hviciPlotting: Generating plot.sas style figures in R

John Ehrlinger Cleveland Clinic

Abstract

We introduce the R package hviciPlotting, a set of tools for creating publication quality graphics in R. The hviciPlotting package is designed to replace the plot.sas macro we currently use in SAS. The package includes both R recipes for generating our standard graphics using ggplot2 commands and a set of themes designed to format those figures for both manuscript and PowerPoint targets.

The goal of this package vignette is to introduce the **hviciPlotting** methodology, as well as to document the best practices of creating our publication quality graphics for both manuscripts and power point presentations.

This document is included with the **hviciPlotting** package as a package vignette, installed into R when the package is installed, and view able using the **vignette("hviciPlotting")** command.

Keywords: publication graphics, powerpoint, ggplot2, plot.sas.

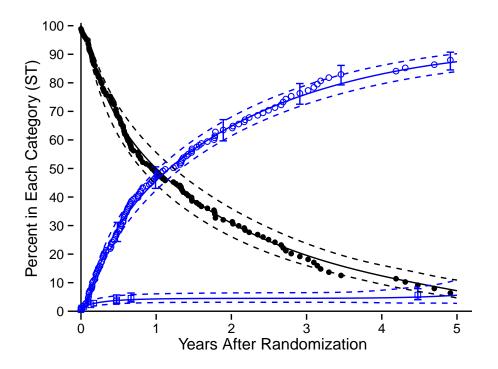


Figure 1: Demonstration figure

1. About this document

This document is an introduction to the R package hviciPlotting, a set of tools for creating publication quality graphics in R. The package and this document describe the process of creating graphics in R that conform to the standards of the clinical investigations statistics group within The Heart & Vascular Institute at the Cleveland Clinic. These graphics are analogous to those generated with the plot.sas macro in SAS.

This document is the package vignette for the **hviciPlotting** package, and as such is the primary documentation for the package. The latest version of the document can be obtained with the

R> vignette("hviciPlotting", package = "hviciplotting")

The goal is to update this document as the package is updated to include all relevant changes for publication.

2. Introduction

For many years, the mainstay for generating graphics for manuscripts and presentations in the statistics group in HVI has been the plot.sas macro using SAS. However, recently, we have had issues migrating this macro to newer versions of SAS (> 8.0) and MicroSoft Office products (> 2003).

In an effort to alleviate the versifying problems, and to standardize the generation of figures within R, we have developed the **hviciPlotting** R package. The goal of the package, and this document, is simplify the creation of publication quality graphics in R. We are specifically encoding the best practices of the HVI Clinical Investigations formatting, so that our statisticians will be able to simply create graphics for publication with a minimal amount of effort.

The hviciPlotting package also implements best practices for R graphics by leveraging the ggplot2 package (Wickham 2009). The ggplot2 package is an implementation of the Grammar of Graphics (Wilkinson 2005), which is a formalization of graphical concepts, and the building of graphical objects from a sequence of independent components. These components can be combined in many different ways.

The plot.sas macro is also an implementation of a graphics grammar. The grammar is derived from the ZETA pen plotters, which used GML (Graphics Machine Language) to control between 4 and and 8 colored pens for generating color line and point figures. Because both systems use a graphics language it is straight forward to translate commands between the two methods.

This document outlines how to generate figures using the **ggplot2** and **hviciPlotting** packages. Our approach is to demonstrate the R commands to generate the same elements created with plot.sas commands. Section 3 gives an overview of the methodology of the plot.sas macro and Section 4 details how to create line and point plots with **ggplot2**. A key part of **hviciPlotting** package is custom themes for figures. Once you have created your figure, Section 6 details how to get the formatting correct for manuscripts or presentations. Section 7 describes functions for saving the figures to simplify the import into publication documents.

