

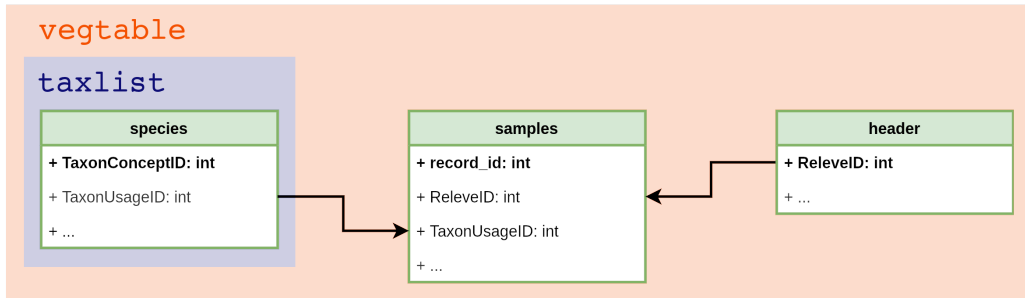
# Basics on the work with vegetation-plots in vegetable

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# Introduction

## vegetable-taxlist complex



**species:** Taxonomic list with taxon attributes (taxonomic ranks, functional traits, chorology, ...)

**header:** Attributes associated to plots (size, location, ecological attributes, ...)

**samples:** Records of taxa in plots (frequency, abundance, ...)

## Why R?

- Increasing popularity among scientists
- Collaborative data assessment
  - R-script + R-image
  - R-markdown (+ R-script) + R-image (+ Data)
  - Project Folder (ZIP file) or a repository (GitHub, GitLab, RStudio Cloud, etc.)
- Documenting and ensuring repeatability.
- Teaching purposes.

Some important information can be extracted with following commands:

```
# How many plots are in the data set?
```

```
nrow(Kenya_veg$header)
```

```
## [1] 1946
```

```
# How many plots per data source?
```

```
aggregate(ReleveID ~ REFERENCE, data=Kenya_veg$header,  
          FUN=function(x) length(unique(x)))
```

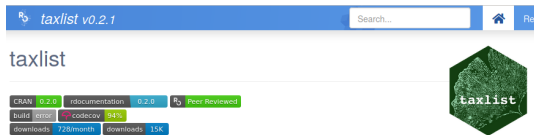
```
##   REFERENCE ReleveID  
## 1      2974      262  
## 2      3011      693  
## 3      3012      610  
## 4      2331      325  
## 5      3506       56
```

# Introduction

<https://docs.ropensci.org/taxlist/>

<https://github.com/kamapu/vegetable>

- Both packages developed on **GitHub**
- Both packages accessible at **CRAN**



## Introduction

`taxlist` is a package designed to handle and assess taxonomic lists in **R**, providing an object class and functions in **S4** language. The homonymous object class `taxlist` was originally designed as a module for taxa recorded in vegetation-plot observations (see `vegetable`), but became as an independent object with the ability of contain not only lists of species but also synonymy, hierarchical taxonomy, and functional traits (attributes of taxa).

Alternative installing commands:

```
## Installing from CRAN
install.packages("vegetable", dependencies=TRUE)

## Installing from GitHub
library(devtools)
install_github("ropensci/taxlist",
               build_vignettes=TRUE)
install_github("kamapu/vegetable")
```

## vegetable in retrospective

- **2015:** Experiments on the basis of `vegdata`
- **2017:** First version at **CRAN**
- **2018:** Publication of `taxlist` by Alvarez & Luebert in *Biodiversity Data Journal*
- **2020:** `taxlist` accepted in *rOpenSci*



Biodiversity Data Journal 6: e23635  
doi: 10.3897/BDJ.6.e23635



R Package

## The `taxlist` package: managing plant taxonomic lists in R

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ZooBank: [um:tsid:zoobank.org/pub:E16342C5-6C4D-4690-8042-7329E43D2292](https://doi.org/10.3897/BDJ.6.e23635)

## Abstract

Taxonomic lists are crucial elements of vegetation-plot databases and provide the links between original entries, reference taxon views and different taxon concepts. We introduce the R package `taxlist` in the context of object-oriented modelling for taxonomic lists. This package provides a data structure based on species lists in Turboveg, which is a software broadly used for the storage of vegetation-plot databases and implements functions for importing and handling them prior to statistical analysis. We also present a schema for relational databases, compatible with `taxlist` objects and recommend its use for handling diversity records.

## Keywords

ecoinformatics, database, taxon concept, taxon view, Turboveg, vegetable

# Dissecting vegetable

**S4** objects (definition, prototype, validation, methods) organized in slots.

Resembling **Turboveg**'s dbf-files and relational databases.

