

Linear models

Modeling... linearly!

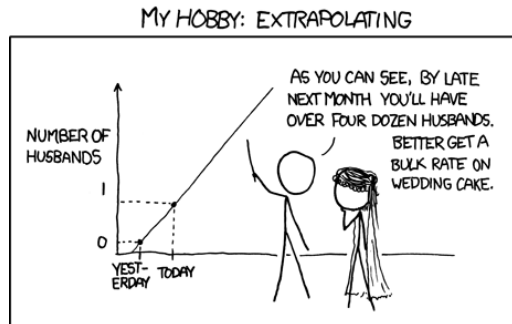
Samuel Robinson, Ph.D.

Sep. 22, 2023

Part 1: How do they work?

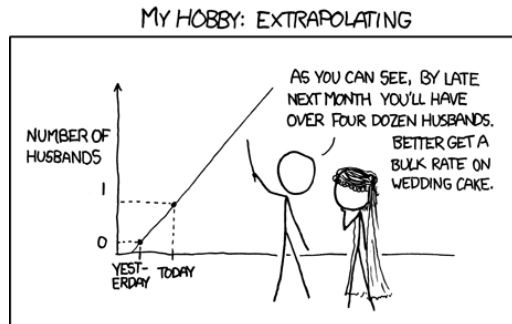
Outline

- What are linear models? How do I fit them?



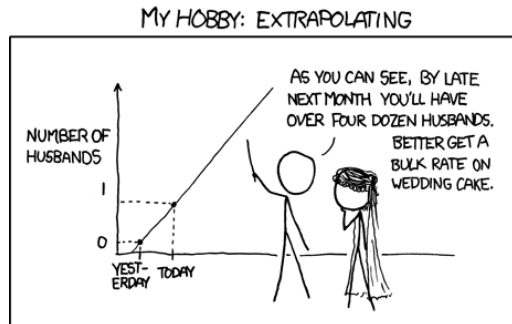
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- What are linear models? How do I fit them?
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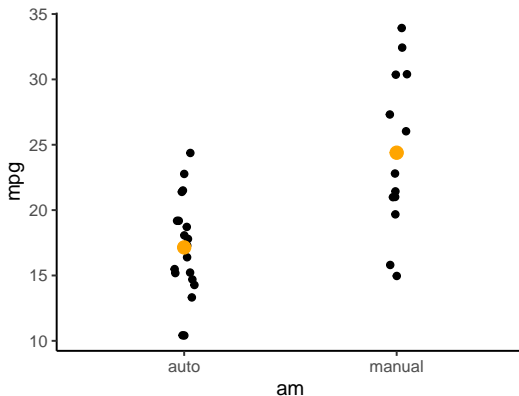
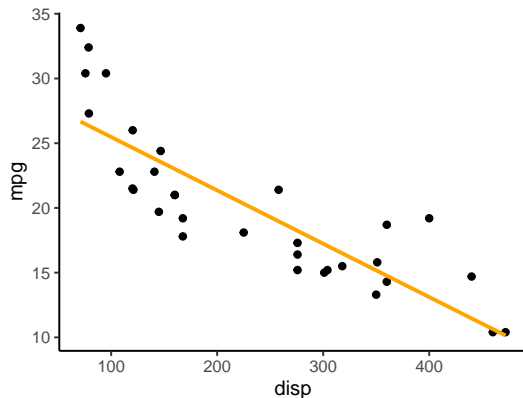
Outline

- What are linear models? How do I fit them?
- Making sure the model is working properly
- Plotting and interpreting model results



Motivation

- *I measured 2 things and I want to know if they're related to each other*
- *I have groups of data, and I want to know whether the means are different*



Terminology

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I use a set of terminology that I find very helpful, from [Berliner \(1996\)](#). I'll be using it here, as well as for describing more complex models.

Model terminology

All linear models take the form:

$$\hat{y} = b_0 + b_1x_1 + b_2x_2 \dots + b_ix_i$$

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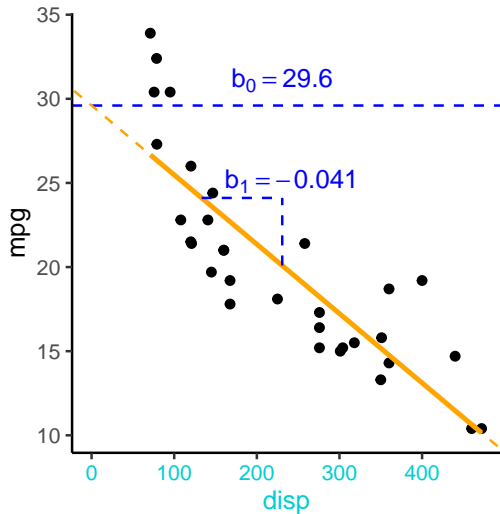
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This may look terrifying, but let's use a simple example:

Example

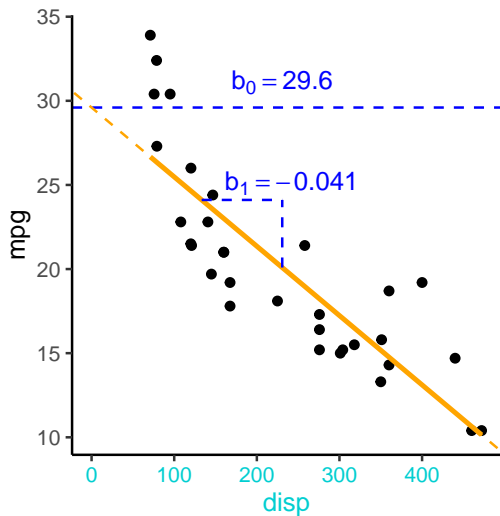


$$\hat{mpg} = b_0 + b_1 disp$$

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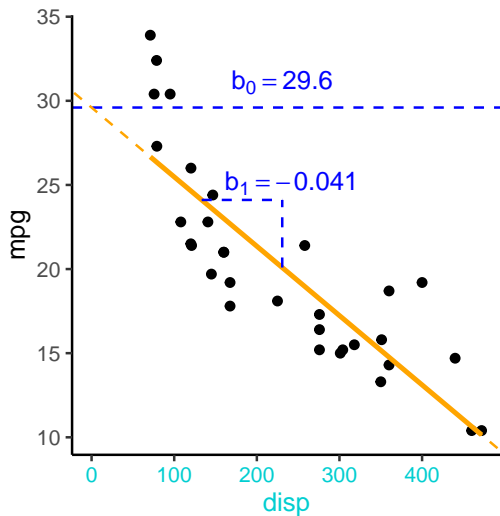


