

Conference Abstract

Introducing ‘The *bdverse*’: a family of R packages for biodiversity data

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Abstract

[The *bdverse*](#) is a collection of packages that form a general framework for facilitating biodiversity science in R. We build it to serve as a sustainable and agile infrastructure that enhances the value of biodiversity data by allowing users to conveniently employ R, for data exploration, quality assessment, data cleaning, and standardization. The *bdverse* supports users with and without programming capabilities. It includes six unique packages in a hierarchical structure — representing different functionality levels (Fig. 1). Major features of three core packages will be highlighted and demonstrated: (i) [bdDwC](#) provides an interactive Shiny app and a set of functions for standardizing field names in compliance with Darwin Core (DwC) format; (ii) [bdchecks](#) is an infrastructure for performing, filtering and managing various biodiversity data checks; (iii) [bdclean](#) is a user-friendly data cleaning Shiny app for the inexperienced R user. It provides features to manage complete workflow for biodiversity data cleaning, including data upload; user input - in order to adjust cleaning procedures; data cleaning; and finally, generation of various reports and versions of the data.

We are now working on submitting the *bdverse* packages to [rOpenSci software review](#), and as soon as the packages meet core requirements, we will officially release the *bdverse*. The *bdverse* project won the [2nd prize in the 2018 Ebbe Nielsen Challenge](#).

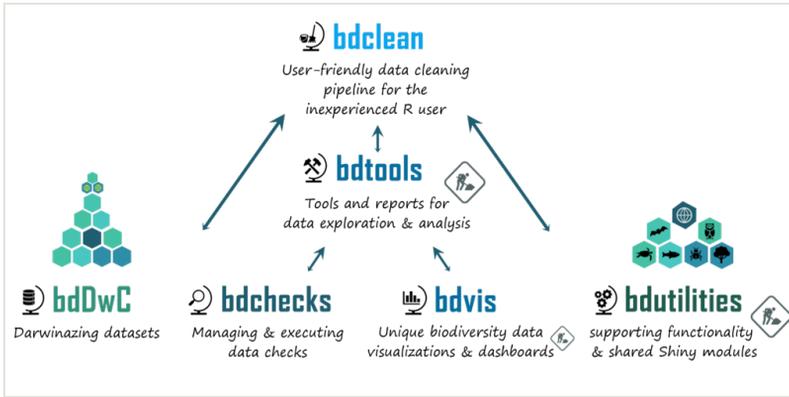


Figure 1.

A schematic representation of the *bdverse*, a toolbelt of packages for handling biodiversity data in R. Repositories of all packages can be publicly accessed via GitHub (<https://github.com/bd-R>).

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