**description** (metadata)

**species** (a taxlist object)

**header** (plot information)

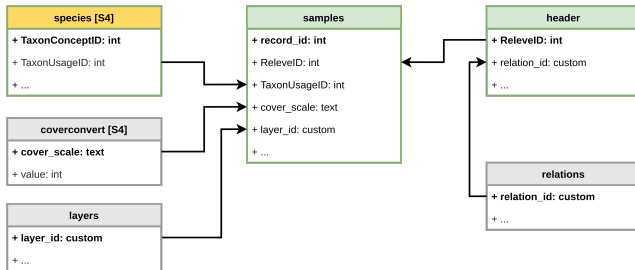
**relations** (tables with relationships to *header*)

**samples** (record table)

**layers** (tables with relationships to *samples*)

**coverconvert** (coverage/abundance conversions)

## Slots in vegetable objects



# Dissecting vegetable

## Slot `species` (`obj@species`)

Taxonomic information of *recorded organisms* stored in a `taxlist` object.

- Relationships of usage names and taxon concepts
- Parent-child relationships and taxonomic ranks
- Functional traits
- Connected through **TaxonUsageID**

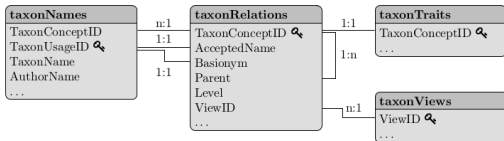
`taxlist` supports different degrees of information completeness/resolution

```
Kenya_veg@species
```

```
## object size: 554 Kb
## validation of 'taxlist' object: TRUE
##
## number of taxon usage names: 3164
## number of taxon concepts: 2392
## trait entries: 102
## number of trait variables: 1
## taxon views: 3
##
## concepts with parents: 2237
## concepts with children: 957
##
## hierarchical levels: form < variety < subspecies < species
## number of concepts in level form: 0
## number of concepts in level variety: 65
## number of concepts in level subspecies: 52
## number of concepts in level species: 1422
## number of concepts in level complex: 0
## number of concepts in level genus: 699
## number of concepts in level family: 154
```

# Dissecting vegetable

## Slot species (obj@species)



Alvarez & Luebert (2018)

```
summary(Kenya_veg@species, "Cyperus involucratus",  
        secundum="secundum")
```

```
## -----  
## concept ID: 51757  
## view ID: 1 - African Plant Database (2012)  
## level: species  
## parent: 54853 Cyperus L.  
##  
## # accepted name:  
## 51757 Cyperus involucratus Rottb.  
##  
## # synonyms (1):  
## 53973 Cyperus flabelliformis Rottb.  
## -----
```

```
summary(Kenya_veg@species, 54853, secundum="secundum")
```

```
## -----  
## concept ID: 54853  
## view ID: 2 - Taxonomic Name Resolution Service (2018)  
## level: genus  
## parent: 55959 Cyperaceae NA  
##  
## # accepted name:  
## 54855 Cyperus L.  
## -----
```



# Dissecting vegetable

## Slot header (obj@header)

Main table (data frame) including information on plot observations (relevés).

- Identifiers
- Time and location of records
- Environmental information (e.g. soil sample analyses)
- Remarks
- *Statistics*

Variable **ReleveID** is mandatory in header.

```
head(Kenya_veg@header)
```

```
##      ReleveID COUNTRY REFERENCE TABLE_NR NR_IN_TAB
## 358      358      KE      2974         1         1
## 359      359      KE      2974         1         2
## 360      360      KE      2974         1         3
## 361      361      KE      2974         1         4
## 362      362      KE      2974         1         5
## 363      363      KE      2974         1         6
##      COVERSCALE DATE SURF_AREA ALTITUDE EXPOSITION
## 358          01 <NA>      NA      NA      <NA>
## 359          01 <NA>      NA      NA      <NA>
## 360          01 <NA>      NA      NA      <NA>
## 361          01 <NA>      NA      NA      <NA>
## 362          01 <NA>      NA      NA      <NA>
## 363          01 <NA>      NA      NA      <NA>
##      INCLINATIO COV_TOTAL TREE_HIGH      REMARKS
## 358          NA      NA      NA Mount Nyiro
## 359          NA      NA      NA Mount Nyiro
## 360          NA      NA      NA Mount Nyiro
## 361          NA      NA      NA Mount Nyiro
## 362          NA      NA      NA Mount Nyiro
## 363          NA      NA      NA Mount Nyiro
##      LONGITUDE LATITUDE PH_H2O
## 358    36.8167    2.1833      NA
## 359    36.8167    2.1833      NA
## 360    36.8167    2.1833      NA
```

# Dissecting vegetable

## Slot header (obj@samples)

Data frame including the records of taxa (inserted as taxon usage names) in plot observations (relevés).

