# The **tables** Package\*

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# 1 Introduction

This is a short introduction to the **tables** package. Inspired by my 20 year old memories of SAS PROC TABULATE, I decided to write a simple utility to create nice looking tables in Sweave documents. For example, we might display summaries of some of Fisher's iris data using the code

```
> tabular( (Species + 1) ^{\sim} (n=1) + Format(digits=2)* + (Sepal.Length + Sepal.Width)*(mean + sd), data=iris )
```

		Sepal.Length		Sepal.Width	
Species	n	mean	sd	mean	sd
setosa	50	5.01	0.35	3.43	0.38
versicolor	50	5.94	0.52	2.77	0.31
virginica	50	6.59	0.64	2.97	0.32
All	150	5.84	0.83	3.06	0.44

You can also pass the output through the Hmisc::latex() function (Harrell, 2011, Harrell et~al., 2011) to produce LATEX output, which when processed by pdflatex will produce the following table:

		Sepal.I	Length	Sepal.	Width
Species	n	mean	$\operatorname{sd}$	mean	$\operatorname{sd}$
setosa	50	5.01	0.35	3.43	0.38
versicolor	50	5.94	0.52	2.77	0.31
virginica	50	6.59	0.64	2.97	0.32
All	150	5.84	0.83	3.06	0.44

If you prefer the style of table that the LaTeX **booktabs** package (Fear, 2005) produces, you can choose that style instead. I mostly like it, so I have used

#### > booktabs()

for the rest of this document. This gives

		Sepal.I	Length	Sepal.V	Width
Species	$\mathbf{n}$	mean	$\operatorname{sd}$	mean	$\operatorname{sd}$
setosa versicolor virginica All	50 50 50 150	5.01 5.94 6.59 5.84	0.35 0.52 0.64 0.83	3.43 2.77 2.97 3.06	0.38 0.31 0.32 0.44

Details on booktabs() are given in section 2.1.5 below.

There is also the html.tabular method for the Hmisc::html() generic; it produces output in HTML format. Finally, see section 2.1.3 for other output formats.

The idea of a table in the **tables** package is a rectangular array of values, with each row and column labelled, and possibly with groups of rows and groups of columns also labelled. These arrays are specified by "table formulas".

Table formulas are R formula objects, with the rows of the table described before the tilde ("~"), and the columns after. Each of those is an expression containing "\*", "+", "=", as well as functions, function calls and variables, and parentheses for grouping. There are also various directives included in the formula, entered as "pseudo-functions", i.e. expressions that look like function calls but which are interpreted by the tabular() function.

For example, in the formula

the rows are given by (Species + 1). The summation here is interpreted as concatenation, i.e. this says rows for Species should be followed by rows for 1.

In the iris dataframe, Species is a factor, so the rows for it correspond to its levels.

The 1 is a place-holder, which in this context will mean "all groups". The columns in the table are defined by

```
(n=1) + Format(digits=2)*(Sepal.Length + Sepal.Width)*(mean + sd)
```

Again, summation corresponds to concatenation, so the first column corresponds to (n=1). This is another use of the placeholder, but this time it is labelled as n. Since we haven't specified any other statistic to use, the first column contains the counts of values in the dataframe in each category.

The second term in the column formula is a product of three factors. The first, Format(digits=2), is a pseudo-function to set the format for all of the entries to come. (For more on formats, see section 2.4.1 below.) The second factor, (Sepal.Length + Sepal.Width), is a concatenation of two variables. Both of these variables are numeric vectors in iris, and they each become the variable to be analyzed, in turn. The last factor, (mean + sd) names two R functions. These are assumed to be functions that operate on a vector and produce a single value, as mean and sd do. The values in the table will be the results of applying those functions to the two different variables and the subsets of the dataset.

# 2 Reference

For the examples below we use the following definitions:

```
> set.seed(100)
> X <- rnorm(10)
> X

[1] -0.50219235  0.13153117 -0.07891709  0.88678481
[5]  0.11697127  0.31863009 -0.58179068  0.71453271
[9] -0.82525943 -0.35986213

> A <- sample(letters[1:2], 10, rep=TRUE)
> A
```

```
[1] "b" "b" "b" "a" "a" "a" "b" "b" "a"
> F <- factor(A)
> F

[1] b b b b a a b b b a
Levels: a b

2.1 Function syntax
2.1.1 tabular()
tabular(table, ...)
```

The tabular function is a generic function. The default method uses as.formula() to try to convert the table argument to a formula, then passes it and all the other arguments to tabular.formula() method, which does

the first two are used, and a warning is issued if anything is passed in the ... arguments.

tabular.default(table, ...)

table The table argument is the table formula, described in detail below.

most of the work. That method has 4 arguments plus ..., but usually only

- data The data argument is a dataframe or environment in which to look for the data referenced by the table.
- n The tabular function needs to know the length of vectors on which it operates, because some formulas (e.g. 1 ~ 1) contain no data. Normally n is taken as the number of rows in data, or the length of the first referenced object in the formula, but sometimes the user will need to specify it. Once specified, it can't be modified: all data in the table should be the same length.
- suppressLabels By default, tabular adds a row or column label for each term, but this does sometimes make the table messy. Setting suppressLabels to a positive integer will cause that many labels to be suppressed at the start of each term. The pseudo-function Heading() can achieve the same effect, one term at a time.

