Plotting with ggplot2

Exploratory Data Analysis

Roger D. Peng, Associate Professor of Biostatistics Johns Hopkins Bloomberg School of Public Health

What is ggplot2?

- An implementation of the *Grammar of Graphics* by Leland Wilkinson
- Written by Hadley Wickham (while he was a graduate student at Iowa State)
- A "third" graphics system for R (along with base and lattice)
- Available from CRAN via install.packages()
- Web site: http://ggplot2.org (better documentation)

What is ggplot2?

- Grammar of graphics represents and abstraction of graphics ideas/objects
- Think "verb", "noun", "adjective" for graphics
- Allows for a "theory" of graphics on which to build new graphics and graphics objects
- "Shorten the distance from mind to page"

Grammar of Graphics

"In brief, the grammar tells us that a statistical graphic is a mapping from data to aesthetic attributes (colour, shape, size) of geometric objects (points, lines, bars). The plot may also contain statistical transformations of the data and is drawn on a specific coordinate system" from ggplot2 book

The Basics: qplot()

- Works much like the plot function in base graphics system
- Looks for data in a data frame, similar to lattice, or in the parent environment
- Plots are made up of aesthetics (size, shape, color) and geoms (points, lines)

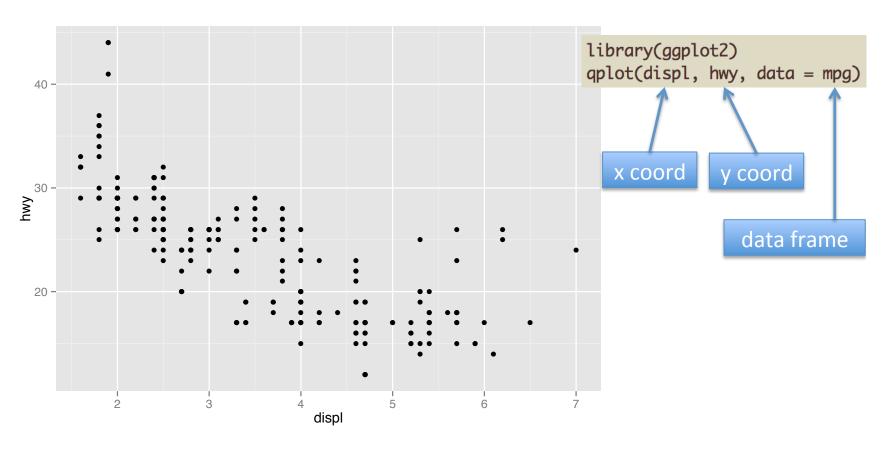
The Basics: qplot()

- Factors are important for indicating subsets of the data (if they are to have different properties); they should be labeled
- The qplot() hides what goes on underneath, which is okay for most operations
- ggplot() is the core function and very flexible for doing things qplot() cannot do

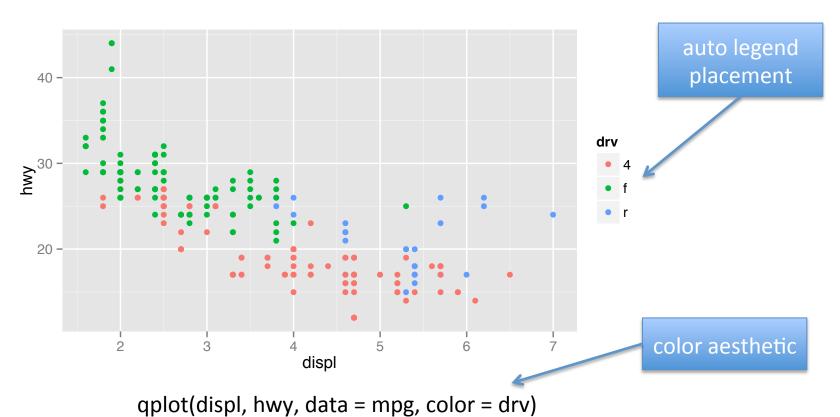
Example Dataset

```
> library(ggplot2)
                                                            Factor label information
> str(mpg)
                                                            important for annotation
'data.frame': 234 obs. of 11 variables:
$ manufacturer: Factor w/ 15 levels "audi", "chevrolet", ...: 1 1 1 1 1 1 1 1 1 ...
$ model : Factor w/ 38 levels "4runner 4wd",..: 2 2 2 2 2 2 3 3 3 ...
$ displ : num 1.8 1.8 2 2 2.8 2.8 3.1 1.8 1.8 2 ...
             : int 1999 1999 2008 2008 1999 1999 2008 1999 1999 2008 ...
$ year
$ cyl : int 4 4 4 4 6 6 6 4 4 4 ...
              : Factor w/ 10 levels "auto(av)". "auto(l3)",...: 4 9 10 1 4 9 1 9 4 10
$ trans
              : Factor w/ 3 levels "4", "f", "r": 2 2 2 2 2 2 1 1 1 ...
$ drv
$ cty
              : int 18 21 20 21 16 18 18 18 16 20 ...
$ hwy
             : int 29 29 31 30 26 26 27 26 25 28 ...
              : Factor w/ 5 levels "c", "d", "e", "p", ...: 4 4 4 4 4 4 4 4 4 4 ...
$ fl
$ class
              : Factor w/ 7 levels "2seater", "compact", ...: 2 2 2 2 2 2 2 2 2 2 ...
```

