

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, data visualization software, software application, software resource

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: 20260905T132624+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 12345 mentions in open access literature.

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

Campbell AS, et al. (2026) Molecular insights into electroreceptor ribbon synapses from differential gene expression in sturgeon lateral line organs. *Journal of anatomy*, 248(5), 784.

Daniel K, et al. (2026) Pre-existing neutralizing antibodies against cattle-transmitted influenza A virus H5N1 are detectable in unexposed individuals. *Immunity*, 59(2), 494.

Mao C, et al. (2026) Chromosome-level genome assembly and transcriptomes of the leaf insect *Cryptophyllum westwoodii* provide insights into the evolution of leaf-like masquerade. *GigaScience*, 15.

Dong Z, et al. (2026) Piezo3 is a novel mechanosensitive Piezo ion channel in vertebrates. *bioRxiv : the preprint server for biology*.

Hao L, et al. (2026) Imagin: An Integrase-Like Gene Conserved Across Malacostracan Crustaceans Derived From a Ginger1 DNA Transposon. *Genome biology and evolution*, 18(2).

Liu S, et al. (2026) Comparative Analysis of Mitochondrial Genomes and Phylogeny of Barbastelle Bats Across China. *Ecology and evolution*, 16(1), e72949.

Zhou W, et al. (2026) Comparative analysis of gene family evolution demonstrates expansion of digestive, immunity and olfactory functions in the black soldier fly (*Hermetia illucens*) lineage. *Heredity*, 135(3), 187.

Shi H, et al. (2026) Emergence and evolution of HoBi-like pestivirus clades a and c in sheep from central China. *BMC veterinary research*, 22(1), 82.

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

Feng H, et al. (2026) The complete chloroplast genome and phylogenetic analysis of *Fraxinus paxiana* Lingelsh. 1907 (Oleaceae). *Mitochondrial DNA. Part B, Resources*, 11(2), 232.

Tsuruga K, et al. (2026) A new species of *Curtobacterium* from apple fruit: *Curtobacterium pomorum* sp. nov. resistant to heavy metals. *Microbiology spectrum*, 14(2), e0203125.

van Jaarsveld WJ, et al. (2026) Adverse effects of grapevine trunk diseases on canes produced from infected rootstock mother blocks. *Pest management science*, 82(4), 2973.

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.

Krupa EA, et al. (2026) First detection of one of the tick-borne lymphadenopathy (TIBOLA) etiological agent in ticks from a highly frequented sub-urban forest near Paris, France. *PLOS global public health*, 6(1), e0005705.

Westphalen MC, et al. (2026) Unveiling the hidden diversity of neotropical *Steccherinum* and allied genera (Steccherinaceae, Basidiomycota). *IMA fungus*, 17, e182915.

Zhou T, et al. (2026) Pleistocene Climate Oscillations and Geographic Barriers Shaped the Phylogeographic Structure of *Anaplecta omei* (Blattodea, Blattoidea, Anaplectidae) in Southern China: Evidence From Mitochondrial Genomes. *Ecology and evolution*, 16(2), e73086.

Choudhury ST, et al. (2026) Genome-wide co-occurrence patterns link mobile genetic elements, antimicrobial resistance and defense systems in *Pseudomonas aeruginosa*. *BMC genomics*, 27(1), 226.

Ahsan T, et al. (2026) Metabolomic profiling of *Bacillus amyloliquefaciens* MU-10 and its antagonistic effects on *Cercospora arachidicola*. *BMC microbiology*, 26(1).

Chang CX, et al. (2026) Morphological and phylogenetic analyses reveal three new species of *Gibellula* (Cordycipitaceae, Hypocreales) from spiders. *MycKeys*, 127, 135.

Felicio D, et al. (2026) Retrotransposition Events Shape the Evolution of the Ataxin-3 Gene Family in Primates. *Genome biology and evolution*, 18(3).