

Resource Summary Report

Generated by [NIF](#) on Aug 9, 2026

Reaper - Demultiplexing trimming and filtering sequencing data

RRID:SCR_001144

Type: Tool

Proper Citation

Reaper - Demultiplexing trimming and filtering sequencing data (RRID:SCR_001144)

Resource Information

URL: <http://www.ebi.ac.uk/~stijn/reaper/reaper.html>

Proper Citation: Reaper - Demultiplexing trimming and filtering sequencing data (RRID:SCR_001144)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 23, 2022. Software program for demultiplexing, trimming and filtering short read sequencing data.

Abbreviations: Reaper

Resource Type: software resource

Keywords: c, alignment, sequence, demultiplex, trim, filter, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Reaper - Demultiplexing trimming and filtering sequencing data

Resource ID: SCR_001144

Alternate IDs: biotools:reaper, OMICS_02157

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

Record Creation Time: 20220129T080205+0000

Record Last Update: 20260808T185738+0000

Ratings and Alerts

No rating or validation information has been found for Reaper - Demultiplexing trimming and filtering sequencing data.

No alerts have been found for Reaper - Demultiplexing trimming and filtering sequencing data.

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 [NIF](#) .

Dufour V, et al. (2015) Chimpanzee drumming: a spontaneous performance with characteristics of human musical drumming. Scientific reports, 5, 11320.