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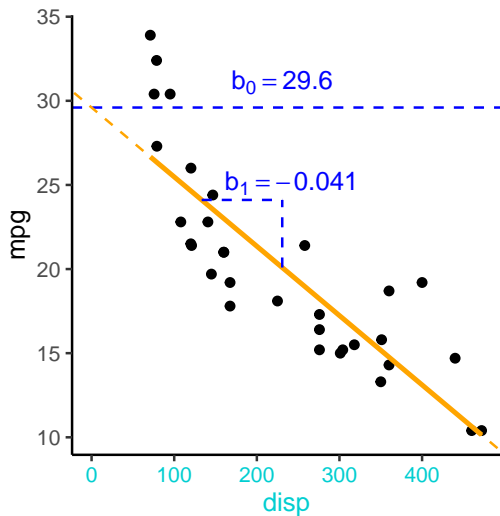


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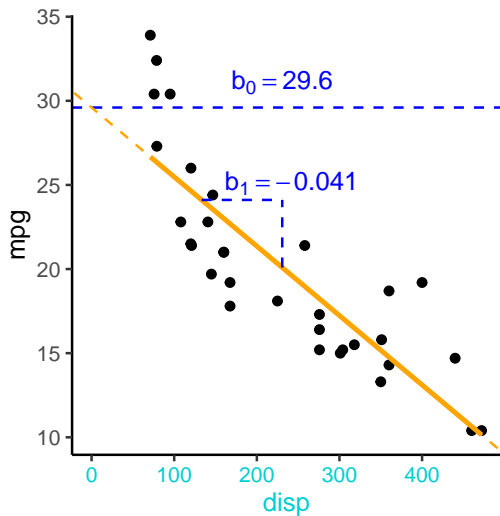


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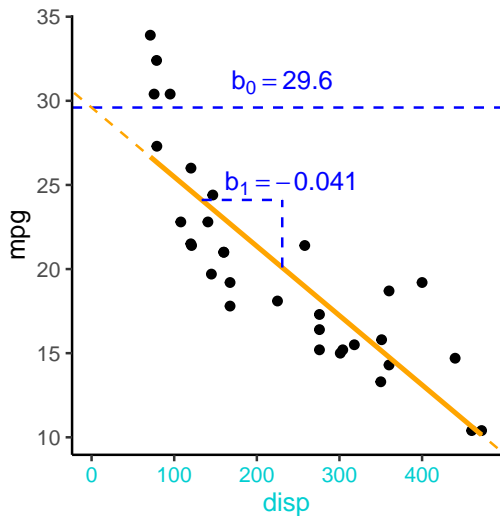


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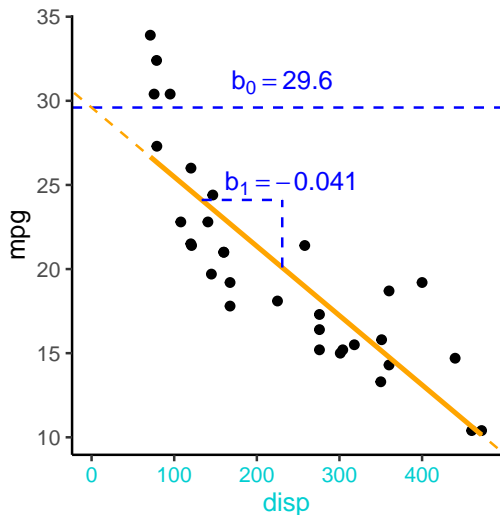


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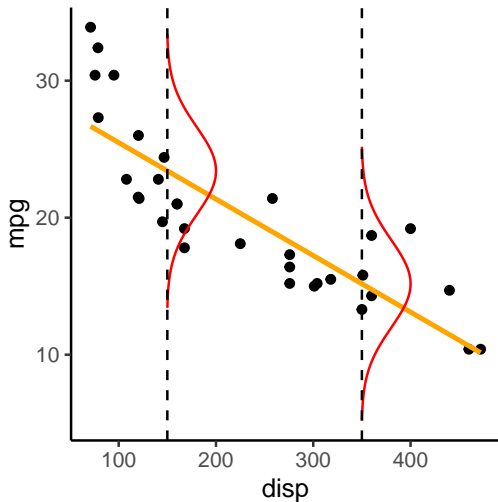
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- σ isn't displayed on the figure. Where is it?