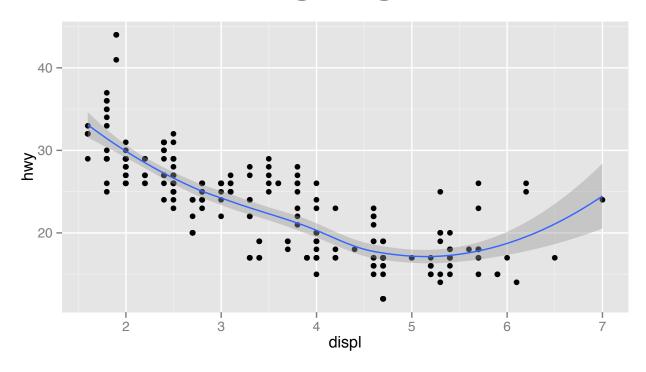
ggplot2 "Hello, world!"



Modifying aesthetics

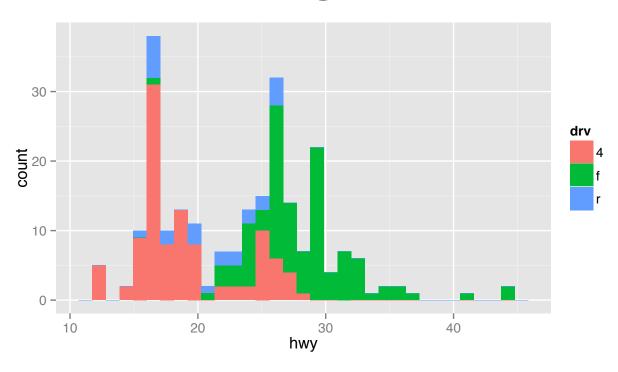


Adding a geom



qplot(displ, hwy, data = mpg, geom = c("point", "smooth"))

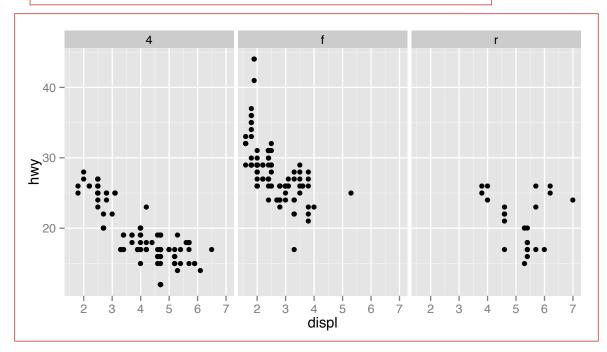
Histograms

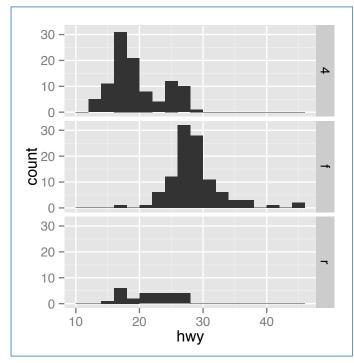


qplot(hwy, data = mpg, fill = drv)