The value returned is a list-mode matrix corresponding to the entries in the table, with a number of attributes to help with formatting. See the ?tabular help page for more details.

```
2.1.2 format(), print(), latex()
format(x, digits=4, justification="n", ...)
print(x, ...)
latex(x, file="", options=NULL, ...)
```

The tables package provides methods for the format(), print() and Hmisc::latex() generics. The arguments are:

x The tabular object returned from tabular().

digits The default number of digits to use when formatting.

- justification The default text justification to use when printing. For text display, the recognized values are "n", "l", "c", "r", standing for none, left, center and right justification respectively. For LATEX the justification is specified via the table\_options() function (section 2.1.5).
- file The default method for the Hmisc::latex() generic writes the LATEX code to a file; latex.tabular() can optionally do the same, but it defaults to writing to screen, for use in Sweave documents like this one.
- options A list of options to pass to table\_options(). These will be set only for the duration of the call to latex().

```
2.1.3 as.matrix(), write.csv.tabular(), write.table.tabular()
as.matrix(x, format = TRUE,
    rowLabels = TRUE, colLabels = TRUE, justification = "n", ...)
write.csv.tabular(x, file = "", justification = "n", row.names=FALSE,
    write.options=list(), ...)
write.table.tabular(x, file="",
    justification = "n", row.names=FALSE, col.names=FALSE,
    write.options=list(), ...)
```

These functions export tables for further computations. The arguments are:

**x** The tabular object.

format Whether to format the entries. See the help page for alternatives.

rowLabels, colLabels If formatting, whether to include the labels or not.

**justification** The default text justification to use when formatting.

file Where to write the output.

row.names, col.names, write.options Additional parameters to pass to write.csv() or write.table().

```
2.1.4 as.tabular()
as.tabular(x, ...)
as.tabular.default(x, like=NULL, ...)
as.tabular.data.frame(x, ...)
```

These functions create tables from existing matrices or dataframes of values. The dimnames of the input are used to construct default row and column names. If more elaborate labelling is wanted, use a tabular object as the like argument. The labelling for like will be used on the newly constructed result.

#### 2.1.5 table\_options(), booktabs()

The table\_options() function sets a number of formatting defaults for the latex() method:

justification This is the default justification for data columns and their headers. Any justification string will be accepted; it should be one that the LaTeX \tabular environment (or substitute) accepts. If a vector of strings is specified they will be recycled across the columns of the table.

rowlabeljustification This is the default justification for row labels. A vector of strings will be recycled across the row label columns.

tabular The environment to use in LaTeX. Alternatives to "tabular" such as "longtable" can be used here. Those often also need modifications within the table; the Literal() (section 2.5.3) function may be helpful.

toprule, midrule, bottomrule The LATEX macros to draw the top, middle and bottom lines in the table. By default these are all "\\hline".

titlerule An optional LATEX macro to draw a line under multicolumn titles.

doBegin, doHeader, doBody, doFooter, doEnd These logical values control the inclusion of specific parts of the output table.

```
$justification
[1] "c"
$rowlabeljustification
[1] "1"
$tabular
[1] "tabular"
$toprule
[1] "\\hline"
$midrule
[1] "\\hline"
$bottomrule
[1] "\\hline"
$titlerule
NULL
$doBegin
[1] TRUE
$doHeader
[1] TRUE
```

\$doBody
[1] TRUE

The defaults are

```
$doFooter
[1] TRUE
$doEnd
[1] TRUE
$latexleftpad
[1] TRUE
$latexrightpad
[1] TRUE
$latexminus
[1] TRUE
$doHTMLheader
[1] FALSE
$doCSS
[1] FALSE
$doHTMLbody
[1] FALSE
$CSS
[1] "<style>\n#ID .left { text-align:left; }\n#ID .center { text-align:center; }
$HTMLhead
[1] "<!DOCTYPE html>\n<html>\n<head>\n<meta charset=\"CHARSET\">\n"
$HTMLbody
[1] "<body>\n"
$HTMLattributes
[1] "frame=\"hsides\" rules=\"groups\""
$HTMLcaption
```

```
NULL
$HTMLfooter
NULL
$HTMLleftpad
[1] FALSE
$HTMLrightpad
[1] FALSE
$HTMLminus
[1] FALSE
```

Some options only apply to HTML output; see the help page ?table\_options for details.

If you are using the LATEX **booktabs** package, the booktabs() function will set different options. Currently those are:

		Sepal.I	Length	Sepal.	Width
Species	n	mean	$\operatorname{sd}$	mean	$\operatorname{sd}$
setosa	50	5.01	0.35	3.43	0.38
versicolor	50	5.94	0.52	2.77	0.31
virginica	50	6.59	0.64	2.97	0.32
All	150	5.84	0.83	3.06	0.44

We can use the doXXXX options to insert raw LATEX into a table:

		Sepal.I	Length	Sepal.V	Width
Species	$\mathbf{n}$	mean	$\operatorname{sd}$	mean	$\operatorname{sd}$
setosa versicolor virginica Overall, we	50 50 50 e see	5.01 5.94 6.59 the follo	0.35 0.52 0.64 wing:	3.43 2.77 2.97	0.38 0.31 0.32
All	150	5.84	0.83	3.06	0.44

#### 2.1.6 latexNumeric()

The latexNumeric() function converts character representations of numbers into a format suitable for display in LATEX documents. There are two goals:

- If chars is a vector with constant width, then the output will also be constant width. This means the default centering used in tabular() will not misalign decimal points (if they were aligned in chars).
- Minus signs will be displayed with the proper symbol rather than a hyphen.

