

R Microplots in Tables with the `latex()` Function

Richard M. Heiberger, `rmh@temple.edu`
<http://astro.ocis.temple.edu/~rmh/>
Statistical Science, Fox School, Temple University

Microplots (sparklines) are often used within cells of a tabular array.

We describe several R functions that simplify the use of microplots within \LaTeX tables constructed in R with `Hmisc::latex` or a similar function. within **HTML** tables constructed with the **htmlTable** package.

We show examples using **base** graphics, **lattice** graphics, and **ggplot2** graphics.

These functions work in \LaTeX documents constructed directly in \LaTeX , with the R packages **Sweave**, **knitr**, or **rmarkdown**, and with the **Emacs** package **org-mode**.

1 Boxplots of iris data with lattice and latticeExtra

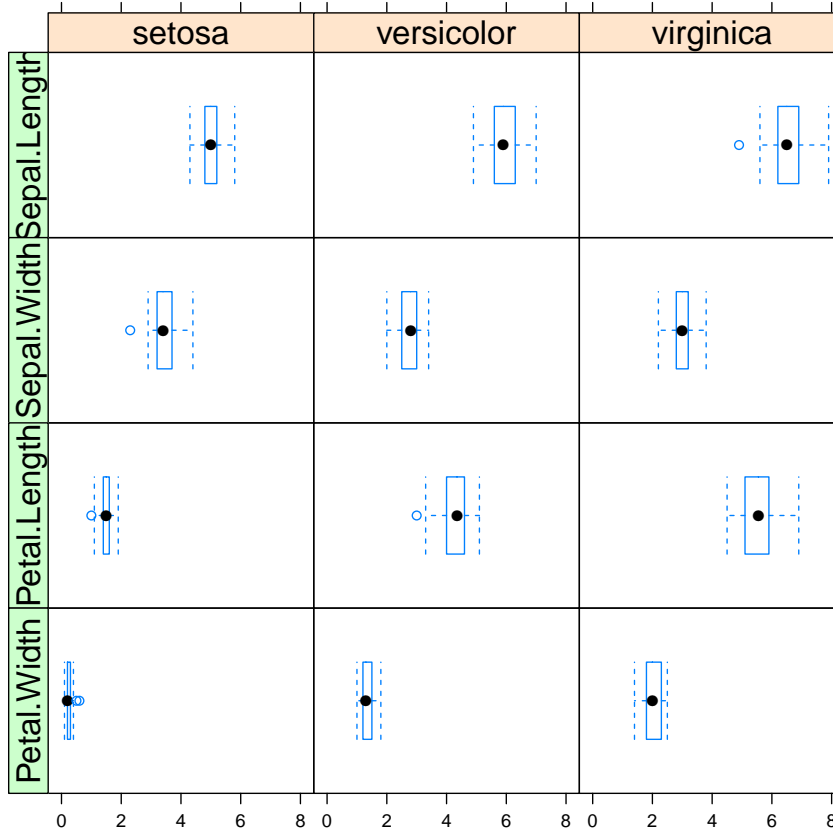














Figure 1: `useOuterStrips(`
`bwplot(~ Sepal.Length + Sepal.Width + Petal.Length + Petal.Width`
`| Species, data=iris, outer=TRUE, as.table=TRUE))`

