

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 2, 2026

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

Availability: Free, Available for download, Freely available

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: 20260801T190228+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 [PRECISE-TBI](#) .

Villamor-Pay?? 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.