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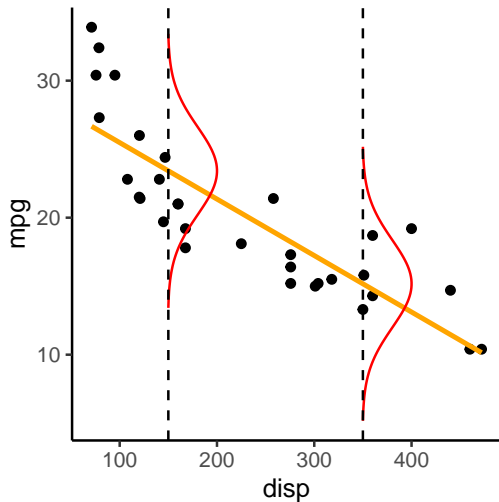
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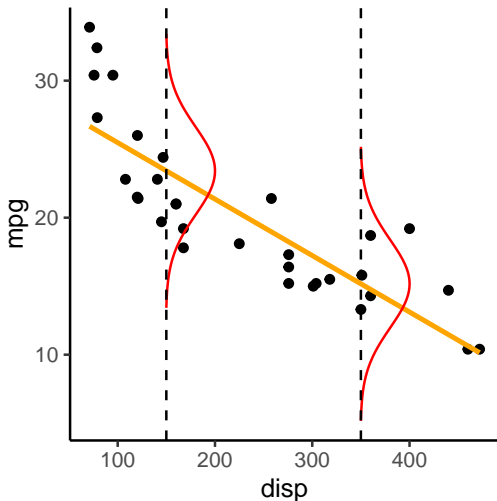
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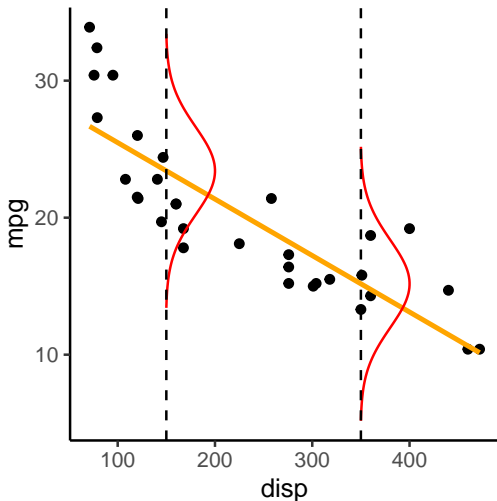
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- If you took a vertical slice at each part of the x-axis, the distribution would be *Normal*

How do I get R to fit this model?

lm is one of the main functions used for linear modeling:

```
#Formula= y ~ x, data = Name of the dataframe containing mpg & disp  
mod1 <- lm(mpg ~ disp, data = mtcars); summary(mod1)
```

```
##  
## Call:  
## lm(formula = mpg ~ disp, data = mtcars)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max   
## -4.8922 -2.2022 -0.9631  1.6272  7.2305   
##  
## Coefficients:  
##              Estimate Std. Error t value Pr(>|t|)      
## (Intercept) 29.599855   1.229720  24.070 < 2e-16 ***  
## disp        -0.041215   0.004712  -8.747 9.38e-10 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 3.251 on 30 degrees of freedom  
## Multiple R-squared:  0.7183, Adjusted R-squared:  0.709   
## F-statistic: 76.51 on 1 and 30 DF,  p-value: 9.38e-10
```

For a detailed breakdown of lm's output, click [here](#)

Simulate data

Now that we know how linear models work, we can simulate our own data:

```
#Parameters:
```

```
b0 <- 1 #Intercept
```

```
b1 <- 2 #Slope
```

```
sigma <- 3 #SD
```

```
#Make up some data:
```

```
x <- 0:30 #Predictor values
```

```
#Predicted y values
```

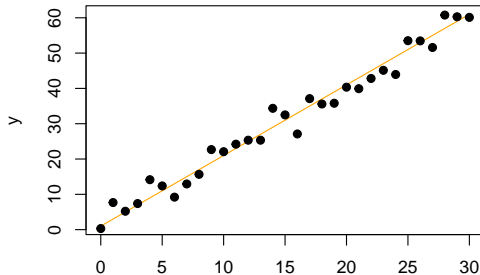
```
pred_y <- b0 + b1*x
```

```
#Add "noise" around pred_y
```

```
actual_y <- rnorm(n = length(pred_y),  
                 mean = pred_y,  
                 sd= sigma)
```

```
#Plot the data we just made
```

```
plot(x,pred_y,col='orange',pch=19,type='l',  
      ylab='y')  
points(x,actual_y,col='black',pch=19)
```



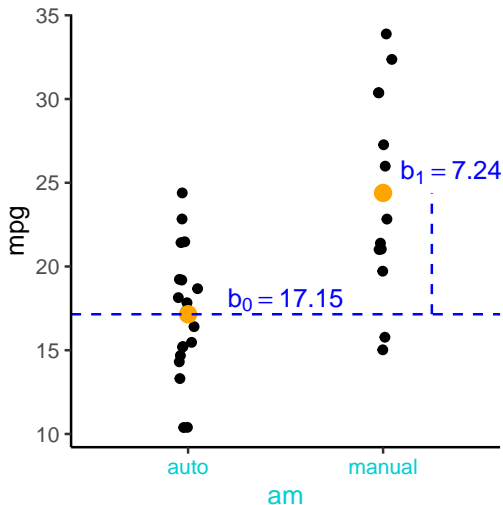
Fit a model from simulated data

How does R do at finding the coefficients? Remember: $b_0 = 1, b_1 = 2, \sigma = 3$

```
fakeDat <- data.frame(x = x, y = actual_y, pred = pred_y) #Simulated data in a dataframe
mod1sim <- lm(y ~ x, data = fakeDat); summary(mod1sim) #Fit model
```

```
##
## Call:
## lm(formula = y ~ x, data = fakeDat)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -5.7568 -1.7623 -0.2176  1.9419  5.3572
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.02974     1.00445   2.021  0.0526 .
## x            1.92670     0.05751  33.499 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.864 on 29 degrees of freedom
## Multiple R-squared:  0.9748, Adjusted R-squared:  0.9739
## F-statistic: 1122 on 1 and 29 DF, p-value: < 2.2e-16
```

What about categorical data?



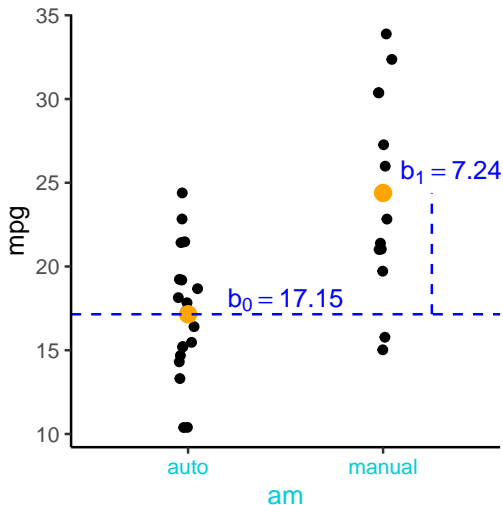
This uses *exactly the same* math!

