

Package: phylocanvas (via r-universe)

September 10, 2024

Title Interactive Phylogenetic Trees Using the 'Phylocanvas'
JavaScript Library

Version 0.1.3

Description Create and customize interactive phylogenetic trees using the 'phylocanvas' JavaScript library and the 'htmlwidgets' package. These trees can be used directly from the R console, from 'RStudio', in Shiny apps, and in R Markdown documents. See <http://phylocanvas.org/> for more information on the 'phylocanvas' library.

URL <https://github.com/zachcp/phylocanvas>, <http://phylocanvas.org/>

BugReports <https://github.com/zachcp/phylocanvas/issues>

Depends R (>= 3.3.1)

Imports ape (>= 4.0), phylobase, htmlwidgets, methods (>= 3.3.0)

Suggests magrittr, knitr, rmarkdown

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Encoding UTF-8

LazyData true

RoxygenNote 6.0.1

VignetteBuilder knitr

Collate 'load.tree.R' 'as_tree.R' 'get.descendants.R'
'nodes.branches.R' 'phylocanvas.R'

Repository <https://zachcp.r-universe.dev>

RemoteUrl <https://github.com/zachcp/phylocanvas>

RemoteRef HEAD

RemoteSha afc7dc0646f699873c4ff675ae5f69bcb923bd36