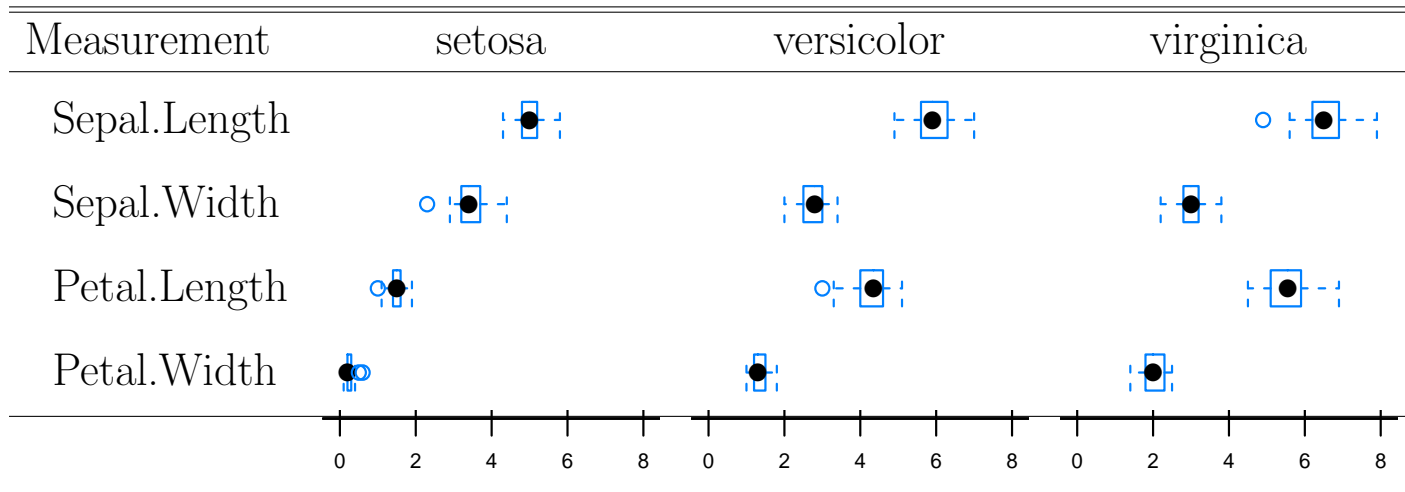
2 Individual boxes placed into a \LaTeX tabular environment

Table 1: Measurement by Species

Measurement	setosa	versicolor	virginica
Sepal.Length			
Sepal.Width			
Petal.Length			
Petal.Width			













3 Individual boxes in a table with the x -scale displayed

Table 2: Measurement by Species, with x -scale



4 Transposed L^AT_EX table

Table 3: Species by Measurement

Species	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
setosa				
versicolor				
virginica				

5 Individual boxes embedded into a more interesting table

Table 4: Five Number Summary and Boxplots for each Species and Measurement

Species	Measurement	Five Number Summary					Box Plots
		min	Q1	med	Q3	max	
setosa							
	Sepal.Length	4.3	4.8	5.00	5.2	5.8	
	Sepal.Width	2.3	3.2	3.40	3.7	4.4	
	Petal.Length	1.0	1.4	1.50	1.6	1.9	
	Petal.Width	0.1	0.2	0.20	0.3	0.6	
versicolor							
	Sepal.Length	4.9	5.6	5.90	6.3	7.0	
	Sepal.Width	2.0	2.5	2.80	3.0	3.4	
	Petal.Length	3.0	4.0	4.35	4.6	5.1	
	Petal.Width	1.0	1.2	1.30	1.5	1.8	
virginica							
	Sepal.Length	4.9	6.2	6.50	6.9	7.9	
	Sepal.Width	2.2	2.8	3.00	3.2	3.8	
	Petal.Length	4.5	5.1	5.55	5.9	6.9	
	Petal.Width	1.4	1.8	2.00	2.3	2.5	

6 How does it work?

There are two tasks. The **microplot** package provides functions for each task.

1. Isolate the contents of each panel of a multipanel graph into its own **pdf** file.

lattice: functions `layoutHeightsCollapse` and `layoutWidthsCollapse`
and argument `layout=c(1,1)`

