

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: 20260821T193633+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 301 mentions in open access literature.

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

Xiao Z, et al. (2026) Comparative analysis of multi-chromosomal mitochondrial genome of *Bletilla striata*: homologous recombination mediated by repetitive sequences. *BMC plant biology*, 26(1).

Yu S, et al. (2026) Mitochondrial genome of *Iodes seguinii* reveals repeat-mediated recombination and phylogenetic insights in *Icacinales*. *BMC genomics*, 27(1).

Tian Z, et al. (2026) Assembly and comparative analysis of the complete mitochondrial genome of two species of *Argentina* (*Rosaceae*). *BMC genomics*, 27(1).

Zheng J, et al. (2026) Mitochondrial genome of the invasive weed *Amaranthus palmeri*: structural features, RNA editing, and evolutionary insights. *BMC plant biology*, 26(1).

Jimenez-Moreno AC, et al. (2026) Patient Preferences in Neuromuscular Diseases: Insights for Future Drug Development. *JIMD reports*, 67(4), e70100.

Zhang Q, et al. (2026) The complete chloroplast genome and phylogenetic analysis of *Guilandina minax* (Hance) G. P. Lewis (*Fabaceae*). *Mitochondrial DNA. Part B, Resources*, 11(3), 434.

Zhang H, et al. (2026) Complete mitochondrial genome of *Salix variegata* Franch.: assembly, characterization, and comparative analysis. *Frontiers in plant science*, 17, 1804453.

Bonjor N, et al. (2026) Characteristics of the mitochondrial genome of the invasive plant *Tagetes minuta* (*Asteraceae*) on the Qinghai-Tibet Plateau and its evolutionary significance. *BMC genomics*, 27(1).

Lust EER, et al. (2026) Preferences of Pregnant Women and Healthcare Professionals on First-Trimester Ultrasound Screening for Fetal Anomalies: A Discrete Choice Experiment. *Prenatal diagnosis*, 46(4), 523.

Li Y, et al. (2026) Structural characterization of the plastid genome and comparative mitochondrial genomics of *Eupatorium lindleyanum* (*Asteraceae*): evolutionary dynamics and phylogenetic insights. *BMC plant biology*, 26(1).

Thoral PJ, et al. (2026) Behaviour artificial intelligence technology to support the decision-making process of continuation or withdrawal of life-sustaining therapy. *Intensive care medicine experimental*, 14(1).

Zhang S, et al. (2026) Assembly and comparative analysis of the complete mitochondrial genome of the *Maclura tricuspidata*. *BMC genomics*, 27(1), 129.

Marsh K, et al. (2026) Valuing Maintenance of Visual Function in Retinitis Pigmentosa: A Discrete Choice Experiment with Patients and the General Public. *PharmacoEconomics - open*, 10(3), 543.

Deng Y, et al. (2026) Characterization of the Complete Mitochondrial Genome of *Pedicularis henryi* and Its Phylogenetic Implications in Lamiales. *Biology*, 15(7).

Wang J, et al. (2026) Mitochondrial genome assembly, comparative analysis of two *caprifoliaceae* species, and insights into adaptive evolution. *BMC plant biology*, 26(1).

Wright SJ, et al. (2026) Understanding the preferences of younger women for the delivery of a service to predict breast cancer risk: a discrete choice experiment. *BJC reports*, 4(1).

Lei M, et al. (2026) Assembly and Comparative Analysis of the Complete Mitochondrial Genome of *Corydalis ophiocarpa* (Papaveraceae). *Current issues in molecular biology*, 48(6).

Cao S, et al. (2026) Complete chloroplast genome of *Cupressus funebris* and comparative analyses within Cupressaceae. *BMC genomics*, 27(1).

Postma A, et al. (2026) Genome and transcriptome-based identification and expression profiling of chemosensory gene families across developmental stages and tissues in *Sirex noctilio* (Hymenoptera: Siricidae). *Insect molecular biology*, 35(3), 270.

Xu B, et al. (2026) The Mitochondrial Genome of *Curcuma longa*: A Large and Structurally Complex Genome with Extensive Intracellular DNA Transfer. *Genes*, 17(2).