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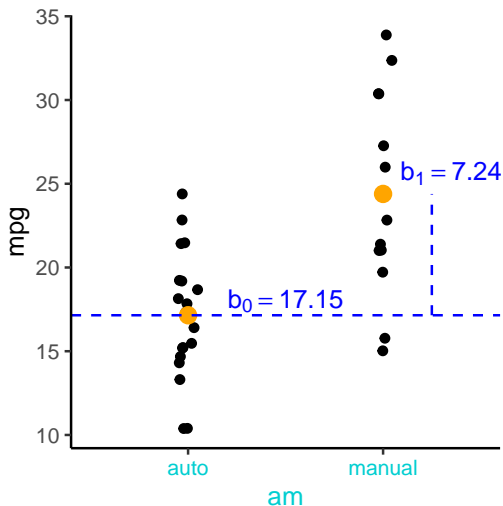
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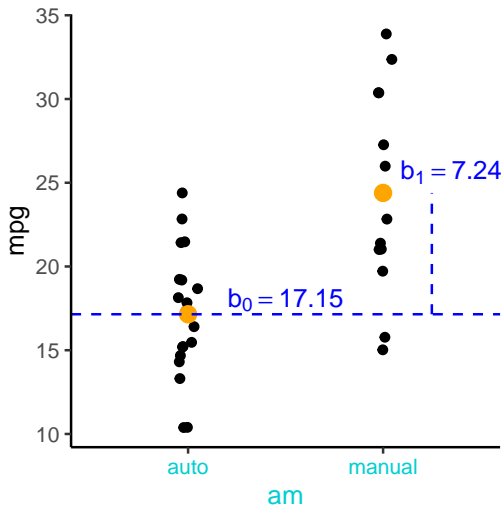
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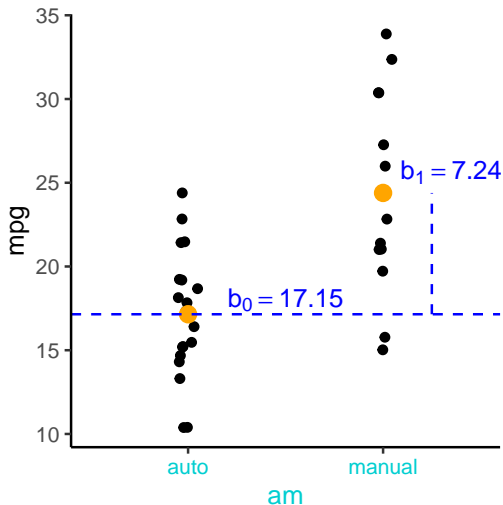
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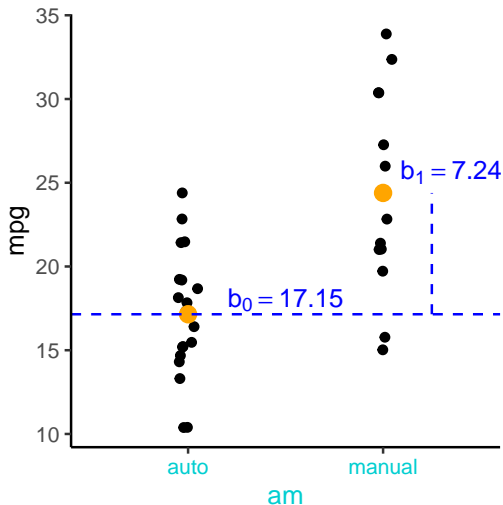
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- Where is σ ?

How do I get R to fit this model?

