

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

Arvados

RRID:SCR_002223

Type: Tool

Proper Citation

Arvados (RRID:SCR_002223)

Resource Information

URL: <https://arvados.org/>

Proper Citation: Arvados (RRID:SCR_002223)

Description: Bioinformatics platform for storing, organizing, processing, and sharing genomic and other biomedical big data. Designed to make it easier for bioinformaticians to develop analyses, developers to create genomic web applications and IT administrators to manage large-scale compute and storage genomic resources. Designed to run on top of cloud operating systems such as Amazon Web Services and OpenStack. Currently, there are implementations that work on AWS and Xen+Debian/Ubuntu. Functionally, Arvados has two major sets of capabilities: (a) data management and (b) compute management.

Abbreviations: arvados

Resource Type: service resource, storage service resource, data repository

Keywords: mapreduce/hadoop, genomic, biomedical, data sharing, compute, data management, cloud

Availability: Free, Freely available

Resource Name: Arvados

Resource ID: SCR_002223

Alternate IDs: OMICS_01835

Alternate URLs: <https://sources.debian.org/src/arvados/>

License: Creative Commons Attribution-ShareAlike 4.0 International license

Record Creation Time: 20220129T080212+0000

Record Last Update: 20260728T043128+0000

Ratings and Alerts

No rating or validation information has been found for Arvados.

No alerts have been found for Arvados.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Bray S, et al. (2023) The Planemo toolkit for developing, deploying, and executing scientific data analyses in Galaxy and beyond. *Genome research*, 33(2), 261.

Brown AV, et al. (2018) Ten quick tips for sharing open genomic data. *PLoS computational biology*, 14(12), e1006472.

Chojnacki S, et al. (2017) Programmatic access to bioinformatics tools from EMBL-EBI update: 2017. *Nucleic acids research*, 45(W1), W550.