- Occurrence, frequency, abundance, sociability, etc.
- Relationships to layers.
- Relationships to collected specimens.
- Relationships to individuals.

Variables **ReleveID** and **TaxonUsageID** are mandatory.

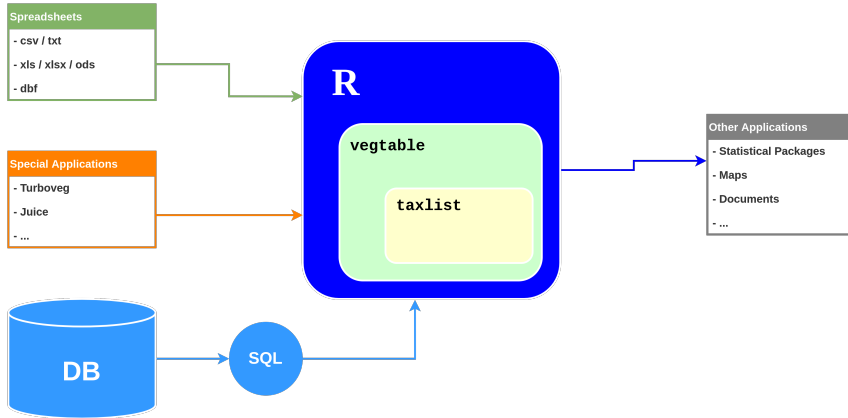
```
head(Kenya_veg@samples)
```

##	ReleveID	TaxonUsageID	COVER_CODE	LAYER
## 5658	358	18	+	0
## 5659	358	220	r	0
## 5660	358	233	+	0
## 5661	358	287	+	0
## 5662	358	407	r	0
## 5663	358	54983	r	0

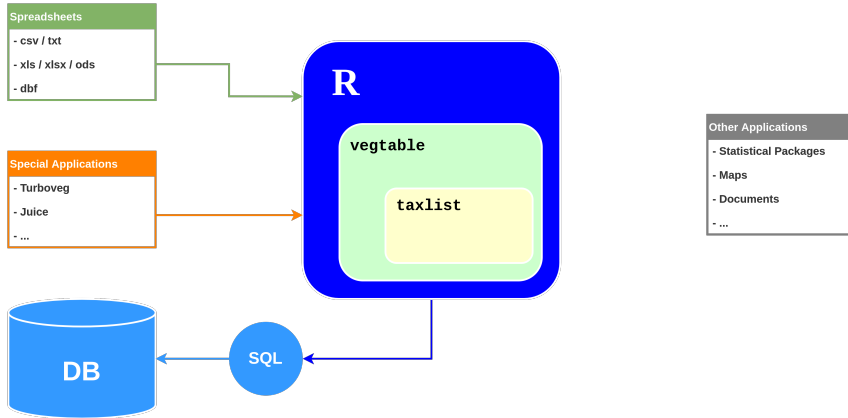
  

##	SOCIABILIT	INDIVID	br_b1	b_bbds
## 5658	<NA>	<NA>	+	<NA>
## 5659	<NA>	<NA>	r	<NA>
## 5660	<NA>	<NA>	+	<NA>
## 5661	<NA>	<NA>	+	<NA>
## 5662	<NA>	<NA>	r	<NA>
## 5663	<NA>	<NA>	r	<NA>

# Recipe

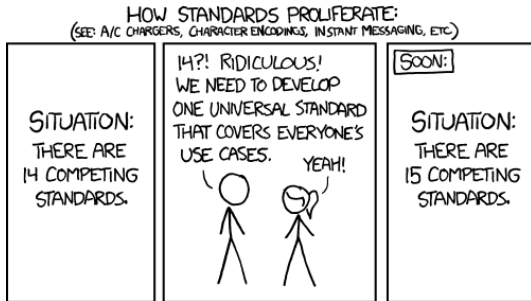


# Recipe



# For the Future

- Multiple taxon views implemented in taxlist
- Import/export functions for **Veg-X**
- Better interaction with **Juice** (ideally in Linux)
- Better interaction with **Turboveg**
- Implementation in **sPlot**
- Contributions to **GIVD**
- Calculation of diversity indices (Shannon's H, Constancy values, Mean cover, etc.)
- Summaries for communities



[xkcd.com](http://xkcd.com)

# THANKS!

## For discussion:

<https://stackoverflow.com/users/5846398/miguel-alvarez>

[https://www.researchgate.net/profile/Miguel\\_Alvarez18](https://www.researchgate.net/profile/Miguel_Alvarez18)

<https://www.facebook.com/groups/ecologyinr>

<https://github.com/kamapu/vegetable/issues>



...and also visit me at <https://kamapu.github.io/>