Syntax is exactly the same for this model

```
#Formula structure: y ~ x  
mod2 <- lm(mpg ~ am, #mpg depends on am  
           data = mtcars) #Name of the dataframe containing mpg & am  
summary(mod2)
```

```
##  
## Call:  
## lm(formula = mpg ~ am, data = mtcars)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max   
## -9.3923 -3.0923 -0.2974  3.2439  9.5077   
##  
## Coefficients:  
##              Estimate Std. Error t value Pr(>|t|)      
## (Intercept)   17.147      1.125   15.247 1.13e-15 ***  
## am              7.245      1.764    4.106 0.000285 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 4.902 on 30 degrees of freedom  
## Multiple R-squared:  0.3598, Adjusted R-squared:  0.3385   
## F-statistic: 16.86 on 1 and 30 DF,  p-value: 0.000285
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- Use `lm` to fit a model to the data you just simulated

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 - Useful command: `rep` (replicate)
 - e.g. `rep(x=c(0,1),each=10)`
 - Useful command: `rnorm` (generate normally-distributed data)
 - e.g. `rnorm(n=100,mean=0,sd=1)`
- Use `lm` to fit a model to the data you just simulated
 - How does R do at guessing your coefficients?

Modeling philosophy

All parametric models are approximating a **generative process**

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When we're fitting the model $\mathbb{1}_m(y \sim x)$, our implicit model of the process is:

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All parametric models are approximating a **generative process**

When we're fitting the model $\mathbb{1}_m(y \sim x)$, our implicit model of the process is:

- “I think that there is some kind of *average value* of **y**, that I'll call \hat{y} ” (parametric)
- “I think that \hat{y} responds to **x** in a **straight line**” (linear in parameters)

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All of these can be changed, as we'll see during the following weeks!

Modeling philosophy (cont.)

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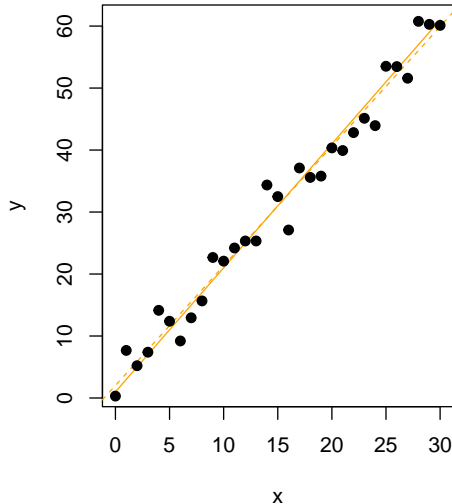
- When we gather data, we're seeing the outcome of this generative process, and trying to guess what the underlying process is.
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$$\hat{y} = b_0 + b_1 x$$

$$y \sim \text{Normal}(\hat{y}, \sigma)$$

$b_0 = 1, b_1 = 2, \sigma = 3$: "True" values

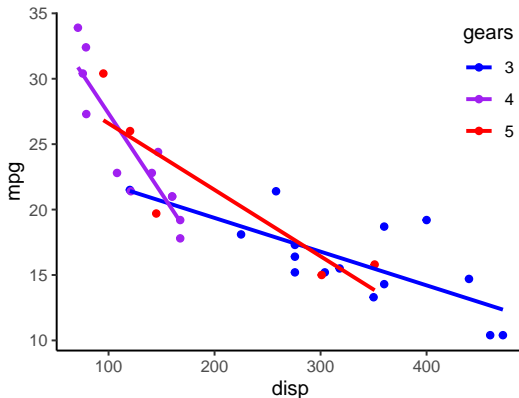
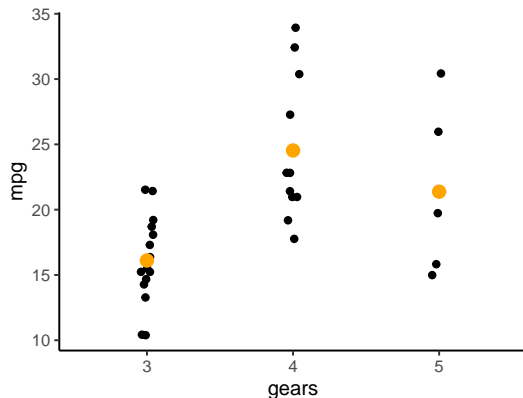
$\hat{b}_0 = 2.0, \hat{b}_1 = 1.9, \hat{\sigma} = 2.9$: Estimated values



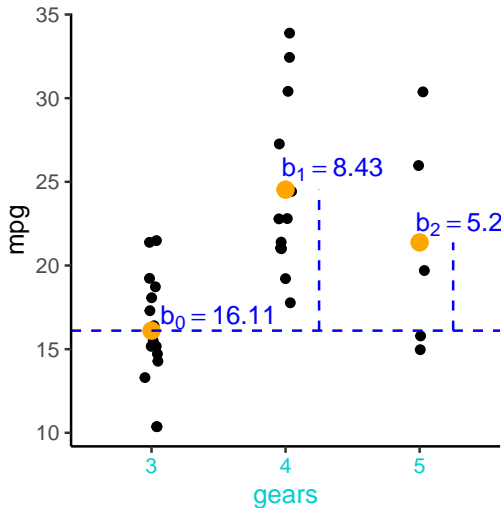
Part 2: More bells and whistles

Motivation

- *I have 2+ groups of data, and I want to know whether the means are different*
- *I have 2+ groups of bivariate data, and I want to know whether the relationships differ between groups*



Categorical data, 3 categories



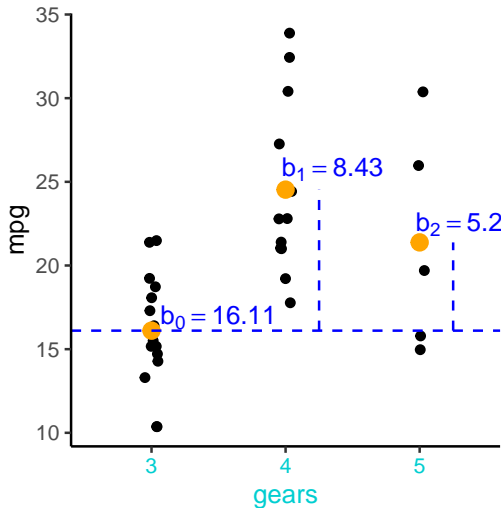
The more factor levels, the more coefficients:

$$\hat{mpg} = b_0 + b_1 \text{gears}_4 + b_2 \text{gears}_5$$

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- mpg is the thing you're interested in predicting

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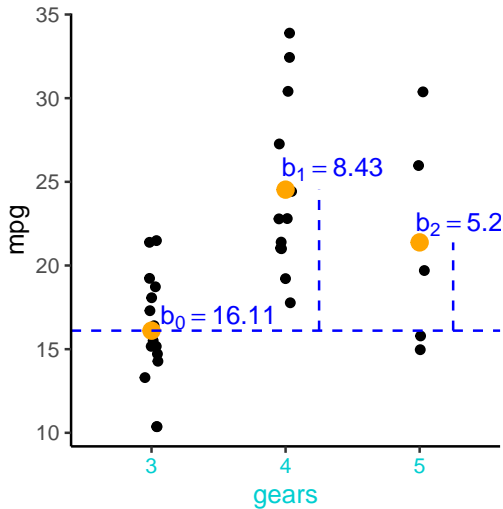
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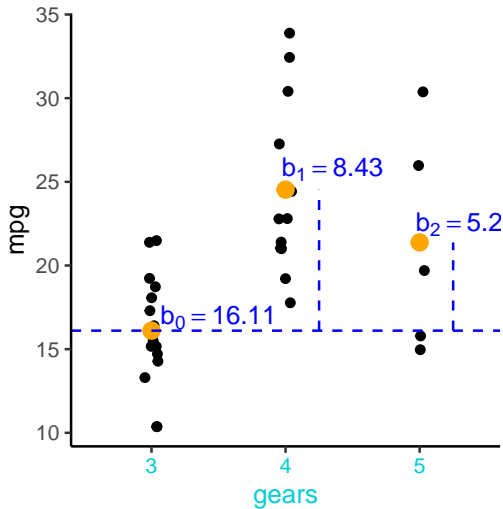
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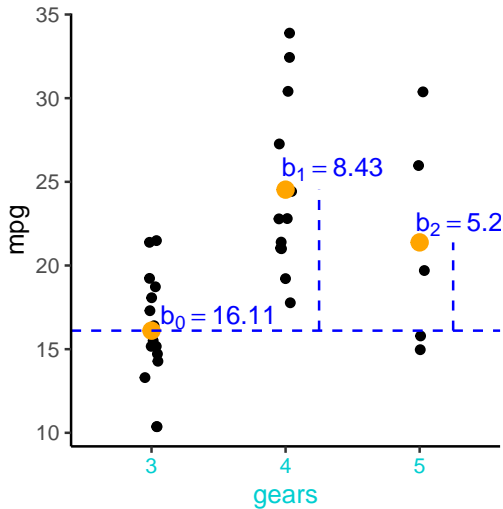
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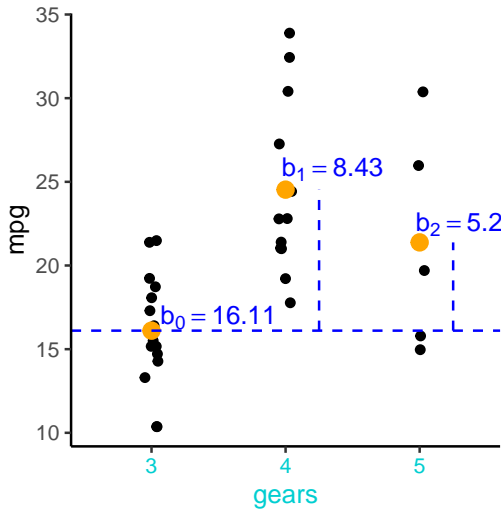
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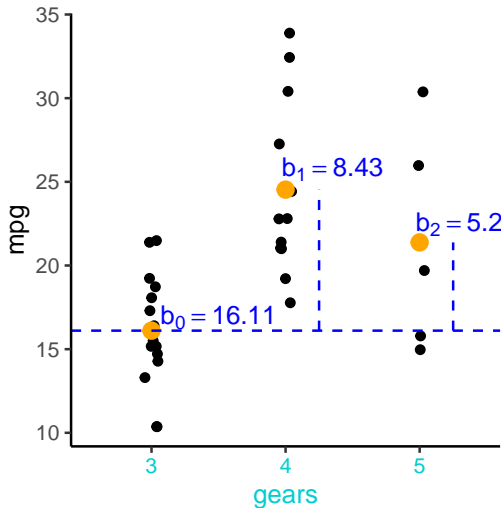
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 - set of 0s and 1s
 - gears_4 = "is this data point from a 4-gear car?"
- b_0 = *intercept* (first level of gear factor)
- $[b_1, b_2]$ = are *coefficients* for $gears$

How do I get R to fit this model?

