Linking JUICE and (R):



New developments in visualization of unconstrained ordination analysis



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JUICE 7.0

is a Windows application for editing, classification and analysis of large phytosociological tables and other ecological data. It has been by Lubomír (Masaryk University Brno) since 1998, written in Visual Basic and optimized for Windows OS (but operating also under Linux via Wine emulator). JUICE can also autoor semi-automatically operate other commercial or freeware software, such as PC-ORD, CANOCO, SYNTAX or Mulva.

developed

For more information about JUICE and free program download, visit http://www.sci.muni.cz/botany/juice/ or http://tinyurl.com/cwtrf9.

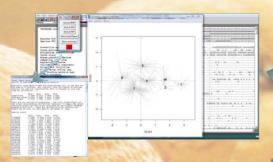
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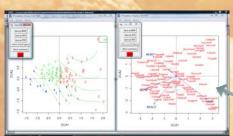
Linking JUICE and R offers quick and effective availability of selected R analytical and visualization functions: JUICE offers user friendly environment of vegetation data handling and import, R works as an engine for analysis and visualization.

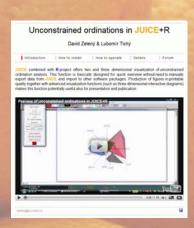
Recently implemented unconstrained ordination analyses are Detrended Correspondence Analysis, Principal Component Analysis and Non-metric Multidimensional Scaling, all available in vegan library written by Oksanen et al.

R-project

- is a free software environment statistical computing graphics, running on a wide variety of platforms. Following R packages are used by JUICE:
- vegan (Oksanen, Kindt, Legendre, O'Hara et al.) - number of tools for analysis of ecological communities, diversity and multivariate analysis;
- ordijuice (Zelený) connecting JUICE and R and offering some additional visualization tools;
- rgl (Adler & Murdoch) three dimensional interactive ordination diagrams;
- geometry (Grasman & Gramacy) calculation of convex hulls;
- tcltk and tkrgl (Murdoch & Chen) user interface for interaction with figures.



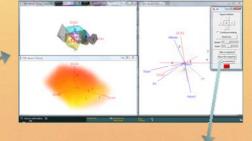


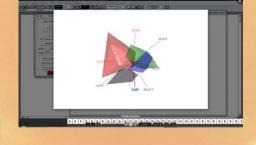












Technical solutions

Link between JUICE and R is mediated by sharing the files via R/bin directory and using libraries ordijuice. JUICE offers wizard for selection of appropriate analysis parameters and type of graphical output. Data are saved into R/bin file, and R is initialized to run in a batch mode by sending the script with selected parameters into Windows command line. R draws the figure onto graphical device (X11 or rgl) and opens Tcl/Tk panel; after either the device or Tcl/Tk bar is closed, R is termi-

nated. This design allows several R sessions to run simultaneously using the same data, but different parameters.

Updates of ordijuice library are available from R repository in http://www.sci.muni.cz/botany/zeleny/R. If ordinations are in use, once per day is checked for new updates and user is