

# Resource Summary Report

Generated by [ASWG](#) on Aug 4, 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:** software resource, data processing software, software application, data analysis software

**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:** 20260804T043839+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 3 mentions in open access literature.

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

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

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.

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.