```
#Formula structure: y ~ x
mod1 <- lm(mpg ~ factor(gear), #mpg depends on gears
           data = mtcars) #Name of the dataframe containing mpg & gears
summary(mod1)
```

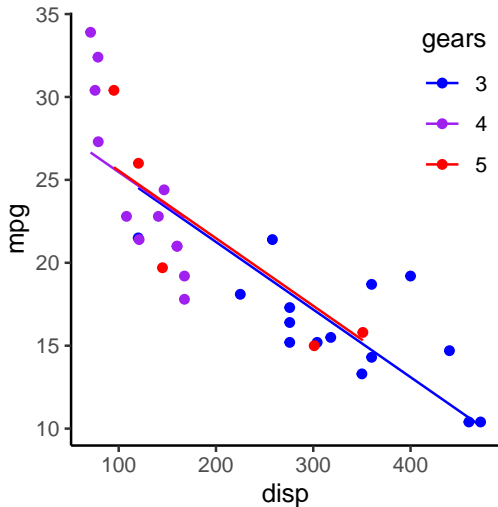
```
##
## Call:
## lm(formula = mpg ~ factor(gear), data = mtcars)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -6.7333 -3.2333 -0.9067  2.8483  9.3667
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    16.107      1.216   13.250 7.87e-14 ***
## factor(gear)4     8.427      1.823    4.621 7.26e-05 ***
## factor(gear)5     5.273      2.431    2.169  0.0384 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 4.708 on 29 degrees of freedom
## Multiple R-squared:  0.4292, Adjusted R-squared:  0.3898
## F-statistic: 10.9 on 2 and 29 DF, p-value: 0.0002948
```

Dummy variables

