

Resource Summary Report

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FigTree

RRID:SCR_008515

Type: Tool

Proper Citation

FigTree (RRID:SCR_008515)

Resource Information

URL: <http://tree.bio.ed.ac.uk/software/figtree>

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Description: A graphical viewer of phylogenetic trees and a program for producing publication-ready figures. It is designed to display summarized and annotated trees produced by BEAST.

Synonyms: FigTree

Resource Type: data processing software, software application, software resource, data visualization software

Keywords: data visualization software, graphical viewer, phylogenetic tree, annotate

Resource Name: FigTree

Resource ID: SCR_008515

Alternate IDs: OMICS_04268, nif-0000-30567

Alternate URLs: <https://sources.debian.org/src/figtree/>

Record Creation Time: 20220129T080247+0000

Record Last Update: 20260731T042755+0000

Ratings and Alerts

No rating or validation information has been found for FigTree.

No alerts have been found for FigTree.

Data and Source Information

Usage and Citation Metrics

We found 11323 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Grimwood RM, et al. (2026) A Novel Crustavirus as a Candidate Aetiology of Tail Fan Necrosis in New Zealand Red Rock Lobsters, *Jasus edwardsii*. *Journal of fish diseases*, 49(2), e70033.

Ansari F, et al. (2026) Characterisation of a Novel Jumbo Lytic *Aeromonas dhakensis* Bacteriophage P19 and Its Endolysin. *Microbial biotechnology*, 19(1), e70289.

Kincer LP, et al. (2026) Neurosymptomatic Cerebrospinal Fluid Escape Is a Central Nervous System-Focused Form of Antiretroviral Therapy Treatment Failure. *Open forum infectious diseases*, 13(1), ofaf659.

Shi M, et al. (2026) Genomic architecture and evolution of heterostyly: new insights from *Cordia subcordata* (Boraginaceae). *Molecular biology and evolution*, 43(1).

Wang G, et al. (2025) Genomic evidence for hybridization and introgression between blue peafowl and endangered green peafowl and molecular foundation of leucistic plumage of blue peafowl. *GigaScience*, 14.

Hata Y, et al. (2025) snRNA-seq analysis of the moss *Physcomitrium patens* identifies a conserved cytokinin-ESR module promoting pluripotent stem cell identity. *Developmental cell*, 60(13), 1884.

Záhonová K, et al. (2025) Diplonemid protists possess exotic endomembrane machinery, impacting models of membrane trafficking in modern and ancient eukaryotes. *Current biology : CB*, 35(7), 1508.

Han B, et al. (2025) Evolutionarily diverse caveolins share a common structural framework built around amphipathic disks. *The Journal of cell biology*, 224(9).

Rakesh S, et al. (2025) Unveiling the structural and functional implications of uncharacterized NSPs and variations in the molecular toolkit across arteriviruses. *NAR genomics and bioinformatics*, 7(2), lqaf035.

Nath R, et al. (2025) Lineage-Specific Class-A GPCR Dynamics Reflect Diverse Chemosensory Adaptations in Lophotrochozoa. *Molecular biology and evolution*, 42(3).

Arshad F, et al. (2025) A comprehensive water buffalo pangenome reveals extensive structural variation linked to population-specific signatures of selection. *GigaScience*, 14.

Fierro Morales JC, et al. (2025) Reduced cell-substrate adhesion formation promotes cell migration in *Dictyostelium*. *Molecular biology of the cell*, 36(8), ar98.

Garsmeur O, et al. (2025) The genomic footprints of wild *Saccharum* species trace domestication, diversification, and modern breeding of sugarcane. *Cell*, 188(25), 7252.

Nguyen HH, et al. (2025) WaveSeekerNet: accurate prediction of influenza A virus subtypes and host source using attention-based deep learning. *GigaScience*, 14.

Liu S, et al. (2025) Mitogenomes of museum specimens provide new insight into species classification and recently reduced diversity of highly endangered *Nomascus gibbons*. *Integrative zoology*, 20(3), 674.

Lobos-Ovalle D, et al. (2025) Morphological and molecular data on the bat flies, *Basilia silvae* (Brèthes, 1913) and *Trichobius parasiticus* Gervais, 1844 parasitizing Chilean bats. *Medical and veterinary entomology*, 39(3), 422.

Fan Q, et al. (2025) Insights into the molecular network characteristics of major HIV-1 subtypes in developed Eastern China: a study based on comprehensive molecular surveillance data. *Infection*, 53(3), 905.

Janssen AB, et al. (2025) PneumoBrowse 2: an integrated visual platform for curated genome annotation and multiomics data analysis of *Streptococcus pneumoniae*. *Nucleic acids research*, 53(D1), D839.

Guarneri I, et al. (2025) *Amphioxus* (*Branchiostoma lanceolatum*) in the North Adriatic Sea: ecological observations and spawning behavior. *Integrative zoology*, 20(2), 331.

Zhou Z, et al. (2025) Ferruginous hemeprotein HhuH facilitates the cadmium adsorption and chromium reduction in *Stenotrophomonas* sp. SY1. *Applied and environmental microbiology*, 91(1), e0209724.