

Resource Summary Report

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Apollo

RRID:SCR_001936

Type: Tool

Proper Citation

Apollo (RRID:SCR_001936)

Resource Information

URL: <http://gmod.org/wiki/Apollo>

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Description: A standalone Java application with a GUI (graphical user interface) for editing genome annotations. Like GBrowse, it allows users to scroll and zoom in on areas of interest in a sequence; authorized users can edit annotations and write the changes back to the underlying database. Apollo can run off GFF3 or a Chado database, and it can also integrate with remote services, such as BLAST and Primer BLAST analyses.

Abbreviations: Apollo

Resource Type: software application, software resource

Defining Citation: [PMID:19439563](#), [PMID:12537571](#), [DOI:10.1186/gb-2002-3-12-research0082](#)

Keywords: java, genome annotation, genome, annotation, windows, mac os x, linux, solaris, unix, bio.tools, FASEB list

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Apollo

Resource ID: SCR_001936

Alternate IDs: OMICS_11761, biotools:apollo, OMICS_01933

Alternate URLs: <https://bio.tools/apollo>, <https://sources.debian.org/src/aragorn/>

Record Creation Time: 20220129T080210+0000

Record Last Update: 20260727T040311+0000

Ratings and Alerts

No rating or validation information has been found for Apollo.

No alerts have been found for Apollo.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 274 mentions in open access literature.

Listed below are recent publications. The full list is available at [dkNET](#) .

Mouter N, et al. (2025) Preferences for the Societal Impacts of a Pandemic when it Transitions into an Endemic: A Discrete Choice Experiment. *The patient*, 18(1), 49.

Xiao Z, et al. (2025) De novo assembly of the complete mitochondrial genomes of two *Camellia*-oil tree species reveals their multibranch conformation and evolutionary relationships. *Scientific reports*, 15(1), 2899.

Vlahakis N, et al. (2025) Fast event-based electron counting for small-molecule structure determination by MicroED. *Acta crystallographica. Section C, Structural chemistry*, 81(Pt 3), 116.

Pouncey AL, et al. (2025) Impact of patient sex on selection for abdominal aortic aneurysm repair: a discrete choice experiment. *BMJ open*, 15(2), e091661.

Xu D, et al. (2025) Comparative analysis of mitochondrial genomes of *Stemona tuberosa* Lour. reveals heterogeneity in structure, synteny, intercellular gene transfer, and RNA editing. *BMC plant biology*, 25(1), 23.

Tan M, et al. (2025) WOX11-OsPRX130 module confers rice drought tolerance by maintaining ROS homeostasis in rice root. *Rice (New York, N.Y.)*, 18(1), 55.

Turmel M, et al. (2025) Comparative Genomics of *Chloropicon primus* and *Chloropicon roscoffensis* Provide Insights into the Evolutionary Dynamics and Ecological Success of These Tiny Green Algae in Marine Environments. *Genome biology and evolution*, 17(7).

Arslan IG, et al. (2025) The role of environmental impact in healthcare providers' choices of inhalers for treatment of asthma and COPD: a discrete choice experiment. *BMC primary care*, 26(1), 278.

Chen Y, et al. (2025) CITED2 Binding to EP300 Regulates Human Spermatogonial Stem Cell Proliferation and Survival Through HSPA6. *Stem cells international*, 2025, 2362489.

Collins S??, et al. (2025) Relationship of Pulmonary Vascular Structure and Function With Exercise Capacity in Health and COPD. *Chest*, 167(2), 402.

Xia JB, et al. (2025) FoxO3 controls cardiomyocyte proliferation and heart regeneration by regulating Sfrp2 expression in postnatal mice. *Nature communications*, 16(1), 2532.

Bai X, et al. (2025) Unraveling the mitochondrial genome of the medicinal Chinese motherwort (*Leonurus japonicus*, Lamiaceae): structural dynamics, organelle-to-nuclear gene transfer, and evolutionary implications. *Frontiers in plant science*, 16, 1546449.

Li Y, et al. (2025) Mitochondrial genome structural variants and candidate cytoplasmic male sterility-related gene in sugarcane. *BMC genomics*, 26(1), 28.

Gui L, et al. (2025) Mitogenome of *Uncaria rhynchophylla*: genome structure, characterization, and phylogenetic relationships. *BMC genomics*, 26(1), 199.

Becking E, et al. (2025) Preferences of pregnant women and obstetric healthcare professionals on using noninvasive prenatal testing to predict adverse pregnancy outcomes: a discrete choice experiment. *BMJ open*, 15(1), e088304.

Makimoto K, et al. (2025) Upper-lobe CT imaging features improve prediction of lung function decline in COPD. *ERJ open research*, 11(3).

Buckell J, et al. (2025) Preferences for multi-cancer tests (MCTs) in primary care: discrete choice experiments of general practitioners and the general public in England. *British journal of cancer*, 133(3), 394.

van Beest EH, et al. (2025) Tracking neurons across days with high-density probes. *Nature methods*, 22(4), 778.

Cerqueira de Araujo A, et al. (2025) Genome sequences of four *Ixodes* species expands understanding of tick evolution. *BMC biology*, 23(1), 17.

Xu C, et al. (2025) Assembly and comparative analysis of the complete mitochondrial of *Spodiopogon sagittifolius*, an endemic and protective species from Yunnan, China. *BMC plant biology*, 25(1), 373.