Facets

qplot(displ, hwy, data = mpg, facets = . ~ drv)





qplot(hwy, data = mpg, facets = drv ~ ., binwidth = 2)

MAACS Cohort

- Mouse Allergen and Asthma Cohort Study
- Baltimore children (aged 5—17)
- Persistent asthma, exacerbation in past year
- Study indoor environment and its relationship with asthma morbidity
- Recent publication: http://goo.gl/WqE9j8

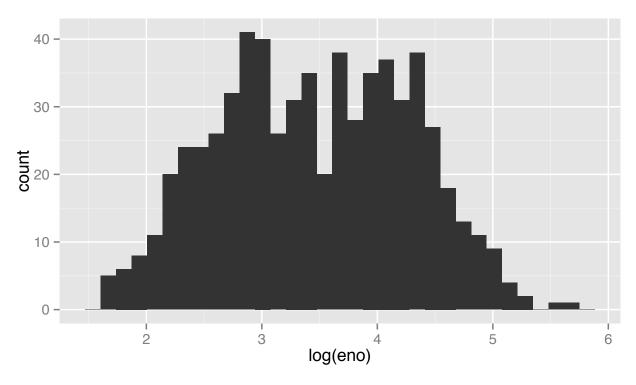
Exhaled nitric oxide

Example: MAACS

Fine particulate matter

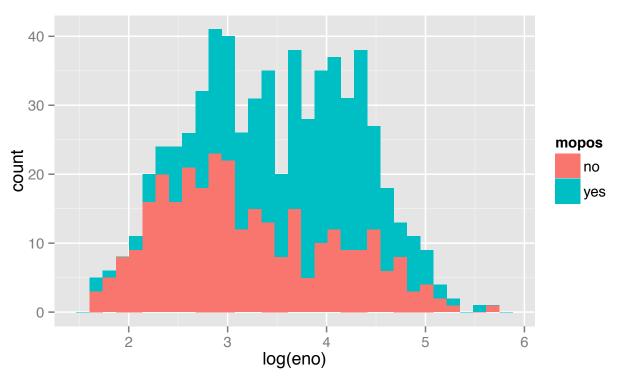
Sensitized to mouse allergen

Histogram of eNO



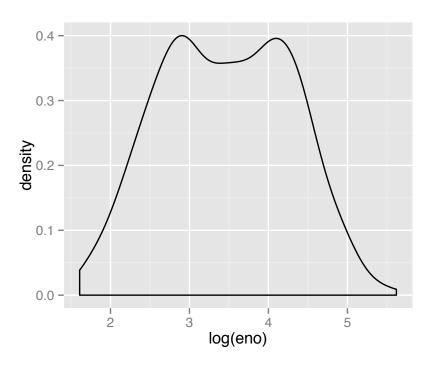
qplot(log(eno), data = maacs)

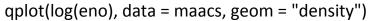
Histogram by Group

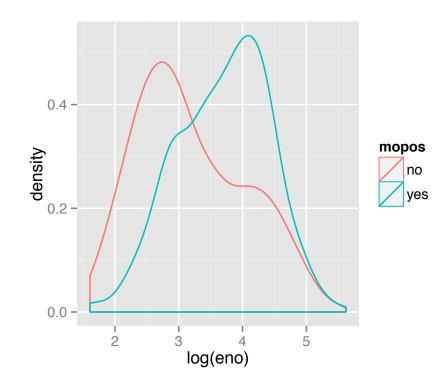


qplot(log(eno), data = maacs, fill = mopos)

Density Smooth

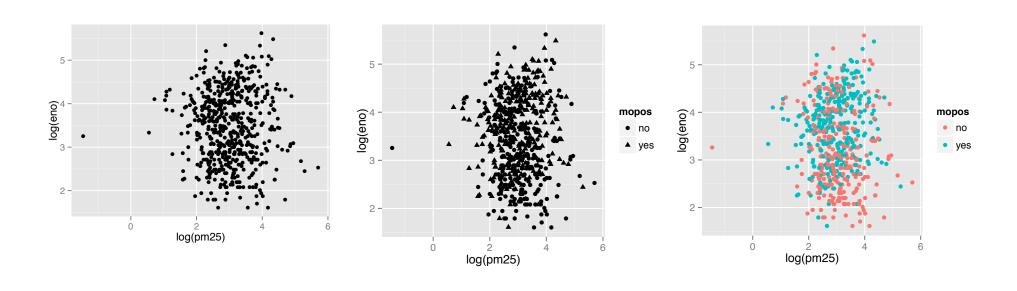






qplot(log(eno), data = maacs, geom = "density", color = mopos)

