

# Package ‘foodwebWrapper’

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**Version** 1.2.0

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**Title** Enhanced Wrapper to Show Which Functions Call What

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**Depends** R (>= 4.2.0)

**LazyData** true

**Imports** tools, utils, tibble, dplyr, R2HTML, textshaping, magrittr,  
tidyverse, stringr

**Description** Enhances the functionality of the mvbutils::foodweb() program. The matrix-format output of the original program contains identical row names and column names, each name representing a retrieved function. This format is enhanced by using the find\_funs() program [see Sebastian (2017) <[https://sebastiansauer.github.io/finds\\_funs/](https://sebastiansauer.github.io/finds_funs/)>] to concatenate the package name to the function name. Each package is assigned a unique color, that is used to color code the text naming the packages and the functions. This color coding is extended to the entries of value ``1" within the matrix, indicating the pattern of ancestor and descendent functions.

**License** GPL (>= 2)

**Encoding** UTF-8

**VignetteBuilder** knitr

**Suggests** knitr, rmarkdown, testthat (>= 3.0.0)

**RoxygenNote** 7.3.3

**Config/testthat/edition** 3

**NeedsCompilation** no

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---

addStyle

*addStyle*

---

### Description

insert tags into HTML code to implement rotating table text

**Usage**

```
addStyle(x, m, colorMap, pawn)
```

**Arguments**

x	character vector containing HTML code
m	character matrix containing table that is represented in x
colorMap	character array of colors
pawn	Boolean if TRUE use chess symbols rather than rectangles in html table

**Value**

returns modified HTML code

**Examples**

```
if(interactive()){  
  #load("data/x_x2.RData")  
  #load("data/x_m5.RData")  
  #load("data/x_colorMap.RData")  
  y<-addStyle(x_x2,x_m5,x_colorMap,pawn=TRUE)  
}
```

---

attachedFunctions      *attachedFunctions*

---

**Description**

print a list of attached packages and their functions for the user to select from

**Usage**

```
attachedFunctions(verbose)
```

**Arguments**

verbose	Boolean if TRUE output several user messages
---------	--

**Value**

returns a list whose components are

- 1 list of user-selected packages and corresponding functions
- where character vector of selected packages

**Examples**

```
if(interactive()){  
  attachedFunctions(verbose=TRUE)  
}
```

---

attachedFunctionsBatch  
*attachedFunctionsBatch*

---

**Description**

same as attachedFunctions() but no user interaction needed

**Usage**

```
attachedFunctionsBatch(packs)
```

**Arguments**

packs            list of character strings containing the names of packages package name is like "pack", not like "package:pack"

**Value**

returns a list whose components are

- l list of user-selected packages and corresponding functions
- where character vector of selected packages

**Examples**

```
if(interactive()){  
  attachedFunctionsBatch(c("SherlockHolmes", "textBoxPlacement"))  
}
```

---

attachedPackages	<i>attachedPackages</i>
------------------	-------------------------

---

**Description**

print a list of attached packages for the user to select from

**Usage**

```
attachedPackages()
```

**Value**

returns a character vector of selected packages

**Examples**

```
if(interactive()){
  attachedPackages()
}
```

---

colorTag	<i>colorTag</i>
----------	-----------------

---

**Description**

add tag to color function name in column 2, based on package in column 1

**Usage**

```
colorTag(v1, v2, nc, x, colorMap, pawn)
```

**Arguments**

v1	character vector first column of m (excluding first 2 entries of m)
v2	character vector second column of m (excluding first 2 entries of m)
nc	integer number of columns of m
x	return value of replaceRotTag()
colorMap	character array of colors
pawn	Boolean if TRUE use chess symbols rather than rectangles in html table

**Details**

v1 is first column of m (excluding first 2 entries of m) v2 is second column of m (excluding first 2 entries of m)

**Value**

returns

**Examples**

```
#load("data/x_m5.RData")
#load("data/x_colorMap.RData")
#load("data/x_x3.RData")
x<-colorTag(x_m5[c(-1,-2),1],x_m5[c(-1,-2),2],ncol(x_m5),x_x3,x_colorMap,pawn=TRUE)
```

---

concatPackFunc2

*concatPackFunc2*

---

**Description**

match the package names with the function names

**Usage**

```
concatPackFunc2(m, v)
```

**Arguments**

m                    character matrix return value component \$funmat of foodweb()  
v                    character vector of package names returned by find\_funz()

