

# Resource Summary Report

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## FigTree

RRID:SCR\_008515

Type: Tool

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### Proper Citation

FigTree (RRID:SCR\_008515)

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### Resource Information

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

**Proper Citation:** FigTree (RRID:SCR\_008515)

**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:** software resource, data visualization software, data processing software, software application

**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:** 20260803T040526+0000

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### Ratings and Alerts

No rating or validation information has been found for FigTree.

No alerts have been found for FigTree.

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### 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 [ASWG](#) .

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.