Scatterplots: eNO vs. PM_{2.5}

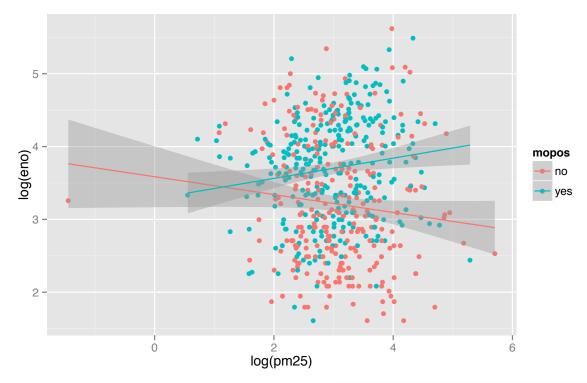


qplot(log(pm25), log(eno), data =
maacs)

qplot(log(pm25), log(eno), data =
maacs, shape = mopos)

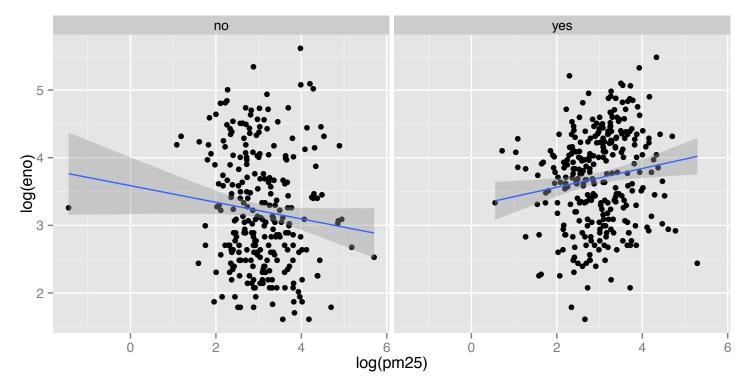
qplot(log(pm25), log(eno), data =
maacs, color = mopos)

Scatterplots: eNO vs. PM_{2.5}



qplot(log(pm25), log(eno), data = maacs, color = mopos, geom = c("point", "smooth"), method = "lm")

Scatterplots: eNO vs. PM_{2 5}



qplot(log(pm25), log(eno), data = maacs, geom = c("point", "smooth"), method = "lm", facets = . ~ mopos)

Summary of qplot()

- The qplot() function is the analog to plot() but with many built-in features
- Syntax somewhere in between base/lattice
- Produces very nice graphics, essentially publication ready (if you like the design)
- Difficult to go against the grain/customize (don't bother; use full ggplot2 power in that case)

Resources

- The ggplot2 book by Hadley Wickham
- The *R Graphics Cookbook* by Winston Chang (examples in base plots and in ggplot2)
- ggplot2 web site (http://ggplot2.org)
- ggplot2 mailing list (http://goo.gl/OdW3uB),
 primarily for developers

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Basic Components of a ggplot2 Plot

- A data frame
- aesthetic mappings: how data are mapped to color, size
- **geoms**: geometric objects like points, lines, shapes.
- **facets**: for conditional plots.
- **stats**: statistical transformations like binning, quantiles, smoothing.
- scales: what scale an aesthetic map uses (example: male = red, female = blue).
- coordinate system