**Value**

returns augmented character matrix m

**Examples**

```
if(interactive()){
#load("data/x_x.RData")
#load("data/x_v.RData")
m<-concatPackFunc2(x_x$funmat,x_v)
}
```

---

consolidate	<i>consolidate</i>
-------------	--------------------

---

**Description**

create a permutation list of package names for re-ordering rows and columns of matrix m, in decreasing order of function counts per package

**Usage**

```
consolidate(v)
```

**Arguments**

v character vector of package names component of return value of `find_funz()`

**Value**

returns a list whose components are character vector for permuting order of m

**Examples**

```
if(interactive()){  
  #load("data/x_v.RData")  
  l<-consolidate(x_v)  
}
```

---

find_funz	<i>find_funz</i>
-----------	------------------

---

**Description**

determine in which R package a function ‘resides’

**Usage**

```
find_funz(packs, rfuns)
```

**Arguments**

packs list of character strings containing the names of the packages  
rfuns list of character strings containing the names of functions in packs to which the result is to be restricted

**Value**

returns vector of character strings, names are functions and values are packages

**Examples**

```
if(interactive()){
#load("data/x_packages.RData")
#load("data/x_funs.RData")
find_funz(packs=x_packages, rfuns=x_funs)
}
```

---

foodwebWrapper

*foodwebWrapper*

---

**Description**

wrapper for the function foodweb() concatenate the R package name for each retrieved R function

**Usage**

```
foodwebWrapper(
  where = character(0),
  ofile = "~/foodwebWrapper.html",
  zeros = TRUE,
  pawn = FALSE,
  verbose = TRUE
)
```

**Arguments**

where	position(s) on search path, or an environment, or a list of environments
ofile	character string containing path name for output file
zeros	Boolean if TRUE delete rows and cols that contain all 0's
pawn	Boolean if TRUE use chess symbols rather than rectangles in html table
verbose	Boolean if TRUE output several user messages

**Details**

if where is missing, then the user is presented with the option of choosing from a list of attached packages

**Value**

foodweb returns an object of (S3) class foodweb. This has three components:

- funmat a matrix of 0s and 1s showing what (row) calls what (column). The dimnames are the function names.
- x shows the x-axis location of the centre of each function's name in the display, in par("usr") units
- level shows the y-axis location of the centre of each function's name in the display, in par("usr") units. For small numbers of functions, this will be an integer; for larger numbers, there will be some adjustment around the nearest integer

**Examples**

```
if(interactive()){
#load("data/x_packages.RData")
ofile<-sprintf("%s/foodwebWrapper.html",tempdir())
foodwebWrapper(ofile=ofile)
foodwebWrapper(when=x_packages,ofile=ofile)
}
```

---

funs\_examples

*foodwebWrapper data sets*


---

**Description**

foodwebWrapper data sets

**Usage**

```
data(funs_examples)
```

---

mapFunctionsColors

*mapFunctionsColors*


---

**Description**

map functions to color coding

**Usage**

```
mapFunctionsColors(row1, col1, colors)
```

**Arguments**

row1            character vector containing names of packages  
 col1            character vector containing names of packages  
 colors          character vector containing names of colors

**Value**

returns a character vector mapping colors to package names

**Examples**

```
if(interactive()){
  colors<-c("darkmagenta","darkolivegreen","darkorange3","brown4","red","blue")
  #load("data/x_m3.RData")
  colorMap<-mapFunctionsColors(x_m3[1,c(-1,-2)],x_m3[c(-1,-2),1],colors)
}
```

---

rearrangeM

*rearrangeM*


---

**Description**

rearrange the order of rows or columns of matrix based on entries in a vector

**Usage**

```
rearrangeM(m, v2)
```

**Arguments**

m                character matrix return value of concatPackFunc2()  
 v2               list whose components are package names for permuting order of m, return value  
                   of consolidate()

**Value**

returns rearranged version of m

**Examples**

```
if(interactive()){
  #load("data/x_m.RData")
  #load("data/x_v2.RData")
  m2<-rearrangeM(x_m,x_v2)
}
```

---

removeZeroRowsCols      *removeZeroRowsCols*

---

**Description**

delete rows and cols of matrix m that contain all "0"s

**Usage**

```
removeZeroRowsCols(m)
```

**Arguments**

m                      character matrix whose entries are either "0" or "1"

**Value**

returns an altered version of character matrix m with removed rows and columns

**Examples**

```
if(interactive()){  
  #load("data/x_m2.RData")  
  m3<-removeZeroRowsCols(x_m2)  
}
```

---

replaceRotTag              *replaceRotTag*

---

**Description**

insert html tags for rotating text

**Usage**

```
replaceRotTag(x, l, dims)
```

**Arguments**

x                      return value of readLines(), HTML code containing data table  
l                      return values of spanTag()  
dims                    return value of dim()