The arguments are:

**chars** A character vector of formatted numeric values.

minus Whether to pad positive cases with spacing of the same width as a minus sign. If TRUE and some entries are negative, then all positive entries will be padded.

leftpad, rightpad Whether to pad cases that have leading or trailing blanks with spacing matching a digit width per space. If leftpad=TRUE, leading blanks will be converted to spaces the same width as a digit 0. (If minus=TRUE, one leading blank may have been consumed in the sign padding.) The rightpad argument handles trailing blanks similarly.

mathmode Whether to wrap the result in dollar signs, so LATEX will render minus signs properly.

# 2.2 Operators

#### **2.2.1** $e_1 + e_2$

Summing two expressions indicates that they should be displayed in sequence. For rows, this means  $e_1$  will be displayed just above  $e_2$ ; for columns,  $e_1$  will be just to the left of  $e_2$ .

Example:

> latex( tabular(F + 1 ~ 1) )

F	All
a	3
b	7
All	10

### **2.2.2** $e_1 * e_2$

Multiplying two expressions means that each element of  $e_1$  will be applied to each element of  $e_2$ . If  $e_1$  is a factor, then  $e_2$  will be displayed for each element of it. NB: \* has higher precedence than + and evaluation proceeds from left to right. The expression  $(e_1 + e_2) * (e_3 + e_4)$  is equivalent to  $e_1 * e_3 + e_1 * e_4 + e_2 * e_3 + e_2 * e_4$ .

Example:

## > latex( tabular( X\*F\*(mean + sd) ~ 1 ) )

F		All
X a b	mean sd mean sd	0.02525 $0.34842$ $-0.03647$ $0.65611$

## **2.2.3** $e_1 \sim e_2$

The tilde separates row specifications from column specifications, but otherwise acts the same as \*, i.e. each row value applies to each column.

Example:

## > latex( tabular( X\*F ~ mean + sd ) )

	F	mean	sd
X	a b	0.02525 $-0.03647$	0.0101

#### **2.2.4** $e_1 = e_2$

The operator = is used to set the name of  $e_2$  to a displayed version of  $e_1$ . It is an abbreviation for  $\texttt{Heading}(e_1)*e_2$ . NB: because = has lower operator precedence than any other operator, we usually put parentheses around these expressions, i.e.  $(e_1 = e_2)$ .

Example: F is renamed to "Newname".

	Newname	mean	sd
X	a	0.02525	0.3484
	b	-0.03647	0.6561

## 2.3 Terms in Formulas

R parses table formulas into sums, products, and bindings separated by the tilde formula operator. What comes between the operators are other expressions. Other than the pseudo-functions described in section 2.4, these are evaluated and the actions depend on the type of the resulting value.

#### 2.3.1 Closures or other functions

If the expression evaluates to a function (e.g. it is the name of a function), then that function becomes the summary statistic to be displayed. The summary statistic should take a vector of values as input, and return a single value (either numeric, character, or some other simple printable value). If no summary function is specified, the default is length, to count the length of the vector being passed.

Note that only one summary function can be specified for any cell in the table or an error will be reported.

Example: mean and sd are specified functions; n is the renamed default statistic.

		X	X		
F	n	mean	$\operatorname{sd}$		
a	3	0.02525	0.3484		
b	7	-0.03647	0.6561		
All	10	-0.01796	0.5611		

#### 2.3.2 Factors

If the expression evaluates to a factor, the dataset is broken up into subgroups according to the levels of the factor. Most of the examples above have shown this for the factor F, but this can also be used to display complete datasets:

Example: creating a factor to show all data. Use the identity function to display the values in each cell.

i	X	A	F
1	-0.50219	b	b
2	0.13153	b	b
3	-0.07892	b	b
4	0.88678	b	b
5	0.11697	a	a
6	0.31863	a	a
7	-0.58179	b	b
8	0.71453	b	b
9	-0.82526	b	b
10	-0.35986	a	a

### 2.3.3 Logical vectors

If the expression evaluates to a logical vector, it is used to subset the data. Example: creating subsets on the fly.

```
> latex( tabular( (X > 0) + (X < 0) + 1
+ ((n = 1) + X*(mean + sd)) ) )
```

		X	
	n	mean	$\operatorname{sd}$
X > 0	5	0.43369	0.3496
X < 0	5	-0.46960	0.2761
All	10	-0.01796	0.5611

#### 2.3.4 Language Expressions

If the expression evaluates to a language object, e.g. the result of quote() or substitute(), then it will be replaced in the table formula by its result. This allows complicated table formulas to be saved and re-used. For examples, see section 2.5.