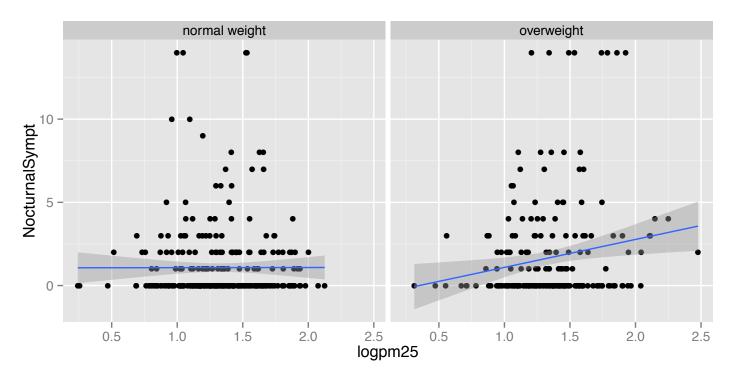
Building Plots with ggplot2

- When building plots in ggplot2 (rather than using qplot) the "artist's palette" model may be the closest analogy
- Plots are built up in layers
 - Plot the data
 - Overlay a summary
 - Metadata and annotation

Example: BMI, PM_{2.5}, Asthma

- Mouse Allergen and Asthma Cohort Study
- Baltimore children (age 5-17)
- Persistent asthma, exacerbation in past year
- Does BMI (normal vs. overweight) modify the relationship between PM_{2.5} and asthma symptoms?

Basic Plot



qplot(logpm25, NocturnalSympt, data = maacs, facets = . ~ bmicat, geom =
c("point", "smooth"), method = "lm")

Building Up in Layers

```
> head(maacs)
    logpm25
                    bmicat NocturnalSympt
2 1.5361795 normal weight
                                                               Data Frame
3 1.5905409 normal weight
                                         0
4 1.5217786 normal weight
                                         0
5 1.4323277 normal weight
                                                       Aesthetics
6 1.2762320
               overweight
                                         8
8 0.7139103
               overweight
                                         0
                                                               Initial call to
                                                                  ggplot
> q <- ggplot(maacs, aes(logpm25, NocturnalSympt))</pre>
> summary(q)
data: logpm25, bmicat, NocturnalSympt [554x3]
                                                               Summary of
mapping: x = logpm25, y = NocturnalSympt
                                                               ggplot object
faceting: facet null()
```

No Plot Yet!

```
> g <- ggplot(maacs, aes(logpm25, NocturnalSympt))
> print(g)
Error: No layers in plot

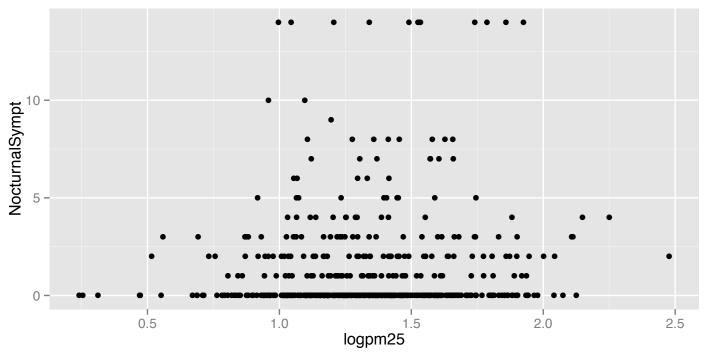
> p <- g + geom_point()
> print(p)

Explicitly save and print
ggplot object

> g + geom_point()

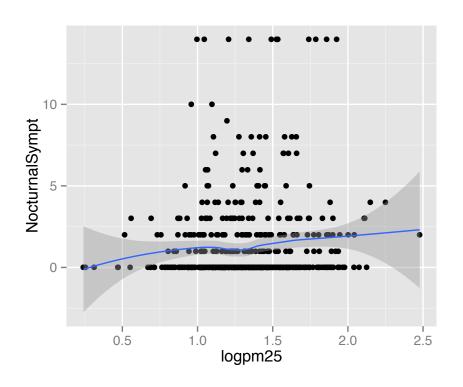
Auto-print plot object
without saving
```

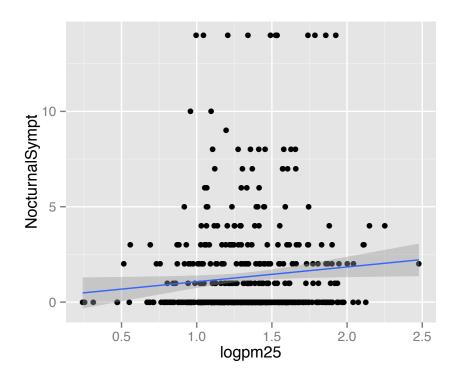
First Plot with Point Layer



```
g <- ggplot(maacs, aes(logpm25, NocturnalSympt))
g + geom_point()</pre>
```

