

Resource Summary Report

Generated by [RRID](#) on Aug 15, 2026

[fqtrim](#)

RRID:SCR_028291

Type: Tool

Proper Citation

fqtrim (RRID:SCR_028291)

Resource Information

URL: <http://ccb.jhu.edu/software/fqtrim/>

Proper Citation: fqtrim (RRID:SCR_028291)

Description: Software tool for filtering and trimming next generation sequencing reads.

Resource Type: source code, software application, software resource

Keywords: filtering, trimming, next generation sequencing reads,

Availability: Free, Available for download, Freely available

Resource Name: fqtrim

Resource ID: SCR_028291

Alternate URLs: <https://github.com/gpertea/fqtrim/tree/v0.9.7>,
<https://zenodo.org/records/1185412>

License: Artistic-2.0 license

Record Creation Time: 20260415T055644+0000

Record Last Update: 20260815T183311+0000

Ratings and Alerts

No rating or validation information has been found for fqtrim.

No alerts have been found for fqtrim.

Data and Source Information

Usage and Citation Metrics

We found 10 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Wang X, et al. (2026) Colonic Engyodontium fungus triggers neutrophil antimicrobial activity to suppress Lactobacillus johnsonii-derived glutamic acid-maintained Tregs. The Journal of clinical investigation, 136(8).

Kong JY, et al. (2026) Cross-site oral and esophageal microbiome signatures define diagnostic patterns and reveal mechanistic drivers of ESCC. iScience, 29(8), 116667.

Zhang Z, et al. (2025) Different Efficacy of Five Soluble Dietary Fibers on Alleviating Loperamide-Induced Constipation in Mice: Influences of Different Structural Features. International journal of molecular sciences, 26(3).

Liu B, et al. (2023) Gastrointestinal Fermentable Polysaccharide Is Beneficial in Alleviating Loperamide-Induced Constipation in Mice. Nutrients, 15(20).

Liu M, et al. (2023) Effects of different forage proportions in fermented total mixed ration on muscle fatty acid profile and rumen microbiota in lambs. Frontiers in microbiology, 14, 1197059.

Qiao X, et al. (2022) Gut microbiota and fecal metabolic signatures in rat models of disuse-induced osteoporosis. Frontiers in cellular and infection microbiology, 12, 1018897.

Elhasnaoui J, et al. (2022) A Regulatory Axis between Epithelial Splicing Regulatory Proteins and Estrogen Receptor ?? Modulates the Alternative Transcriptome of Luminal Breast Cancer. International journal of molecular sciences, 23(14).

Zhu Y, et al. (2022) Gut Microbiota Correlates With Clinical Responsiveness to Erythropoietin in Hemodialysis Patients With Anemia. Frontiers in cellular and infection microbiology, 12, 919352.

Zhang M, et al. (2022) The gas production, ruminal fermentation parameters, and microbiota in response to Clostridium butyricum supplementation on in vitro varying with media pH levels. Frontiers in microbiology, 13, 960623.

Li W, et al. (2019) Variations in microbial community structure and functional gene expression in bio-treatment processes with odorous pollutants. Scientific reports, 9(1), 17870.