#### 2.3.5 Other vectors

If the expression evaluates to something other than the above, then it is assumed to be a vector of values to be summarized in the table. If you would like to summarize a factor or logical vector, wrap it in I() to prevent special handling.

Note that the following must all be true, or an error will be reported:

- only one value vector can be specified for any cell in the table,
- all value vectors must be the same length,
- is.atomic() must evaluate to TRUE for the vector.

Example: treating a logical vector as values.

> latex( tabular( 
$$I(X > 0) + I(X < 0)$$
  
+  $((n=1) + mean + sd)$  )

	n	mean	$\operatorname{sd}$
I(X > 0)	10	0.5	0.527
I(X < 0)	10	0.5	0.527

### 2.4 "Pseudo-functions"

Several directives to **tables** may be embedded in the table formula. This is done using "pseudo-functions". Syntactically they look like function calls, but reserved names are used. In most cases, their action applies to later factors in the term in which they appear. For example,

will apply the Justify(r) directive to both Y and Z, but the Format(digits=2) directive will only apply to Z, and neither will apply to A.

#### 2.4.1 Format()

By default **tables** formats each column using the standard format() function, with arguments taken from the format.tabular() call (see section 2.1.2).

The Format() pseudo-function does two things: it changes the formatting, and it specifies that all values it applies to will be formatted together. The "call" to Format looks like a call to format, but without specifying the argument x. When tabular() formats the output it will construct x from the entries in the table governed by the Format() specification.

Example: The mean and standard deviation are both governed by the same format, so they are displayed with the same number of decimal places, chosen so that the smallest values (the means) show two significant digits.

```
> latex( tabular( (F+1) ~ (n=1)
+ Format(digits=2)*X*(mean + sd) ) )
```

		X	
F	n	mean	$\operatorname{sd}$
a	3	0.025	0.348
b	7	-0.036	0.656
All	10	-0.018	0.561

For customized formatting, an alternate syntax is to pass a function call to Format(), rather than a list of arguments. The function should accept an argument named x (but as with the regular formatting, x should not be included in the formula), to contain the data. It should return a character vector of the same length as x.

Example: Use a custom function and sprintf() to display a standard error in parentheses.

```
> stderr <- function(x) sd(x)/sqrt(length(x))
> fmt <- function(x, digits, ...) {
+    s <- format(x, digits=digits, ...)
+    is_stderr <- (1:length(s)) > length(s) %/% 2
+    s[is_stderr] <- sprintf("$(%s)$", s[is_stderr])
+    s[!is_stderr] <- latexNumeric(s[!is_stderr])
+    s
+ }
> latex( tabular( Format(fmt(digits=1))*(F+1) ~ X*(mean + stderr) ) )
```

		X				
F	mean	stderr				
a	0.03	(0.20)				
b	-0.04	(0.25)				
All	-0.02	(0.18)				

Character values in cells in the table are handled specially; see section 3.1 below.

#### 2.4.2 .Format()

The pseudo-function .Format() is mainly intended for internal use. It takes a single integer argument, saying that data governed by this call uses the same formatting as the format specification indicated by the integer. In this way entries can be commonly formatted even when they are not contiguous. The integers are assigned sequentially as the format specification is parsed; users will likely need trial and error to find the right value in a complicated table with multiple formats.

Example: Format two separated columns with the same format.

		X	
F	mean	n	$\operatorname{sd}$
a	0.025	3	0.348
b	-0.036	7	0.656
All	-0.018	10	0.561

#### 2.4.3 Heading()

Normally tabular() generates row and column labels by deparsing the expression being tabulated. These can be changed by using the Heading() pseudo-function, which replaces the heading on the next object found. The heading can either be the name of a function or a string in quotes, which will be displayed as entered (so LATEX codes can be used).

If no argument is passed, the next label is suppressed.

There's an optional second argument, which must be either TRUE or FALSE if present. If it is TRUE (or not present), then the heading will override a previously specified heading. If FALSE, it will not. The latter seems likely only to be of use in automatically generated code, and is used in the automatically generated labels for factors.

Example: Replace F with a Greek  $\Phi$ , and suppress the label for X.

```
> latex( tabular( (Heading("$\\Phi$")*F+1) ~ (n=1)
+ Format(digits=2)*Heading()*X*(mean + sd) ) )
```

Φ	n	mean	sd
a	3	0.025	0.348
b	7	-0.036	0.656
All	10	-0.018	0.561

### 2.4.4 Justify()

The Justify() pseudo-function is used to specify the text justification of the headers and data values in the table. If called with one argument, that value is used for both labels and data; if called with two arguments, the first is used for the labels, the second for the data. If no Justify() specification is given, the default passed to format(), print() or latex() will be used. Values may be specified without quotes if they are legal R names; quoted strings may also be used. (The latter is useful for LATEX output, for example Justify("r@{}"), to suppress column spacing on the right.)

Example:

		X	-
F	n	mean	$\operatorname{sd}$
a	3	0.025	0.348
b	7	-0.036	0.656
All	10	-0.018	0.561

#### 2.4.5 Percent()

The Percent() pseudo-function is used to specify a statistic that depends on other values in the table. It has two optional arguments:

denom="all" This specifies how the denominator (argument y to fn below) is set. The most commonly used values are "all", meaning all values are used, "row", meaning only the values in the current row are used, "col", meaning only the values in the current column are used.

