

vegetable: An R object for vegetation-plot data sets

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Why R?

- Collaborative data assessment
 - (R-script + R-image).
- Documenting and ensuring repeatability.
- Teaching purposes.

taxlist-vegetable complex

- taxlist: taxonomic lists
- vegetable: bio-diversity records



Introduction

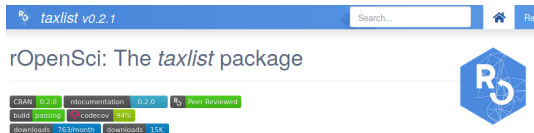
<https://docs.ropensci.org/taxlist/>

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

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

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

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



The screenshot shows the rOpenSci website for the *taxlist* v0.2.1 package. The header includes the package name and a search bar. Below the header, the text "rOpenSci: The *taxlist* package" is displayed. To the right is the R logo. Below the text, there are several status badges: CRAN 0.2.0, rdocumentation 0.2.0, Peer Reviewed, build passing, codecov 94%, and downloads 763/months and 15K. The R logo is a blue hexagon with a white 'R' and a circular arrow.

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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SWEA-Dataveg

<https://kamapu.github.io/sweadataveg>

<https://www.givd.info/ID/AF-00-006>

Conservation Biology



Contributed Paper

Expanding Kenya's protected areas under the Convention on Biological Diversity to maximize coverage of plant diversity

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Applications of vegtable

Suggested Applications

- Cleaning data from (messy) spreadsheet applications previous database storage
- Importing data from relational databases
- Merging data from different databases
- Documenting data assessment (R-scripts or R-markdown documents)

Recommended Functions

- `backup_object()`: Use of R-images as full backups with timestamp
- `clean()`: Get rid of orphan entries
- `count_taxa()`: Counting number of taxa at plot or cluster levels
- `cross2db()`: Formating cross tables into column oriented tables
- `crosstable()`: Writting cross tables
- `df2vegtable()`: Coercing data frames into vegtable objects
- `load_last()`: Load the latest backup written by `backup_object()`
- `match_names()`: Matching strings with a taxonomic list
- `subset()`: Produce sub-sets of vegtable objects
- `summary()`: Briefing of vegtable and taxlist objects
- `trait_stats()`: Statistics for numeric functional traits
- `trait_proportion()`: Proportion of categorical traits
- `tv2vegtable()`: Import data sets from **Turboveg 2**
- `used_concepts()`: List of taxonomic concepts applied in the records
- `used_synonyms()`: List of synonyms applied in the records
- `vegtable2kml()`: Display plot locations in **Google Earth**
- `write_juice()`: Export tables to **Juice**

Dissecting vegetable

vegetable objects

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

- **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)



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**

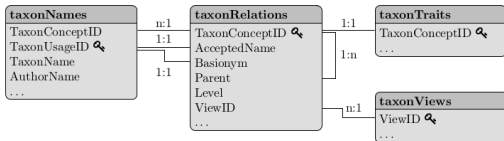
`taxlist` supports different degrees of information completeness/resolution

```
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>

Dissecting vegetable

Slot `coverconvert` (`obj@coverconvert`)

An S4 object used for conversion of abundance scale to cover percentage.

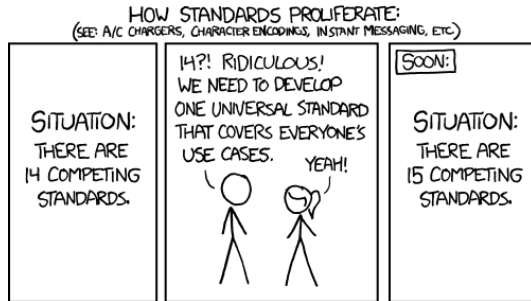
- Cover conversion tables.
- Information on class limits (bottom and top)
- Conversion by three alternative rules (bottom, mean, top)
- Only records that make sense to convert

```
names(Kenya_veg@coverconvert)
## [1] "br_bl" "b_bbd" "ordin."
summary(Kenya_veg@coverconvert$br_bl)

## ## Number of cover scales: 1
##
## * scale 'br_bl':
##   Levels      Range
## 1         r      0 - 1
## 2         +      0 - 1
## 3         1      1 - 5
## 4         2      5 - 25
## 5         3     25 - 50
## 6         4     50 - 75
## 7         5    75 - 100
```

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

THANKS!

For discussion:

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

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

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



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