Resource Summary Report

Generated by RRID on Apr 19, 2025

Pipeliner

RRID:SCR_003171
Type: Tool

Proper Citation

Pipeliner (RRID:SCR_003171)

Resource Information

URL: https://github.com/brunonevado/Pipeliner

Proper Citation: Pipeliner (RRID:SCR_003171)

Description: Software for evaluating the performance of bioinformatics pipelines for Next Generation re-Sequencing.

Resource Type: software resource

Defining Citation: PMID:24890372

Keywords: standalone software, c++, bio.tools

Funding:

Resource Name: Pipeliner

Resource ID: SCR_003171

Alternate IDs: OMICS_04844, biotools:pipeliner

Alternate URLs: https://bio.tools/pipeliner

Record Creation Time: 20220129T080217+0000

Record Last Update: 20250410T064946+0000

Ratings and Alerts

No rating or validation information has been found for Pipeliner.

No alerts have been found for Pipeliner.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID.

Villamor-Payà M, et al. (2024) De novo TLK1 and MDM1 mutations in a patient with a neurodevelopmental disorder and immunodeficiency. iScience, 27(6), 109984.

Lim Y, et al. (2021) Exploration of alcohol use disorder-associated brain miRNA-mRNA regulatory networks. Translational psychiatry, 11(1), 504.

Smith RL, et al. (2021) Experimental Babesia rossi infection induces hemolytic, metabolic, and viral response pathways in the canine host. BMC genomics, 22(1), 619.

Tanwar S, et al. (2020) Type I IFN signaling in T regulatory cells modulates chemokine production and myeloid derived suppressor cells trafficking during EAE. Journal of autoimmunity, 115, 102525.

Federico A, et al. (2019) Pipeliner: A Nextflow-Based Framework for the Definition of Sequencing Data Processing Pipelines. Frontiers in genetics, 10, 614.

Bianco E, et al. (2015) A deep catalog of autosomal single nucleotide variation in the pig. PloS one, 10(3), e0118867.