Artemis: Genome Browser and Annotation Tool
RRID:SCR_004267
Type: Tool

Proper Citation

Artemis: Genome Browser and Annotation Tool (RRID:SCR_004267)

Resource Information

URL: http://www.sanger.ac.uk/resources/software/artemis/

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Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on February 28, 2023. Free genome browser and annotation tool that allows visualization of sequence features, next generation data and the results of analyses within the context of the sequence, and also its six-frame translation. Artemis is free software and is distributed under the terms of the GNU General Public License. Artemis is written in Java, and is available for UNIX, Macintosh and Windows systems. It can read EMBL and GENBANK database entries or sequence in FASTA, indexed FASTA or raw format. Other sequence features can be in EMBL, GENBANK or GFF format.

Abbreviations: Artemis

Resource Type: software resource

Defining Citation: PMID:11120685, DOI:10.1093/bioinformatics/btr703

Keywords: training tool, genome browser, gene annotation, java, bio.tools

Funding Agency: Wellcome Trust

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Artemis: Genome Browser and Annotation Tool

Resource ID: SCR_004267
Alternate IDs: nlx_28554, OMICS_00903, biotools:artemis


Record Creation Time: 20220129T080223+0000

Record Last Update: 20240424T182802+0000

Ratings and Alerts

No rating or validation information has been found for Artemis: Genome Browser and Annotation Tool.

No alerts have been found for Artemis: Genome Browser and Annotation Tool.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 354 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Butburee T, et al. (2024) General Pyrolysis for High-Loading Transition Metal Single Atoms on 2D-Nitro-Oxygeneous Carbon as Efficient ORR Electro catalysts. ACS applied materials & interfaces, 16(8), 10227.


Harris RA, et al. (2024) Complete genomes of Clostridium botulinum type B(F) isolates associated with a 1995 foodborne botulism outbreak from commercial pâté reveals a recombination event disrupting the ntnh gene. Microbial genomics, 10(1).

Demont T, et al. (2024) Exposure to worrisome topics can increase cognitive performance when incentivized by a performance goal. Scientific reports, 14(1), 1204.


Imam NG, et al. (2024) Comprehensive study of nanostructured Bi2Te3 thermoelectric materials - insights from synchrotron radiation XRD, XAFS, and XRF techniques. RSC advances, 14(3), 1875.


Papa-Ezdra R, et al. (2024) Outbreak of Pseudomonas aeruginosa High-Risk Clone ST309 Serotype O11 Featuring blaPER-1 and qnrVC6. Antibiotics (Basel, Switzerland), 13(2).


Jerez SA, et al. (2023) Vibrio type III secretion system 2 is not restricted to the Vibrionaceae and encodes differentially distributed repertoires of effector proteins. Microbial genomics, 9(4).