

Resource Summary Report

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parallel-fastq-dump

RRID:SCR_024150

Type: Tool

Proper Citation

parallel-fastq-dump (RRID:SCR_024150)

Resource Information

URL: <https://github.com/rvalieris/parallel-fastq-dump>

Proper Citation: parallel-fastq-dump (RRID:SCR_024150)

Description: Software wrapper to speed up downloading process by dividing the work into multiple threads.

Resource Type: software library, software resource, software toolkit

Keywords: wrapper, speed up downloading process, dividing the work into multiple threads,

Availability: Free, Available for download, Freely available,

Resource Name: parallel-fastq-dump

Resource ID: SCR_024150

Alternate URLs: <https://sources.debian.org/src/parallel-fastq-dump/>

License: MIT License

Record Creation Time: 20230824T050212+0000

Record Last Update: 20260728T043707+0000

Ratings and Alerts

No rating or validation information has been found for parallel-fastq-dump.

No alerts have been found for parallel-fastq-dump.

Data and Source Information

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Fuma K, et al. (2025) Prenatal inflammation impairs early CD11c-positive microglia induction and delays myelination in neurodevelopmental disorders. *Communications biology*, 8(1), 75.

Montero H, et al. (2025) Convergent losses of arbuscular mycorrhizal symbiosis in carnivorous plants. *The New phytologist*, 248(4), 2040.

Cullen JN, et al. (2025) Charting the equine miRNA landscape: An integrated pipeline and browser for annotating, quantifying, and visualizing expression. *PLoS genetics*, 21(9), e1011835.

Hoetzing M, et al. (2024) Geographic population structure and distinct intra-population dynamics of globally abundant freshwater bacteria. *The ISME journal*, 18(1).

Hwang W, et al. (2024) The Rice Online Expression Profiles Array Database Version 2 (ROADv2): An Interactive Atlas for Rice Functional Genomics. *Rice* (New York, N.Y.), 17(1), 75.

Li R, et al. (2024) Genome-wide characterization and evolution analysis of miniature inverted-repeat transposable elements in Barley (*Hordeum vulgare*). *Frontiers in plant science*, 15, 1474846.

Padwal MK, et al. (2023) Application of Machine Learning in Predicting Hepatic Metastasis or Primary Site in Gastroenteropancreatic Neuroendocrine Tumors. *Current oncology* (Toronto, Ont.), 30(10), 9244.