Graduate School Class Reminders

- ► Maintain six feet of distancing
- ▶ Please sit in the same chair each class time
- ► Observe entry/exit doors as marked
- ► Use hand sanitizer when you enter/exit the classroom
- Use a disinfectant wipe/spray to wipe down your learning space before and after class
- ► Media Services: 414 955-4357 option 2

Documentation on the web

- ► CRAN: http://cran.r-project.org
- ► R manuals: https://cran.r-project.org/manuals.html
- ► SAS: http://support.sas.com/documentation
- ► Step-by-Step Programming with Base SAS 9.4 (SbS): https://documentation.sas.com/api/docsets/basess/ 9.4/content/basess.pdf
- ► SAS 9.4 Programmer s Guide: Essentials (PGE): https://documentation.sas.com/api/docsets/lepg/9.4/content/lepg.pdf
- ► Wiki: https://wiki.biostat.mcw.edu (MCW/VPN)
- ► lattice: http://lmdvr.r-forge.r-project.org/ figures/figures.html

- ► There are two graphics packages for R that have a similar syntax which makes learning both of them convenient
- ► The graphics package for routine daily usage is one of the base packages as described in Section 12
- ► Easy to use, it can create high-quality graphical figures, but there are limits to their customizability
- ➤ You can see the documentation with library(help=graphics)
- Depends on the grDevices package which has support for devices, colors, fonts, etc.

- ► The lattice package which is one of the *recommended* packages can create more appealing customizable graphical figures of publication and grant proposal quality
- ► It builds upon grDevices and graphics
- ► With some added capabilities in latticeExtra
- ► You can see the documentation with library(help=lattice)
- ► But, the lattice package documentation is voluminous and it can take quite a while to read and understand
- ► Yet, the examples are very helpful; there are extra examples online that will dispell some of the mystery http://lmdvr.r-forge.r-project.org/figures/figures.html

- ► DO NOT USE ggplot2!
- ► It is unacceptable for the following reasons
- ► The graphs are terrible: a gray background by default?
- ▶ ggplot2 package doesn't come with R
- ▶ Its syntax doesn't resemble graphics/lattice
- ► It has no 3D capabilities

► Common types of plots and their high-level functions

Type of plot	Dimensions	graphics	lattice
Box and whisker	1	boxplot	bwplot
Histogram	1	hist	histogram
Density	1	plot.density	densityplot
Quantile-quantile	1	qqnorm	qqmath
QQ	2	qqplot	qq
Scatter	2	plot	xyplot
Contour	3	contour	contourplot
Heat map	3	image	levelplot
Surface	3	persp	wireframe

- ► Let's explore the high-level plot function
- Many objects types have their own generic functions
 plot(object) will call the function plot.CLASS where the object is of type CLASS like plot.density
- ▶ But, we are more concerned with the default: plot.default
- ► Here the arguments of primary interest that are shared/comparable between graphics and lattice

```
plot(x, y = NULL, col = 1,
    type = "p", cex = 1, pch = 1, lty = 1, lwd = 1,
    xlim = NULL, ylim = NULL,
    xlab = NULL, ylab = NULL)
```

```
plot(x, y = NULL, col = 1,
    type = "p", cex = 1, pch = 1, lty = 1, lwd = 1,
    xlim = NULL, ylim = NULL,
    xlab = NULL, ylab = NULL)
```

- \triangleright x is a vector for the x-axis
- ▶ y same thing for the y-axis if y is not specified, then x is plotted as a time series
- common type settings: "p" for points, "1" for lines,
 "b" for both points and lines, "h" for vertical lines,
 "s" for stair steps/survival functions and
 "n" does not produce anything
- ► cex for the size of points and pch for the kind
- ▶ specify the line type by lty and the line width by lwd
- ightharpoonup xlim is a length 2 vector for the limits of the x-axis
- ► xlab is the label for the *x*-axis: either a character string or an expression see > ?plotmath
- ▶ ylim/ylab same thing for the y-axis

```
plot(x, y = NULL, col = 1,
    type = "p", cex = 1, pch = 1, lty = 1, lwd = 1,
    xlim = NULL, ylim = NULL,
    xlab = NULL, ylab = NULL)
```

- ► col is the color to be used for the plot points/lines/etc.
- ► A color's name by character string: see colors for a list
- Or it can be an integer marker number
- ► This is based on the current color palette which can be customized by the palette function: default markers
- ► Choosing good colors can be tricky: stick to the powers of 2 $0 = 2^{-\infty} \quad \text{white} \qquad \begin{vmatrix} 1 = 2^0 & black \\ 3 & \text{green} \end{vmatrix} = \begin{vmatrix} 2 = 2^1 & red \\ 4 = 2^2 & blue \end{vmatrix} = \begin{vmatrix} 2 = 2^1 & cyan \\ 5 & cyan \\ 8 = 2^3 & gray \end{vmatrix}$
- ▶ If you need more, then see bass/colors.R

