only available as source code, Available for download, libzpaq is required

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: 20250517T055450+0000

Ratings and Alerts

No rating or validation information has been found for fastqz.

No alerts have been found for fastqz.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.