Adding More Layers: Smooth

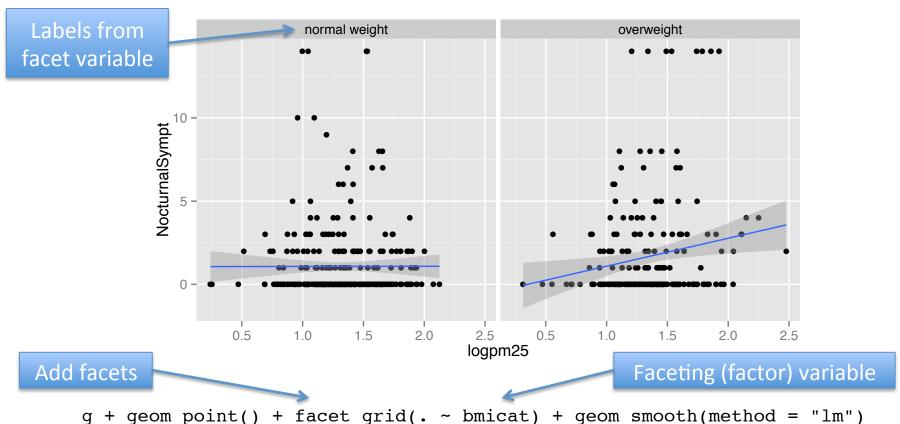




g + geom_point() + geom_smooth()

g + geom_point() + geom_smooth(method = "lm")

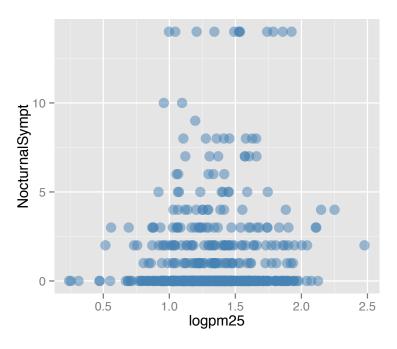
Adding More Layers: Facets

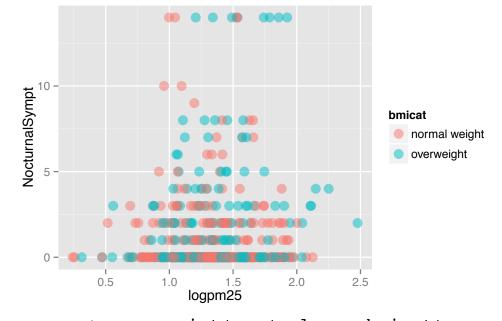


Annotation

- Labels: xlab(), ylab(), labs(), ggtitle()
- Each of the "geom" functions has options to modify
- For things that only make sense globally, use theme()
 - Example: theme(legend.position = "none")
- Two standard appearance themes are included
 - theme_gray(): The default theme (gray background)
 - theme_bw(): More stark/plain

Modifying Aesthetics





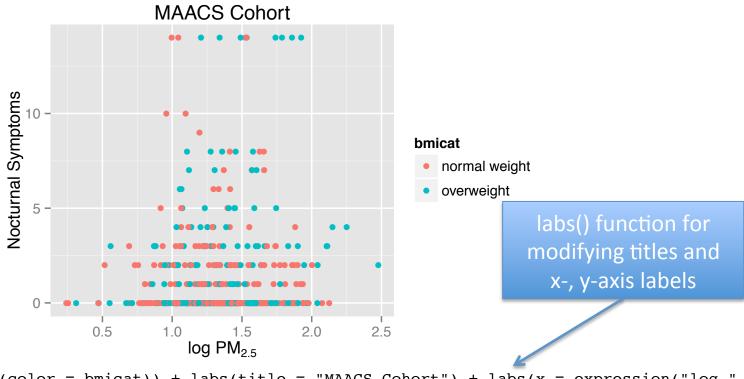
g + geom_point(color = "steelblue",
size = 4, alpha = 1/2)

g + geom_point(aes(color = bmicat),
size = 4, alpha = 1/2)

