## **Resource Summary Report**

Generated by RRID on May 12, 2025

## **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

**Funding:** 

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: 20250420T014225+0000

# Ratings and Alerts

No rating or validation information has been found for Kdetrees.

No alerts have been found for Kdetrees.

### **Data and Source Information**

Source: SciCrunch Registry

# **Usage and Citation Metrics**

We have not found any literature mentions for this resource.