3. The plot.sas macro

To demonstrate the process, we first look at some example code using the plot.sas macro. This code is intended to generate a figure for manuscript publication and was modified to generate Figure 1. Note the first line of the code block indicates the location of the file.

```
%let STUDY=/studies/cardiac/valves/aortic/replacement/
partner_publication_office/partner1b/mortality_5y
* Bring in PostScript plot macro
filename plt "!MACROS/plot.sas"; %inc plt;
filename gsasfile pipe 'lp';
                   POSTSCRIPT PLOTS
*_____;
* Multiple decrement, nonparametric and parametric
filename gsasfile "&STUDY/graphs/ce.states.ST_toJohn.both.ps";
%plot(goptions gsfmode=replace, device=pscolor, gaccess=gsasfile end;
id l="&STUDY/graphs/ce.states.ST_toJohn.sas percent", end;
labelx 1="Years After Randomization", end;
axisx order=(0 to 5 by 1), minor=none, end;
labely 1="Percent in Each Category (ST)", end;
axisy order=(0 to 100 by 10), minor=none, end;
```

We interupt this command here for some explanation. The plot.sas macro call starts with the %plot command. The first line sets global graphic values, including the file where the figure will be saved (see Section 7). Each plot.sas command is terminated with the end; statement. We'll look at each command type individually.

The id 1= command sets the footnote text used for manuscript figures to identify where the figure is saved (see Section 7). The labelx and labely commands set the axis label text (Section 4.4) and the axisx and axisy set the scales for each axis locating text and tics (Section 4.6).

The tuple command builds up graphics objects within the figure plot window. The first set of tuple commands builds up a set of three elements containing both points (Section 4.2) and errorbars (Section 4.3). Each tuple statement operates on a dataset indicated by the set command.

```
/******NON-PARAMETRIC: SYMBOLS AND CONFIDENCE BARS ******/
tuple set=green, symbol=dot, symbsize=1/2, linepe=0, linecl=0,
ebarsize=3/4, ebar=1,
x=iv_state, y=sginit, cll=stlinit, clu=stuinit, color=black,
end;
tuple set=green, symbol=circle, symbsize=1/2, linepe=0, linecl=0,
ebarsize=3/4, ebar=1,
x=iv_state, y=sgdead1, cll=stldead1, clu=studead1, color=blue,
end;
tuple set=green, symbol=square, symbsize=1/2, linepe=0, linecl=0,
ebarsize=3/4, ebar=1,
x=iv_state, y=sgstrk1, cll=stlstrk1, clu=stustrk1, color=blue,
end;
```

Symbols shapes and sizes are specified with the symbol and symbols commands (Section ??). The second set of tuple statements build up a set of three elements containing lines and confidence intervals (Section 4.1).

The plot.sas macro code is closed by the ending); characters, and SAS is instructed to run; the code. Running combines building the figure by combining elements from label, axis and tuple statements and saving it into the file specified by the gsasfile variable.

Note that much of the figure formatting is mixed within the tuple statements using width, color, linepe and linecl commands. In the plot.sas macro, omitting these commands will generate a figure with the default values specified within the device theme (Section 6). A similar set of plot.sas commands is used to create presentation graphics.

Differences include the target device and ftext as well as some handling of figure labels with value instead of label commands. We also have rules for what to and not to include in presentation graphics (Section 8).

4. Generating ggplot2 graphics

4.1. Lines

4.2. Points
4.3. ErrorBars
4.4. Labels
4.5. Colors
4.6. Scales
5. Generating other figure types
5.1. Bar Charts
5.2. Histograms
5.3. Additional Figure Types
6. ggplot2 themes for publication
6.1. Theme for Manuscripts
6.2. Theme for Presentations
7. Saving Publication graphics
7.1. Manuscript graphics
7.2. PowerPoint graphics

8. Graphics rules to live by

9. Conclusions

In this article, we present some functions in the hviciPlotting package for R

References

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Wilkinson L (2005). The Grammar of Graphics (Statistics and Computing). Springer-Verlag New York, Inc., Secaucus, NJ, USA. ISBN 0387245448.

Affiliation:

John Ehrlinger Quantitative Health Sciences Lerner Research Institute Cleveland Clinic 9500 Euclid Ave Cleveland, Ohio 44195

E-mail: john.ehrlinger@gmail.com

URL: http://www.biostat.uzh.ch/aboutus/people/rufibach.html