

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

Treerecs

RRID:SCR_024497

Type: Tool

Proper Citation

Treerecs (RRID:SCR_024497)

Resource Information

URL: <https://gitlab.inria.fr/Phylophile/Treerecs>

Proper Citation: Treerecs (RRID:SCR_024497)

Description: Open source, species and gene tree reconciliation software. Software integrated phylogenetic tool, from sequences to reconciliations. Used to correct, rearrange and reroot gene trees with regard to given species tree.

Resource Type: software application, software resource

Defining Citation: [PMID:33085745](#)

Keywords: species and gene tree reconciliation, integrated phylogenetic, from sequences to reconciliations, correct gene trees, rearrange gene trees, reroot gene trees,

Availability: Free, Available for download, Freely available

Resource Name: Treerecs

Resource ID: SCR_024497

Alternate URLs: <https://project.inria.fr/treerecs/>

License: GNU AGPL v 3.0

Record Creation Time: 20231002T161336+0000

Record Last Update: 20260919T195829+0000

Ratings and Alerts

No rating or validation information has been found for Treerecs.

No alerts have been found for Treerecs.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Dufayard JF, et al. (2021) RapGreen, an interactive software and web package to explore and analyze phylogenetic trees. NAR genomics and bioinformatics, 3(3), lqab088.