The special syntax Equal(...) will record the expressions in ..., and ignore any factor based subsetting if the factor does not appear among the expressions. Similarly Unequal(...) will use values which differ in any of the expressions in ... from the values in the current cell.<sup>1</sup>

If a logical vector is given, it is used to select which values form the denominator. Anything else is just passed to fn as given.

fn=percent This is the function which actually does the computation. The
 default definition is function(x, y) 100\*length(x) /length(y), giv ing the percentage count, but any other two argument function could
 be used.

These two examples are different ways of producing the same table:

<sup>&</sup>lt;sup>1</sup> In fact, the mechanism is more general. The expressions in Equal(...) or Unequal(...) are departed and treated as strings. Any logical vector elsewhere in the table may be labelled with a string using the labelSubset function and those labels will be respected. Unlabelled logical vectors in the table formula will always be used for subsetting.

		Carburetors						
Gears		1	2	3	4	6	8	All
3	n	3	4	3	5	0	0	15
	Percent	9	12	9	16	0	0	47
	RowPct	20	27	20	33	0	0	100
	ColPct	43	40	100	50	0	0	47
4	n	4	4	0	4	0	0	12
	Percent	12	12	0	12	0	0	38
	RowPct	33	33	0	33	0	0	100
	ColPct	57	40	0	40	0	0	38
5	n	0	2	0	1	1	1	5
	Percent	0	6	0	3	3	3	16
	RowPct	0	40	0	20	20	20	100
	ColPct	0	20	0	10	100	100	16
All	n	7	10	3	10	1	1	32
	Percent	22	31	9	31	3	3	100
	RowPct	22	31	9	31	3	3	100
	ColPct	100	100	100	100	100	100	100

#### 2.4.6 Arguments()

The Arguments() pseudo-function is an exception to the rule that pseudo-functions apply to later factors in the table. What it does is to specify (additional) arguments to the summary function (see section 2.3.1). For example, the weighted.mean() function takes two arguments: x and w. To use it in a table, you would specify the values to use as x via the usual mechanism for the analysis variable (section 2.3.5), and include a term Arguments(w=weights) either before or after it. The function will be called as weighted.mean(x[subset], w=weights[subset]), where subset is a logical vector indicating which rows of data belong in the current cell.

It is actually a little more complicated than as described above. The arguments to Arguments are evaluated in full, then only those which are length n are subsetted. And if no analysis variable has been specified, but Arguments() has been, then the function will be called without the x[subset] argument. Finally, the Arguments() entry will not create a heading.

For example:

	weighted.mean
gp	X
1	3.500
2	3.360
All	3.453

The same table (without the x heading) can be produced using

The order of the weighted.mean and Arguments() factors makes no difference.

#### 2.5 Formula Functions

Currently several examples of formula functions are provided. Not all are particularly robust; e.g. Hline() only works for LATEX output and must be in a particular position in the formula. Users can provide their own as well. Such functions should return a language object, which will be substituted into the formula in place of the formula function call.

#### 2.5.1 All()

This function expands all the columns from a dataframe into separate variables in the table. It has syntax

The arguments are

df A dataframe or matrix whose columns are to be displayed

numeric, character, logical, factor, complex and raw Whether to include columns of the corresponding types in the table.

other Whether to include columns that match none of the previous types.

texify Whether to escape LaTeX special characters. See section 3.1.

If functions are given for any of the selection arguments, the columns will be transformed according to the specified function before inclusion. For example, using factor=as.character will convert factors into character vectors in the table.

Example: Show the means of the numeric columns in the iris data.

#### > latex( tabular( Species ~ Heading()\*mean\*All(iris), data=iris) )

Species	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
setosa	5.006	3.428	1.462	0.246
versicolor	5.936	2.770	4.260	1.326
virginica	6.588	2.974	5.552	2.026

### 2.5.2 Hline()

This function produces horizontal lines in the table. It only works for LaTeX output, and must be the first factor in a term in the table formula. It has syntax

#### Hline(columns)

The argument is

columns An optional vector listing which columns should get the line.

Example:

Species	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
setosa versicolor virginica	5.006 5.936 6.588	3.428 2.770 2.974	1.462 4.260 5.552	0.246 1.326 2.026
All	5.843	3.057	3.758	1.199

### 2.5.3 Literal()

This function inserts literal text as a label. It has syntax

#### Literal(x)

The single argument is the text to insert. It is used by the Hline() function to insert the text.

#### 2.5.4 PlusMinus()

This function produces table entries like  $x \pm y$  with an optional header. It has syntax

PlusMinus(x, y, head, xhead, yhead, digits=2, ...)

The arguments are

x, y These are expressions which should each generate a single column in the table. The x value will be flush right, the y value will be flush left, with the  $\pm$  symbol between.

head If not missing, this header will be put over the pair of columns.

xhead, yhead If not missing, these will be put over the individual columns.

digits, ... These arguments will be passed to the standard format() function.

