

Resource Summary Report

Generated by [NIF](#) on Sep 10, 2026

MeQA

RRID:SCR_000317

Type: Tool

Proper Citation

MeQA (RRID:SCR_000317)

Resource Information

URL: <http://life.tongji.edu.cn/meqa/>

Proper Citation: MeQA (RRID:SCR_000317)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on August 31, 2022. Software for pre-processing, quality assessment, read distribution and methylation estimation for MeDIP-sequence datasets. It has the ability to quickly analyze sequence data for DNA methylation. This software integrates customized scripting and existing utilities tools that work on both paired end and single end data.

Resource Type: software resource

Defining Citation: [PMID:22199384](#)

Keywords: quality assessment, read distribution, methylation estimation, medip, medip-sequence, dna methylation, utilities tool, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: MeQA

Resource ID: SCR_000317

Alternate IDs: biotools:meqa, OMICS_00616

Alternate URLs: <https://bio.tools/meqa>

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260905T132418+0000

Ratings and Alerts

No rating or validation information has been found for MeQA.

No alerts have been found for MeQA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.