

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

TREE-PUZZLE

RRID:SCR_024382

Type: Tool

Proper Citation

TREE-PUZZLE (RRID:SCR_024382)

Resource Information

URL: <http://www.tree-puzzle.de>

Proper Citation: TREE-PUZZLE (RRID:SCR_024382)

Description: Software tool to reconstruct phylogenetic trees from molecular sequence data by maximum likelihood. Allows analysis of large data sets and automatically assigns estimations of support to each internal branch. Computes pairwise maximum likelihood distances as well as branch lengths for user specified trees. Conducts statistical tests on the data set.

Synonyms: tree-puzzle

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

Keywords: reconstruct phylogenetic trees, molecular sequence data, maximum likelihood, compute pairwise maximum likelihood distances, conduct statistical tests, data analysis,

Availability: Free, Available for download, Freely available,

Resource Name: TREE-PUZZLE

Resource ID: SCR_024382

Alternate IDs: OMICS_18458

Alternate URLs: <https://sources.debian.org/src/tree-puzzle/>

Old URLs: <https://sources.debian.org/src/tree-ppuzzle/>

Record Creation Time: 20230830T050217+0000

Record Last Update: 20260905T133030+0000

Ratings and Alerts

No rating or validation information has been found for TREE-PUZZLE.

No alerts have been found for TREE-PUZZLE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Nylund A, et al. (2023) Genotyping tool for salmonid gill pox virus (SGPV) obtained from farmed and wild Atlantic salmon (*Salmo salar*). Archives of virology, 168(10), 249.

Mull N, et al. (2023) Novel Ozark Orthohantavirus in Hispid Cotton Rats (*Sigmodon hispidus*), Arkansas, USA. Emerging infectious diseases, 29(12), 2576.

White SK, et al. (2018) A new "American" subgroup of African-lineage Chikungunya virus detected in and isolated from mosquitoes collected in Haiti, 2016. PloS one, 13(5), e0196857.

Sánchez-Busó L, et al. (2013) Genetic Characterization of *Legionella pneumophila* Isolated from a Common Watershed in Comunidad Valenciana, Spain. PloS one, 8(4), e61564.