Resource Summary Report

Generated by T1D on Jul 31, 2026

Bambino

RRID:SCR_005649
Type: Tool

Proper Citation

Bambino (RRID:SCR_005649)

Resource Information

URL: https://cgwb.nci.nih.gov/goldenPath/bamview/documentation/index.html

Proper Citation: Bambino (RRID:SCR_005649)

Description: A variant detector and graphical alignment viewer for next-generation sequencing data in the SAM/BAM format, which is capable of pooling data from multiple source files. Bambino may be launched online via Java Web Start or downloaded and run locally.

Abbreviations: Bambino

Synonyms: Bambino: a variant detector and alignment viewer for next-generation sequencing data in the SAM/BAM format

Resource Type: service resource, data analysis service, software resource, production service resource, analysis service resource

Defining Citation: PMID:21278191

Keywords: java, next-generation sequencing, bio.tools

Availability:
Https://cgwb.nci.nih.gov/goldenPath/bamview/documentation/Bambino_Click-thru-agreement-for-executable_final.txt

Resource Name: Bambino

Resource ID: SCR_005649

Alternate IDs: OMICS_00876, biotools:bambino

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

Record Creation Time: 20220129T080231+0000

Record Last Update: 20260729T044124+0000

Ratings and Alerts

No rating or validation information has been found for Bambino.

No alerts have been found for Bambino.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at T1D .

Umeda M, et al. (2025) Fusion oncoproteins and cooperating mutations define disease phenotypes in NUP98-rearranged leukemia. medRxiv : the preprint server for health sciences.

Wen J, et al. (2021) The landscape of coding RNA editing events in pediatric cancer. BMC cancer, 21(1), 1233.

Brady SW, et al. (2020) Pan-neuroblastoma analysis reveals age- and signature-associated driver alterations. Nature communications, 11(1), 5183.

Castel SE, et al. (2018) Modified penetrance of coding variants by cis-regulatory variation contributes to disease risk. Nature genetics, 50(9), 1327.

Xing Z, et al. (2018) Analysis of mutations in primary and metastatic synovial sarcoma. Oncotarget, 9(96), 36878.