**Value**

returns modified version of HTML code containing data table

**Examples**

```

if(interactive()){
#load("data/x_x.RData")
#load("data/x_l.RData")
#load("data/x_m3.RData")
x<-replaceRotTag(x_x,x_l,dim(x_m3))
}

```

---

rotStyle

*rotStyle*


---

**Description**

add html style definition for rotation

**Usage**

```
rotStyle()
```

**Value**

returns character string containing html style definition for rotation

**Examples**

```
r<-rotStyle()
```

---

rotTag

*rotTag*


---

**Description**

add html tag to rotate function name

**Usage**

```
rotTag(v1, v2, colorMap)
```

**Arguments**

v1	character vector containing first row of matrix m (excluding first 2 entries of m)
v2	character vector containing second row of matrix m (excluding first 2 entries of m)
colorMap	character array of colors

**Details**

see <https://stackoverflow.com/questions/47261100/how-to-rotate-text-90-degrees-inline> also need to increase height of row to accommodate rotated text see <https://resultuniversity.com/html/html-table-width-height#:~:text=To%20set%20the%20height%20of%20a%20specific%20row%20in%20an,property%20in%20pixels%20>

**Value**

returns character vector containing inserted html tags

**Examples**

```
if(interactive()){
#load("data/x_m5.RData")
#load("data/x_colorMap.RData")
rt<-rotTag(x_m5[1,c(-1,-2)],x_m5[2,c(-1,-2)],x_colorMap)
}
```

---

spanTag	<i>spanTag</i>
---------	----------------

---

**Description**

Add html tag for package name to span multiple columns. Also insert hyperlink to CRAN package and function documentation.

**Usage**

```
spanTag(v, direction = "COLSPAN", colorMap)
```

**Arguments**

v	character vector representing first row of m (excluding first 2 entries of m)
direction	character string COLSPAN or ROWSPAN
colorMap	character array of colors

**Details**

see <https://www.pierobon.org/html/span.htm#:~:text=Cells%20within%20HTML%20tables%20can,span%20more%20than>

**Value**

returns a list whose components are

- u return value of `unique(v)`
- tab return value of `table(v)`
- v2 character vector modified version of v containing html span tags

**Examples**

```

if(interactive()){
#load("data/x_m5.RData")
#load("data/x_colorMap.RData")
l<-spanTag(x_m5[1,c(-1,-2)],"COLSPAN",x_colorMap)
}

```

---

x\_colorMap

*foodwebWrapper data sets*


---

**Description**

foodwebWrapper data sets

**Usage**

```
data(x_colorMap)
```

---

x\_examples

*foodwebWrapper data sets*


---

**Description**

foodwebWrapper data sets

**Usage**

```
data(x_examples)
```

---

x\_f

*foodwebWrapper data sets*


---

**Description**

foodwebWrapper data sets

**Usage**

```
data(x_f)
```

---

x\_funs *foodwebWrapper data sets*

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_funs)

---

x\_m *foodwebWrapper data sets*

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_m)

---

x\_m2 *foodwebWrapper data sets*

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_m2)

---

x\_m3 *foodwebWrapper data sets*

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_m3)

---

x_m4	<i>foodwebWrapper data sets</i>
------	---------------------------------

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_m4)

---

x_m5	<i>foodwebWrapper data sets</i>
------	---------------------------------

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_m5)

---

x_packages	<i>foodwebWrapper data sets</i>
------------	---------------------------------

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_packages)

---

x_v	<i>foodwebWrapper data sets</i>
-----	---------------------------------

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_v)

---

x_v2	<i>foodwebWrapper data sets</i>
------	---------------------------------

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_v2)

---

x_where	<i>foodwebWrapper data sets</i>
---------	---------------------------------

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_where)

---

x_x	<i>foodwebWrapper data sets</i>
-----	---------------------------------

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_x)

---

x_x2	<i>foodwebWrapper data sets</i>
------	---------------------------------

---

**Description**

foodwebWrapper data sets

**Usage**

data(x\_x2)

---

`x_x3`*foodwebWrapper data sets*

---

**Description**

foodwebWrapper data sets

**Usage**`data(x_x3)`

---

`x_y`*foodwebWrapper data sets*

---

**Description**

foodwebWrapper data sets

**Usage**`data(x_y)`

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