```
mod1Matrix <- model.matrix(mod1) #Get model matrix (columns used to predict mpg)  
head(mod1Matrix,20) #Show first 20 rows of model matrix
```

##	(Intercept)	factor(gear)4	factor(gear)5
## Mazda RX4	1	1	0
## Mazda RX4 Wag	1	1	0
## Datsun 710	1	1	0
## Hornet 4 Drive	1	0	0
## Hornet Sportabout	1	0	0
## Valiant	1	0	0
## Duster 360	1	0	0
## Merc 240D	1	1	0
## Merc 230	1	1	0
## Merc 280	1	1	0
## Merc 280C	1	1	0
## Merc 450SE	1	0	0
## Merc 450SL	1	0	0
## Merc 450SLC	1	0	0
## Cadillac Fleetwood	1	0	0
## Lincoln Continental	1	0	0
## Chrysler Imperial	1	0	0
## Fiat 128	1	1	0
## Honda Civic	1	1	0
## Toyota Corolla	1	1	0

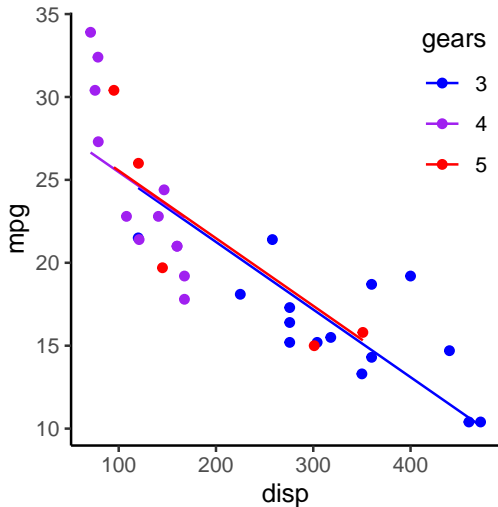
What if 2 things are *both* important?



$$\begin{aligned} \hat{mpg} &= b_0 + b_1 disp \\ &\quad + b_2 gears_4 + b_3 gears_5 \\ mpg &\sim Normal(\hat{mpg}, \sigma) \end{aligned}$$

- Suppose that both *disp* and *gears* are important for predicting *mpg*?

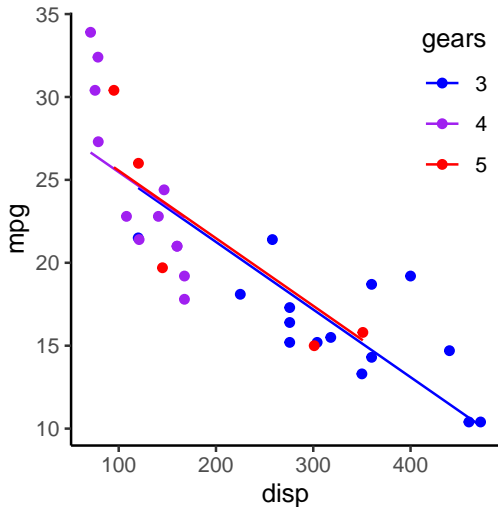
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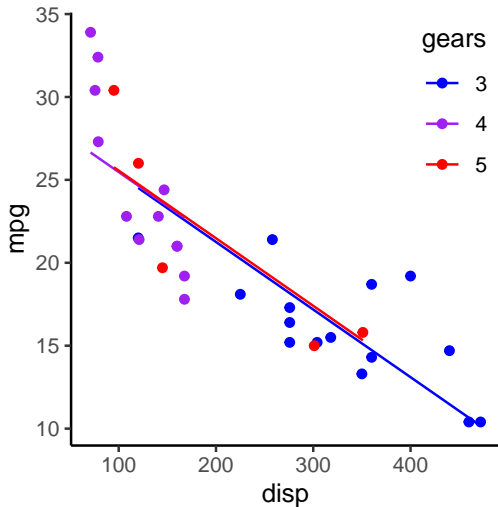
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- This is very similar to the last example, except that now we've added *disp*
- *gears* now changes the intercepts, while *disp* changes the overall slope
- Now that both variables are included, does it look like *gear* is very important?

How do I get R to fit this model?

#mpg depends on disp and gears

```
mod2 <- lm(mpg ~ disp+factor(gear), data = mtcars)
summary(mod2)
```

```
##
## Call:
## lm(formula = mpg ~ disp + factor(gear), data = mtcars)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -4.9155 -2.1892 -0.9054  1.5790  7.2498
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   29.411183    2.627966   11.192 7.58e-12 ***
## disp         -0.040774    0.007601   -5.364 1.03e-05 ***
## factor(gear)4  0.138017    2.021332    0.068  0.946
## factor(gear)5  0.224712    1.976090    0.114  0.910
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 3.365 on 28 degrees of freedom
## Multiple R-squared:  0.7185, Adjusted R-squared:  0.6883
## F-statistic: 23.82 on 3 and 28 DF,  p-value: 7.31e-08
```

Dummy variables

```
mod2Matrix <- model.matrix(mod2) #Get model matrix (columns used to predict mpg)
head(mod2Matrix,20) #Show first 20 rows of model matrix
```

```
##                (Intercept)  disp factor(gear)4 factor(gear)5
## Mazda RX4                1 160.0                1            0
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- `lm` model input:

```
model1 <- lm(y ~ x1 + x2 + ..., data = myDataFrame)
```

```
summary(model1)
```

Interlude: problems with plotting raw data

- Say that I've fit the following model:

```
mpg ~ disp + factor(gear)
```


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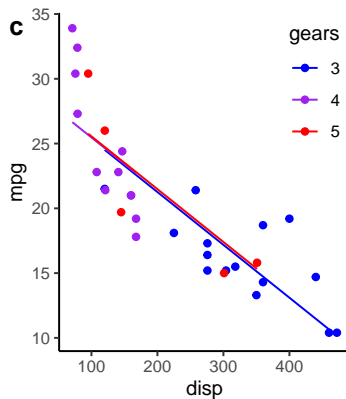
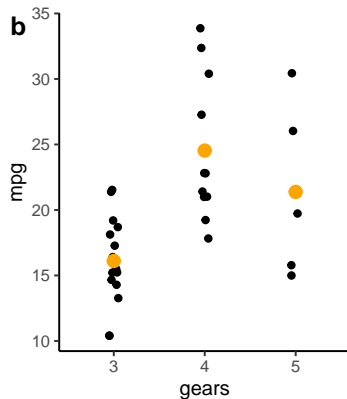
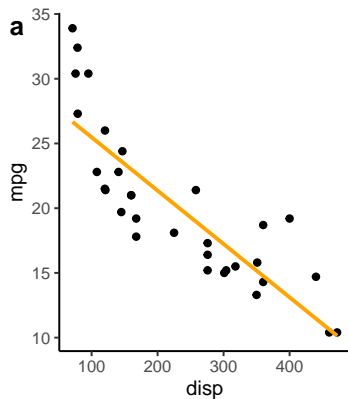
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Interlude: problems with plotting raw data

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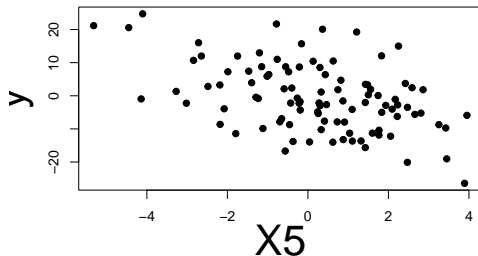
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Incorrect example, using raw data:

```
#Fit model with 5 variables (all important)
simMod <- lm(y~X1+X2+X3+X4+X5,data=pred)
#Plot x5 and y
plot(y~X5,data=pred,pch=19,cex.lab=3)
```



