

# Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

## Kdetrees

RRID:SCR\_004522

Type: Tool

### Proper Citation

Kdetrees (RRID:SCR\_004522)

### Resource Information

**URL:** <http://cran.r-project.org/web/packages/kdetrees/>

**Proper Citation:** Kdetrees (RRID:SCR\_004522)

**Description:** R package using a non-parametric method for estimating distributions of phylogenetic trees, with the goal of identifying trees that are significantly different from the rest of the trees in the sample.

**Synonyms:** kdetrees: Nonparametric method for identifying discordant phylogenetic trees

**Resource Type:** software resource

**Defining Citation:** [PMID:24764459](#)

**Keywords:** applet, mac os x, unix/linux, windows, r, bio.tools

**Availability:** GNU General Public License, v2

**Resource Name:** Kdetrees

**Resource ID:** SCR\_004522

**Alternate IDs:** biotools:kdetrees, OMICS\_04172

**Alternate URLs:** <https://github.com/grady/kdetrees>, <https://bio.tools/kdetrees>

**Record Creation Time:** 20220129T080225+0000

**Record Last Update:** 20260725T190552+0000

### Ratings and Alerts

No rating or validation information has been found for Kdetrees.

No alerts have been found for Kdetrees.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.