ggplot2: function `theme_collapse` and loop through panels

base: adjust `par` arguments and `xlim` and `ylim`

2. Automate construction of the graphics statements.

L^AT_EX: `as.includegraphics` function

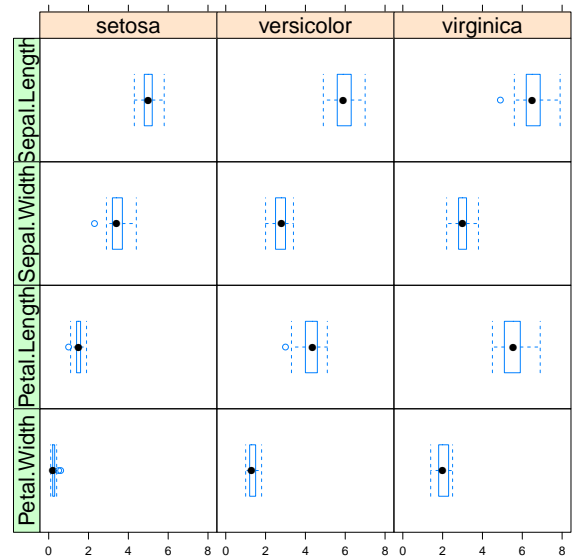
HTML: `as.htmling` function

org-mode: `as.orgtable` and `as.orgfile` functions

7 lattice

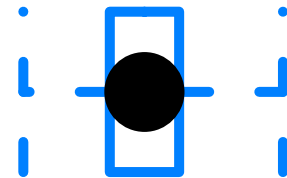
```
## boxplot matrix of iris data
irisBW <-
  bwplot(~ Sepal.Length + Sepal.Width + Petal.Length + Petal.Width |
         Species,
         data=iris, outer=TRUE, as.table=TRUE,
         scales=list(alternating=FALSE),
         xlab=NULL,
         par.strip.text=list(cex=1.5))
```

```
## pdf of boxplot matrix
pdf("irisBW.pdf")
useOuterStrips(irisBW)
suppress <- dev.off()
```




```
## twelve individual boxplots without axes
irisBW.update <-
update(irisBW,
      xlab=NULL,
      par.settings=list(
        layout.heights=layoutHeightsCollapse(),
        layout.widths=layoutWidthsCollapse(),
        axis.line=list(col="transparent")),
      layout=c(1,1)
    )

## create 12 pdf files, one per boxplot
pdf("irisBW%03d.pdf", onefile=FALSE, height=.4, width=1.6) ## inch
irisBW.update
suppress <- dev.off()
```



The first panel in file `irisBW001.pdf` is shown here.

The functions `layoutHeightsCollapse` (shown here) and `layoutWidthsCollapse` set the vertical and horizontal space for everything in a plot, except the panel itself, to 0.

```
> layoutHeightsCollapse
function (...)
{
  x.settings <- lattice::trellis.par.get()$layout.heights
  x.settings[] <- 0
  x.settings$panel = 1
  inputs <- list(...)
  if (length(inputs))
    x.settings[names(inputs)] <- inputs
  x.settings
}
```

The function `as.includegraphics` wraps the graph file names into the format used by the \LaTeX **graphicx** package.

```
> graphnames[1:2]
[1] "irisBW001.pdf" "irisBW002.pdf"
> graphicsnames <- as.includegraphics(graphnames[1:12], wd=".")
> dim(graphicsnames) <- c(4,3)
> graphicsnames[1:2, 1]
[1] "\\includegraphics[height=1em]{./irisBW001.pdf}"
[2] "\\includegraphics[height=1em]{./irisBW002.pdf}"
```

These values are placed into an ordinary matrix or dataframe and sent to the `Hmisc::latex` function to create a latex file fragment that can be input with the \LaTeX `\input` macro.

```
BWMS.latex <- Hmisc::latex(graphicsnames)
BWMS.latex$style <- "graphicx"
```

8 More Information on Microplots

The **microplot** package shows simple examples with **lattice**, **ggplot2**, and **base** graphics.

The **microplot** package shows simple examples in L^AT_EX using the R packages **Sweave**, **knitr**, and **rmarkdown**, and the Emacs package **org-mode**.

The **microplot** package shows simple examples in HTML using the R package **rmarkdown** and the Emacs package **org-mode**.

```
utils::install.packages("microplot", dependencies="r")  
## this includes HH and its dependencies
```

The **HH** package is designed to accompany
Statistical Analysis and Data Display, Second Edition
Richard M. Heiberger and Burt Holland
Springer 2015

<http://www.springer.com/us/book/9781493921218>

