Graphics Devices in R

Biostatistics 140.776

What is a Graphics Device?

A graphics device is something where you can make a plot appear

- ► A window on your computer (screen device)
- ► A PDF file (file device)
- ► A PNG or JPEG file (file device)
- A scalable vector graphics (SVG) file (file device)

When you make a plot in R, it has to be "sent" to a specific graphics device

What is a Graphics Device?

The most common place for a plot to be "sent" is the screen device

- On a Mac the screen device is launched with the quartz()
- ▶ On Windows the screen device is launched with windows()
- On Unix/Linux the screen device is launched with x11()

What is a Graphic Device?

When making a plot, you need to consider how the plot will be used to determine what device the plot should be sent to.

► The list of devices is found in ?Devices; there are also devices created by users on CRAN

For quick visualizations and exploratory analysis, usually you want to use the screen device

- Functions like plot in base, xyplot in lattice, or qplot in ggplot2 will default to sending a plot to the screen device
- ► On a given platform (Mac, Windows, Unix/Linux) there is only one screen device

For plots that may be printed out or be incorporated into a document (e.g. papers/reports, slide presentations), usually a *file device* is more appropriate

 NOTE: Not all graphics devices are available on all platforms (i.e. you cannot launch the windows() on a Mac)

There are two basic approaches to plotting. The first is most common:

- Call a plotting function like plot or qplot
- 2. The plot appears on the screen device
- 3. Annotate plot if necessary (e.g. adding geom_*s)
- 4. Enjoy

```
library(datasets)
library(ggplot2)

## Make plot appear on screen device
## Annotate with a title
qplot(eruptions, waiting, data = faithful)
+ labs(title = "Old Faithful Geyser data")
```

The second approach to plotting is most commonly used for file devices:

- 1. Explicitly launch a graphics device
- 2. Call a plotting function to make a plot (Note: if you are using a file device, no plot will appear on the screen)
- 3. Annotate plot if necessary
- Explicitly close graphics device with dev.off() (this is very important!)

```
library(ggplot2)
## Open PDF device
## Create 'myplot.pdf' in working directory
pdf(file = "myplot.pdf")
## Create plot and send to a file
## (no plot appears on screen)
qplot(eruptions, waiting, data = faithful)
+ labs(title = "Old Faithful Geyser data")
## Close the PDF file device
dev.off()
## Now you can view the file 'myplot.pdf'
```

Graphics File Devices

There are two basic types of file devices: *vector* and *bitmap* devices Vector formats:

- pdf: useful for line-type graphics, resizes well, usually portable, not efficient if a plot has many objects/points
- svg: XML-based scalable vector graphics; supports animation and interactivity, potentially useful for web-based plots
- postscript: older format, also resizes well, usually portable, can be used to create encapsulated postscript files; Windows systems often don't have a postscript viewer

Graphics File Devices

Bitmap formats

- png: bitmapped format, good for line drawings or images with solid colors, uses lossless compression (like the old GIF format), most web browsers can read this format natively, good for plotting many many many points, does not resize well
- jpeg: good for photographs or natural scenes, uses lossy compression, good for plotting many many many points, does not resize well, can be read by almost any computer and any web browser, not great for line drawings
- tiff: Creates bitmap files in the TIFF format; supports lossless compression

Multiple Open Graphics Devices

- ▶ It is possible to open multiple graphics devices (screen, file, or both), for example when viewing multiple plots at once
- Plotting can only occur on one graphics device at a time
- ➤ The currently active graphics device can be found by calling dev.cur()
- ▶ Every open graphics device is assigned an integer ≥ 2 .
- ➤ You can change the active graphics device with dev.set(<integer>) where <integer> is the number associated with the graphics device you want to switch to

Copying Plots

Copying a plot to another device can be useful because some plots require a lot of code and it can be a pain to type all that in again for a different device.

- dev.copy: copy a plot from one device to another
- dev.copy2pdf: specifically copy a plot to a PDF file

NOTE: Copying a plot is not an exact operation, so the result may not be identical to the original.

Copying Plots

```
library(datasets)
library(ggplot2)
## Create plot on screen device
qplot(eruptions, waiting, data = faithful)
 + labs(title = "Old Faithful Geyser data")
## Copy my plot to a PNG file
dev.copy(png, file = "geyserplot.png")
## Don't forget to close the PNG device!
dev.off()
```

Summary

- ▶ Plots must be created on a graphics device
- ► The default graphics device is almost always the screen device, which is most useful for exploratory analysis
- ▶ File devices are useful for creating plots that can be included in other documents or sent to other people
- For file devices, there are vector and bitmap formats
- Vector formats are good for line drawings and plots with solid colors using a modest number of points
- Bitmap formats are good for plots with a large number of points, natural scenes or web-based plots