Constant values

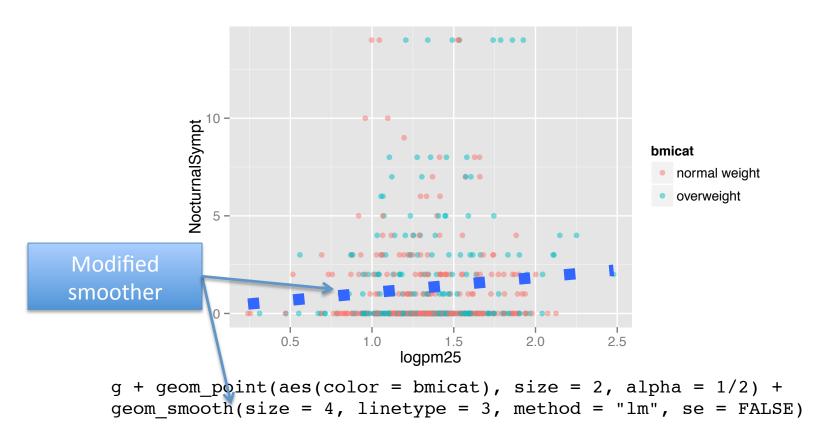
Data variable

Modifying Labels

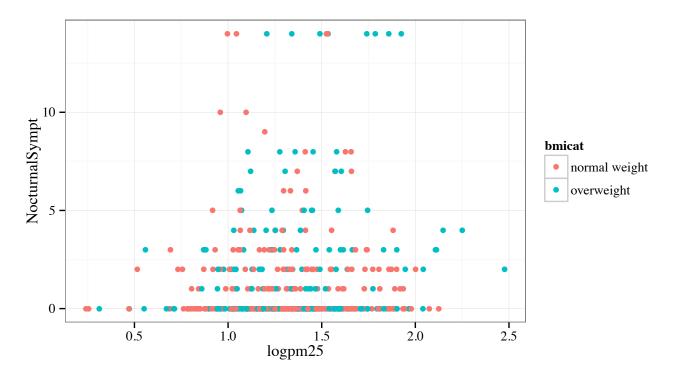


g + geom_point(aes(color = bmicat)) + labs(title = "MAACS Cohort") + labs(x = expression("log "
* PM[2.5]), y = "Nocturnal Symptoms")

Customizing the Smooth

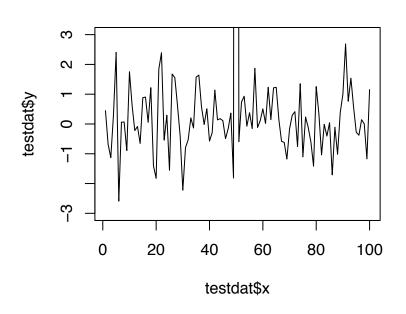


Changing the Theme

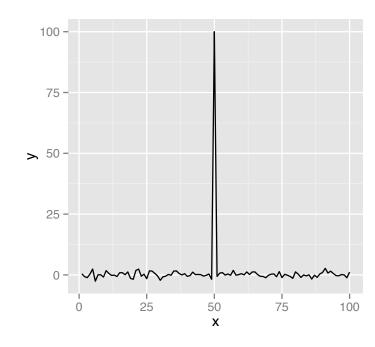


```
g + geom_point(aes(color = bmicat)) + theme_bw(base_family = "Times")
```

A Notes about Axis Limits



```
testdat <- data.frame(x = 1:100, y = rnorm(100))
testdat[50,2] <- 100 ## Outlier!
plot(testdatx, testdaty, type = "1", ylim = c(-3,3))
```



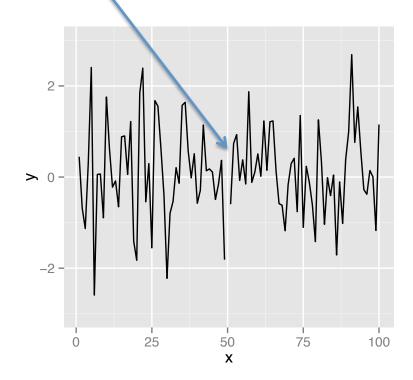
```
g \leftarrow ggplot(testdat, aes(x = x, y = y))

g + geom_line()
```