Example: Display mean  $\pm$  standard error.

```
> stderr <- function(x) sd(x)/sqrt(length(x))
> latex( tabular( (Species+1) ~ All(iris)*
+ PlusMinus(mean, stderr, digits=1), data=iris ) )
```

Species	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
setosa	$5.01 \pm 0.05$	$3.43 \pm 0.05$	$1.46 \pm 0.02$	$0.25 \pm 0.01$
versicolor	$5.94 \pm 0.07$	$2.77 \pm 0.04$	$4.26 \pm 0.07$	$1.33 \pm 0.03$
virginica	$6.59 \pm 0.09$	$2.97 \pm 0.05$	$5.55 \pm 0.08$	$2.03 \pm 0.04$
All	$5.84 \pm 0.07$	$3.06 \pm 0.04$	$3.76 \pm 0.14$	$1.20 \pm 0.06$

#### 2.5.5 Paste()

This function produces table entries made up of multiple values. It has syntax

The arguments are

... Expressions to be displayed in the columns of the table.

head If not missing, this will be used as a column heading for the combined columns.

digits Digits used in formatting. If a single value is given, all columns will be formatted in common. If multiple values are given, each is formatted separately.

justify One or more justifications to use on the individual columns.

prefix, sep, postfix Text to use before, between, and after the columns.

Example: Display a confidence interval.

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
Species	95% CI	95% CI	95% CI	95% CI
setosa versicolor virginica All	[5.79, 6.08] [6.41, 6.77]	[2.68, 2.86] [2.88, 3.07]	[1.41, 1.51] [4.13, 4.39] [5.40, 5.71] [3.47, 4.04]	[1.27, 1.38] [1.95, 2.10]

#### 2.5.6 Factor(), RowFactor() and Multicolumn()

The Factor() function converts its argument into a factor, but keeps the original name for a column heading. RowFactor() is designed to be used only for LATEX output: it produces multiple rows the way a factor does, but with more flexibility in the formatting. The Multicolumn() function is also designed for LATEX output: it displays factor levels in the style where the level is displayed across multiple columns on its own line.

They have syntax

The arguments are

**x** A variable to be treated as a factor.

**name** The name to be used for the factor; by default, the name passed as x.

**levelnames** An optional argument to allow customization of the displayed level names.

texify Whether to escape LATEX special characters. See section 3.1.

spacing Extra spacing is added before every group of spacing lines.

**space** How much extra space to add (in "ex" units).

**nopagebreak** Macro to insert to suppress page breaks except between groups.

width How many columns for the label?

first What is the first column?

justify What justification to use.

Example: Show the first 15 lines of the iris dataset, in groups of 5 lines.

- > subset <- 1:15
- > latex( tabular( RowFactor(subset, "\$i\$", spacing=5)
- + All(iris[subset,], factor=as.character)\*Heading()\*identity ) )

i	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa
7	4.6	3.4	1.4	0.3	setosa
8	5.0	3.4	1.5	0.2	setosa
9	4.4	2.9	1.4	0.2	setosa
10	4.9	3.1	1.5	0.1	setosa
11	5.4	3.7	1.5	0.2	setosa
12	4.8	3.4	1.6	0.2	setosa
13	4.8	3.0	1.4	0.1	setosa
14	4.3	3.0	1.1	0.1	setosa
15	5.8	4.0	1.2	0.2	setosa

To add extra space after each high level group in a multi-way classification, use spacing = 1. For example:

```
> dat <- expand.grid(Block=1:3, Treatment=LETTERS[1:2],</pre>
```

<sup>+</sup> Subset=letters[1:2])

<sup>&</sup>gt; dat\$Response <- rnorm(12)</pre>

<sup>&</sup>gt; latex( tabular( RowFactor(Block, spacing=1)

<sup>\*</sup> RowFactor(Treatment, spacing=1, space=0.5)

Block	Treatment	Subset	Response
1	A	a	-0.02932
		b	0.76406
	В	a	-0.91381
		b	-0.81438
2	A	0	n 2000E
2	А	a b	-0.38885 $0.26196$
	_	D	
	В	a	2.31030
		b	-0.43845
3	A	a	0.51086
9		b	0.77340
	В	a	-0.43809
	D	a b	-0.43809 $-0.72022$
			0.12022

For longer tables, the "longtable" environment allows the table to cross page boundaries. Using this is more complicated, as in the example below. The toprule setting inserts the caption as well as the top rule, because the longtable package requires it to be within the table. The midrule setting gets the headings to repeat on subsequent pages.<sup>2</sup> To avoid extra spacing at the top of those pages, we need to undo the automatic addition of a \normalbaselineskip there, and use suppressfirst=FALSE so that the first page doesn't get messed up. Whew!