- ► Interactively, plot will create a graphics window with a panel
- ➤ Or multiple plot calls yielding multiple panels via par(mfcol=c(rows, cols)) or par(mfrow=c(rows, cols)) but often with very poor quality: see lecture5.R
- ► Low-level functions that overlay graphics on this panel follow
- ► Those that are self-explanatory and similar to lattice
- ► lines(x, y=NULL, col=1, lty=1, lwd=1)
- ▶ points(x, y=NULL, col=1, pch=1, cex=1)
- text(x, y=NULL, labels, col=1, cex=1, pos=NULL)
 pos 3
 2 (x, y) 4 positions go clock-wise starting at 6 o'clock
- ► legend is NOT similar to lattice
- ► legend(x, y=NULL, col=1, legend, lty=1, lwd=1, pch=1, cex=1, horiz=FALSE)

- ► This is the same for the lattice package
- > ?Devices lists the graphical file formats supported
- We are going to restrict our attention to Adobe Portable Document Format (PDF)
- ▶ In an interactive R session, you can capture what is in the graphics window at any time by (in batch, this is an error) > dev.copy2pdf(file="FILENAME.pdf")
- ► In an interactive or batch R session, you can create a graphics file without a graphics window
- pdf(file="FILENAME.pdf")
 PLOTTING STATEMENTS
 dev.off()
- ► The latter can create graphics files with multiple pages each plot creates a page (except for multiple panels where each of these is a page)

```
xyplot(formula, data, groups = NULL, col = 1,
    type = "p", cex = 1, pch = 1, lty = 1, lwd = 1,
    xlim = NULL, ylim = NULL, xlab = NULL, ylab = NULL,
    strip = strip.custom(strip.names = FALSE),
    layout = c(cols, rows), as.table = FALSE)
```

- ▶ formula of the form y~x for a single panel or y~x|g1*g2*... for panels of plots conditioned on the variables g1*g2*... which are typically factors if not factors, then specify factor(g1)*factor(g2)*...
- ▶ data is a data frame with x and y, etc.
- ► groups=z overlays data within the same panel based on different values of the variable z
- ▶ layout controls how multiple panels are configured, if any
- ➤ as.table controls the order multiple plans are drawn as.table = TRUE seems more logical than the default
- strip controls how the panels are labeled
 strip.names = TRUE seems more logical than the default

- ► Interactively, xyplot will create a graphics window
- ► Low-level functions can overlay graphics on this panel
- Those that are similar to graphics follow
- typically, just prefix an 1 on their names for lattice
- ▶ llines(x, y=NULL, col=1, lty=1, lwd=1) adds lines
 ▶ lpoints(x, y=NULL, col=1, pch=1, cex=1) adds points
- ▶ Ipoints(x, y=NULL, col=1, pch=1, cex=1) adds points
 ▶ ltext(x, y=NULL, col=1, labels, cex=1,
- pos=NULL, col=1, labels, cex=1,
 pos=NULL, offset=0.5) adds text
- ► And there is an additional one like plot itself lplot.xy(list(x=x, y=y), col=1, type="p", cex=1, pch=1, lty=1, lwd=1)

cex=1, pch=1, lty=1, lwd=1)

xyplot(formula, data, ## the easy way
panel=function(...){ panel.abline(h=0);panel.xyplot(...)})

xyplot(formula, data) ## or the hard way

xyplot(formula, data) ## or the hard way
update(trellis.last.object(),
 panel=function(...) {
 i=panel.number() ## call panel.FUNCTIONS and
 15/17

- ► As we saw, there are three 3D functions in graphics: contour, image and persp
- ► Each of these functions has an interface like NAME(x, y, Z) where x and y are vectors and Z is a matrix
- ► This is a confusing interface to say the least i.e., how does Z relate to x and y?
- ► lattice has a more intuitive interface for the corresponding functions: contourplot, levelplot and wireframe as follows (with arguments analogous to xyplot)
- ► NAME(z~x*y|g1*g2*..., data) where x, y and z are all vectors in the data frame so their relationship is obvious
- ► Many of the same arguments as xyplot
- groups is a notable exception

- ► There is a legend argument but it seems difficult to use see key instead
- key=list(border="black", x=x, y=y,
 text=list(...), ..., lines=list(...))
 or ..., points=list(...))
 where x and y in [0, 1] of the displayed area
- xlim and ylim are too simplistic for many needs scales is more flexible
- scales=list(
 x=list(limits=c(low, high), log=TRUE),
 y=list(at=c(1, 2, 4, 8),
 labels=c("a", "b", "c", "d")))
- ▶ at are the tick marks drawn: can be increasing or decreasing
- ▶ labels are the values to display there

- ▶ trellis.par.get() to show default settings
- ▶ and trellis.par.set() to alter them
- ► for example, slide 13 shows the default for xyplot is col = 1 however, that is NOT right: more complicated than that

```
trellis.par.get("plot.symbol")
a$col = 1
trellis.par.set("plot.symbol", a)
```

- ► You can overlay two or more trellis objects
- ► For example, doubleYScale to create a second y-axis
- ► Combine two objects with the as.layer function
- ► Combine multiple objects with the c function