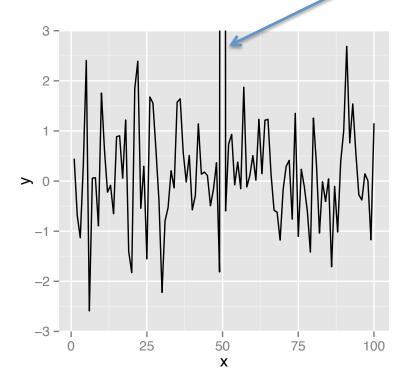


Axis Limits





 $g + geom_line() + ylim(-3, 3)$



g + geom_line() + coord_cartesian(ylim = c(-3, 3))

More Complex Example

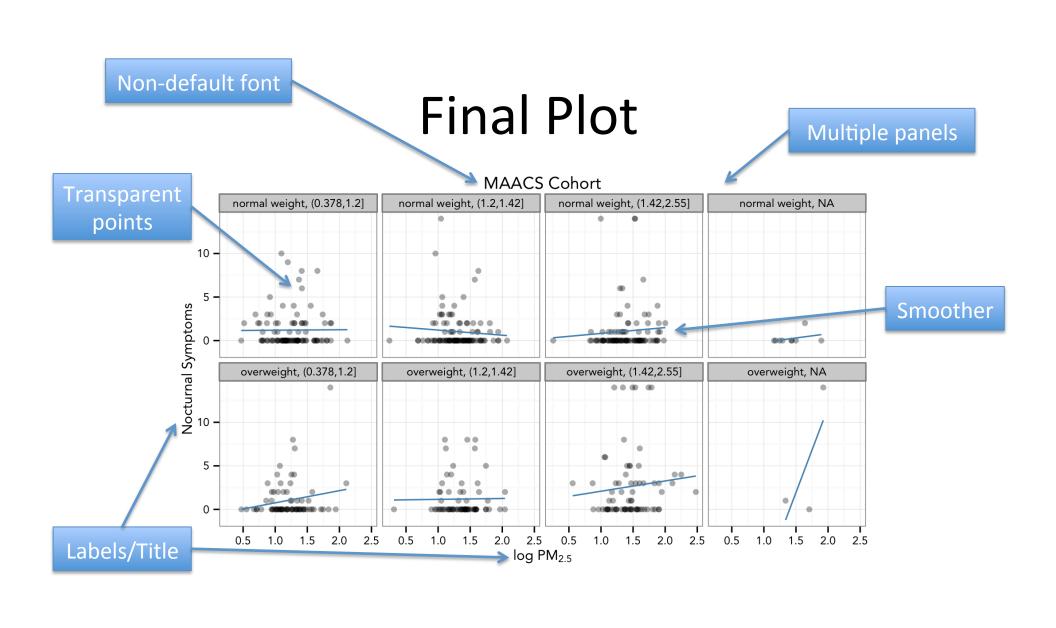
- How does the relationship between PM_{2.5} and nocturnal symptoms vary by BMI and NO₂?
- Unlike our previous BMI variable, NO₂ is continuous
- We need to make NO2 categorical so we can condition on it in the plotting
 - Use the cut() function for this

Making NO₂ Deciles

```
## Calculate the deciles of the data
> cutpoints <- quantile(maacs$logno2_new, seq(0, 1, length = 11), na.rm = TRUE)

## Cut the data at the deciles and create a new factor variable
> maacs$no2dec <- cut(maacs$logno2_new, cutpoints)

## See the levels of the newly created factor variable
> levels(maacs$no2dec)
[1] "(0.378,0.969]" "(0.969,1.1]" "(1.1,1.17]" "(1.17,1.26]"
[5] "(1.26,1.32]" "(1.32,1.38]" "(1.38,1.44]" "(1.44,1.54]"
[9] "(1.54,1.69]" "(1.69,2.55]"
```



Code for Final Plot

Summary

- ggplot2 is very powerful and flexible if you learn the "grammar" and the various elements that can be tuned/modified
- Many more types of plots can be made; explore and mess around with the package (references mentioned in Part 1 are useful)