Resource Summary Report

Generated by ASWG on Aug 3, 2026

PASHA

RRID:SCR_004455
Type: Tool

Proper Citation

PASHA (RRID:SCR_004455)

Resource Information

URL: http://sourceforge.net/projects/pasha/

Proper Citation: PASHA (RRID:SCR_004455)

Description: A parallel short read assembler for large genomes using de Bruijn graphs.

Abbreviations: PASHA

Resource Type: software resource

Defining Citation: PMID:21867511

Resource Name: PASHA

Resource ID: SCR_004455

Alternate IDs: OMICS_00024

Record Creation Time: 20220129T080224+0000

Record Last Update: 20260801T190236+0000

Ratings and Alerts

No rating or validation information has been found for PASHA.

No alerts have been found for PASHA.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Boulanger M, et al. (2023) DeSUMOylation of chromatin-bound proteins limits the rapid transcriptional reprogramming induced by daunorubicin in acute myeloid leukemias. Nucleic acids research, 51(16), 8413.

Yahia Y, et al. (2023) RNA polymerase II CTD is dispensable for transcription and required for termination in human cells. EMBO reports, 24(9), e56150.

Pellow D, et al. (2023) Efficient minimizer orders for large values of k using minimum decycling sets. Genome research, 33(7), 1154.

Bejjani F, et al. (2023) Multiple Fra-1-bound enhancers showing different molecular and functional features can cooperate to repress gene transcription. Cell & bioscience, 13(1), 129.

Bellec M, et al. (2022) The control of transcriptional memory by stable mitotic bookmarking. Nature communications, 13(1), 1176.

Bejjani F, et al. (2021) Fra-1 regulates its target genes via binding to remote enhancers without exerting major control on chromatin architecture in triple negative breast cancers. Nucleic acids research, 49(5), 2488.

Escobar TM, et al. (2019) Active and Repressed Chromatin Domains Exhibit Distinct Nucleosome Segregation during DNA Replication. Cell, 179(4), 953.

Stafford JM, et al. (2018) Multiple modes of PRC2 inhibition elicit global chromatin alterations in H3K27M pediatric glioma. Science advances, 4(10), eaau5935.

Iasillo C, et al. (2017) ARS2 is a general suppressor of pervasive transcription. Nucleic acids research, 45(17), 10229.

Tanifuji G, et al. (2014) Nucleomorph and plastid genome sequences of the chlorarachniophyte Lotharella oceanica: convergent reductive evolution and frequent recombination in nucleomorph-bearing algae. BMC genomics, 15(1), 374.