

Resource Summary Report

Generated by T1D on Sep 5, 2026

Artfastqgenerator

RRID:SCR_015979

Type: Tool

Proper Citation

Artfastqgenerator (RRID:SCR_015979)

Resource Information

URL: <https://sourceforge.net/projects/artfastqgen/>

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Description: Software to evaluate and improve the accuracy of sequencing error under different experimental conditions. It can identify which components of a system may be suboptimal and which regions of the genome may be problematic.

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

Defining Citation: [PMID:23152858](#)

Keywords: software, tool, sequencing, error, aligning, analysis, evaluation, genome, reference

Funding: Bobby Moore Fund for Cancer Research UK C1298/A8362

Availability: Free, Freely available, Available for download

Resource Name: Artfastqgenerator

Resource ID: SCR_015979

License: GNU General Public License v3.0

Record Creation Time: 20220129T080328+0000

Record Last Update: 20260903T235317+0000

Ratings and Alerts

No rating or validation information has been found for Artfastqgenerator.

No alerts have been found for Artfastqgenerator.

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 [T1D](#) .

Zhao L, et al. (2022) Quantitative PCR Assays for the Strain-Specific Identification and Enumeration of Probiotic Strain Lacticaseibacillus rhamnosus X253. Foods (Basel, Switzerland), 11(15).