<sup>&</sup>lt;sup>2</sup>I've done all of this in a way that is compatible with the **booktabs** style; if you want the default style, use \hline in place of the **booktabs** \toprule and \midrule macros in the options settings instead.

```
+ options = list(tabular="longtable",
+ toprule="\\caption{This table crosses page boundaries.}\\\
+ \\toprule",
+ midrule="\\midrule\\\\[-2\\normalbaselineskip]\\endhead\\hline\\endfoot") )
```

Table 1: This table crosses page boundaries.

i	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa
7	4.6	3.4	1.4	0.3	setosa
8	5.0	3.4	1.5	0.2	setosa
9	4.4	2.9	1.4	0.2	setosa
10	4.9	3.1	1.5	0.1	setosa
11	5.4	3.7	1.5	0.2	setosa
12	4.8	3.4	1.6	0.2	setosa
13	4.8	3.0	1.4	0.1	setosa
14	4.3	3.0	1.1	0.1	setosa
15	5.8	4.0	1.2	0.2	setosa
16	5.7	4.4	1.5	0.4	setosa
17	5.4	3.9	1.3	0.4	setosa
18	5.1	3.5	1.4	0.3	setosa
19	5.7	3.8	1.7	0.3	setosa
20	5.1	3.8	1.5	0.3	setosa
21	5.4	3.4	1.7	0.2	setosa
22	5.1	3.7	1.5	0.4	setosa
23	4.6	3.6	1.0	0.2	setosa
24	5.1	3.3	1.7	0.5	setosa
25	4.8	3.4	1.9	0.2	setosa

Table 1: This table crosses page boundaries.

$\overline{i}$	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
26	5.0	3.0	1.6	0.2	setosa
27	5.0	3.4	1.6	0.4	setosa
28	5.2	3.5	1.5	0.2	setosa
29	5.2	3.4	1.4	0.2	setosa
30	4.7	3.2	1.6	0.2	setosa
31	4.8	3.1	1.6	0.2	setosa
32	5.4	3.4	1.5	0.4	setosa
33	5.2	4.1	1.5	0.1	setosa
34	5.5	4.2	1.4	0.2	setosa
35	4.9	3.1	1.5	0.2	setosa
36	5.0	3.2	1.2	0.2	setosa
37	5.5	3.5	1.3	$0.2 \\ 0.2$	setosa
38	4.9	3.6	1.3 $1.4$	0.2	setosa
39	4.4	3.0	1.3	$0.1 \\ 0.2$	setosa
40	5.1	3.4	1.5	$0.2 \\ 0.2$	setosa
10	0.1	5.1	1.0	0.2	ветова
41	5.0	3.5	1.3	0.3	setosa
42	4.5	2.3	1.3	0.3	setosa
43	4.4	3.2	1.3	0.2	setosa
44	5.0	3.5	1.6	0.6	setosa
45	5.1	3.8	1.9	0.4	setosa
46	4.8	3.0	1.4	0.3	setosa
47	5.1	3.8	1.6	0.2	setosa
48	4.6	3.2	1.4	$0.2 \\ 0.2$	setosa
49	5.3	$\frac{3.2}{3.7}$	1.5	$0.2 \\ 0.2$	setosa
50	5.0	3.3	1.4	0.2	setosa

To suppress the row numbering, use suppress=3 in the call to tabular. (It is 3 because we need to suppress the column heading, the rewritten labels for the rows, and the original labels. Trial and error is the best way to determine this!) Unfortunately, the spacing features of RowFactor() won't

work without the row labels.

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa
4.6	3.4	1.4	0.3	setosa
5.0	3.4	1.5	0.2	setosa
4.4	2.9	1.4	0.2	setosa
4.9	3.1	1.5	0.1	setosa

(It is actually possible to get this to work with RowFactor(), but it is ugly: set the name and level names to "", and set the justification to "l@{}" to suppress the intercolumn spacing. Then the column of row labels will be there, but it will be zero width and invisible.)

RowFactor with spacing > 1 will add the nopagebreak macro at the beginning of each label except the first in the group. This can produce LATEX errors in any column except the first one. One workaround for this is to post-process the table to move the macro. For example, if tab contains the result of tabular() and LATEX complains about misplaced \nopagebreak macros, this will allow it to be displayed properly:

```
> code <- capture.output( latex( tab ) )
> code <- sub("^(.*)(\\\nopagebreak )", "\\2\\1", code)
> cat(code, sep = "\n")
```

To get group labels to span multiple columns, the levelnames argument can be used with embedded LATEX code. For example,

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width			
Iris setos	Iris setosa						
mean	5.0060	3.4280	1.4620	0.2460			
$\operatorname{sd}$	0.3525	0.3791	0.1737	0.1054			
Iris vers	icolor						
mean	5.9360	2.7700	4.2600	1.3260			
$\operatorname{sd}$	0.5162	0.3138	0.4699	0.1978			
Iris virgi	inica						
mean	6.5880	2.9740	5.5520	2.0260			
$\operatorname{sd}$	0.6359	0.3225	0.5519	0.2747			

# 3 Further Details

# 3.1 Formatting

As mentioned in 2.4.1, formatting in **tables** depends on the standard **format()** function or other user-selected functions. Here are the details of how it is done.

The format.tabular() method does the first part of the work. First, it constructs the calls to the appropriate formatting functions, and uses them to format all of the non-character entries in the table. The character entries are left as-is, except as described below. This converts the tabular object to a character array.

The procedure goes as follows:

- 1. Entries in the table without specified formatting are formatted first, separately by column using the format() function. This is so that entries in a given column will end up with the same character width and (with the default settings) with the same number of decimal places.
- 2. Entries in the table with specified formatting are grouped according to the format specification. For example, if two columns both share the same Format(), they will be formatted in a single call. This results in such entries ending up with the same character width and (with the default settings) with the same number of decimal places.
- 3. If the latex argument is TRUE, any numeric entries are passed to the latexNumeric() function (see 2.1.6), which replaces blanks and mi-

nus signs with fixed width spaces and LATEX minus signs so that all entries will display in the same width. This means that numeric values will normally have decimal points aligned, unless the formatting function explicitly removes leading spaces. Non-numeric entries are passed through the Hmisc::latexTranslate function so that special characters are displayed properly.

4. If the latex argument is FALSE, an attempt is made to justify the results using simple ASCII spacing, according to the Justify() specification with the justification argument used as a default.

Note that LaTeX special characters will be escaped in data when latex() is called, but row and column headings generated by All(), Factor(), etc. will by default have the escapes done in all cases. Those functions have a texify argument that can be set to FALSE to disable this behaviour (e.g. if the label is meant to be processed by LaTeX). For example, with the definition

- >  $df \leftarrow data.frame(A = factor(c( "$", "\\" ) ), B_label=1:2)$  the code
- > latex( tabular( mean ~ A\*B\_label, data=df ) )

would fail, as the labels would include the special characters. But this will work:

> latex( tabular( mean ~ Factor(A)\*All(df), data=df ) )

	A		
	\$	\	
	$B_{-label}$	$B_{-label}$	
mean	1	2	

As mentioned above, character values in cells in the table are handled specially. If the default format function (or a custom function named format) is used, then those character values are not formatted, they are just copied into the result. (This is so that a column can have mixed numeric and character values, and the numerics are not converted to character before formatting.) If you want to use format on character values, you will need to use a custom formatting function with a different name.

## 3.2 Missing Values

By default, most summary statistics in R return NA if any of the input values are NA, but have ways to treat NA differently. For example, the mean() function has the na.rm argument:

```
> dat <- data.frame( a = c(1, 2, 3, NA), b = 1:4 )
> mean(dat$a)
[1] NA
> mean(dat$a, na.rm=TRUE)
[1] 2
```

The tabular() function itself has no way to specify special NA handling, but there are several ways to do this yourself, depending on how you want them handled. To ignore NA values within the column, define a new function which sets the different behaviour. For example,

An alternative approach is to use na.omit() to work on a subset of your data which has rows with any missing values removed, e.g.

> latex( tabular( mean ~ a + b, data = na.omit(dat) ) ) 
$$\frac{a \quad b}{mean \quad 2 \quad 2}$$

A third possibility is to use the complete.cases() function to remove missings only from some columns, e.g.

	All		Cor	nplete
	a	b	a	b
Mean	2	2.5	2	2

Missing values in factors are normally ignored, i.e. observations whose value is missing won't match any category. If you would like NA to be used as an additional category, use exclude = NULL in a call to factor() when you create the variable, e.g. compare the following two tables:

A	n
1	1
2	1
3	1
All	4

```
> A <- factor(dat$a, exclude = NULL)
> latex( tabular( A + 1 ~ (n=1) ) )
```

A	n
1	1
2	1
3	1
NA	1
All	4

# 3.3 Subsetting and Joining Tables

It is possible to select a subset of a table using the usual R matrix indexing on the table object. For example, this table contains rows with no data in them, and those yield ugly NA and NaN statistics:

```
> q <- data.frame(p = rep(c("A", "B"), each=10, len=30),
+ a = rep(c(1,2,3), each=10), id=seq(30),
+ b = round(runif(30,10,20)),
```

```
+ c = round(runif(30, 40, 70)),
+ stringsAsFactors = FALSE)
> tab <- tabular((Factor(p)*Factor(a)+1)
+ (N = 1) + (b + c)*(mean+sd), data=q)
> latex(tab)
```

			b		С	
p	a	N	mean	$\overline{\mathrm{sd}}$	mean	sd
Α	1	10	14.80	2.781	53.5	8.303
	2	0	NaN	NA	NaN	NA
	3	10	16.90	2.601	56.2	9.987
В	1	0	NaN	NA	NaN	NA
	2	10	14.90	2.283	51.7	8.642
	3	0	NaN	NA	NaN	NA
	All	30	15.53	2.662	53.8	8.892

To omit those rows, use matrix-like subsetting to select the rows where the first column of data (i.e. N) is greater than zero:

## > latex(tab[ tab[,1] > 0, ])

			b		С	
p	a	N	mean	$\operatorname{sd}$	mean	$\operatorname{sd}$
A	1	10	14.80	2.781	53.5	8.303
	3	10	16.90	2.601	56.2	9.987
В	2	10	14.90	2.283	51.7	8.642
	All	30	15.53	2.662	53.8	8.892

Similarly, cbind() can be used to join tables that have identical row labels, and rbind() can be used to join tables with identical column labels. Thus the top part of the table above could be produced in another way:

			b		c	
	a	N	mean	sd	mean	$\operatorname{sd}$
A	1	10	14.8	2.781	53.5	8.303
	3	10	16.9	2.601	56.2	9.987
В	2	10	14.9	2.283	51.7	8.642

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