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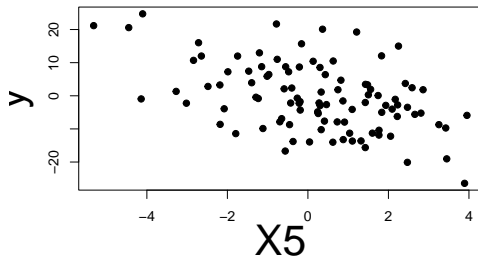
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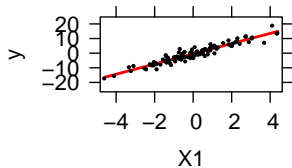


The effect of $X5$ is actually **very** strong ($p < 0.0001$), but it doesn't look like it from this plot!

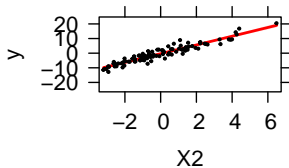
Partial effects plots - using *effects*

```
library(effects) #Load effects package
simModEff <- predictorEffects(simMod,partial.residuals=TRUE) #Calculate partial effects
#Plot partial effects
plot(simModEff,lines=list(col='red'), partial.residuals=list(pch=19,col='black',cex=0.25))
```

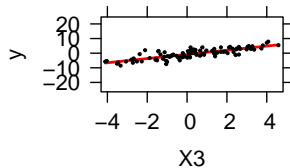
X1 predictor effect plot



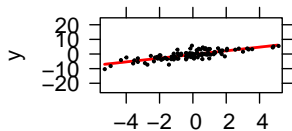
X2 predictor effect plot



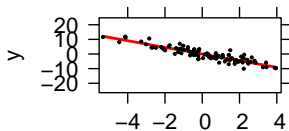
X3 predictor effect plot



X4 predictor effect plot



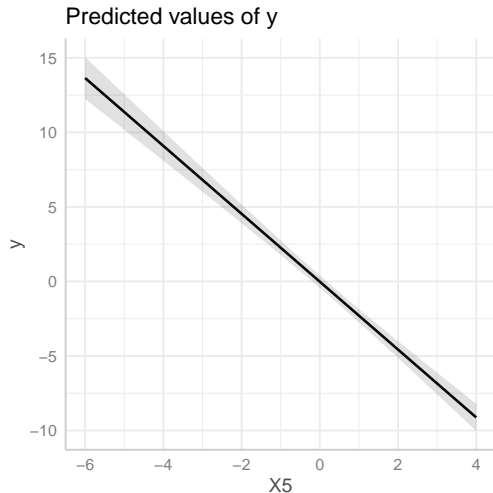
X5 predictor effect plot



Partial effects plots - using *ggeffects*

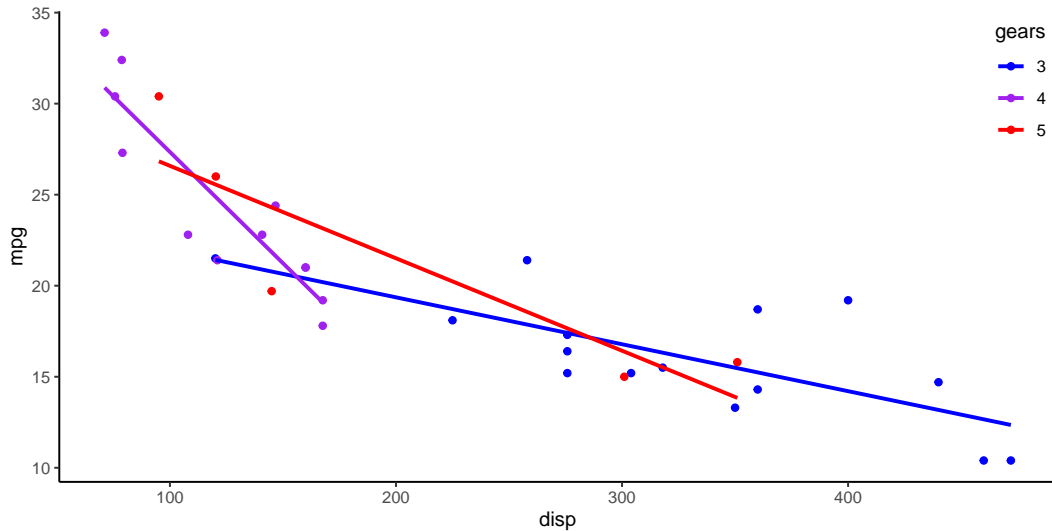
```
#Load ggeffects package  
library(ggeffects)  
  
#Calculate partial effects for X5  
simModEff2 <- ggeffect(simMod, terms=c('X5'))  
  
#Plot the effect of X5  
plot(simModEff2)
```

- You can also turn ggeffect objects into a dataframe and make your own custom plots

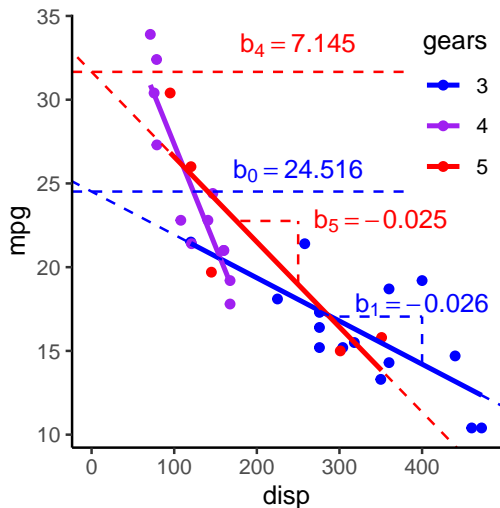


Interactions

What if the slopes *and* intercepts differ between groups?



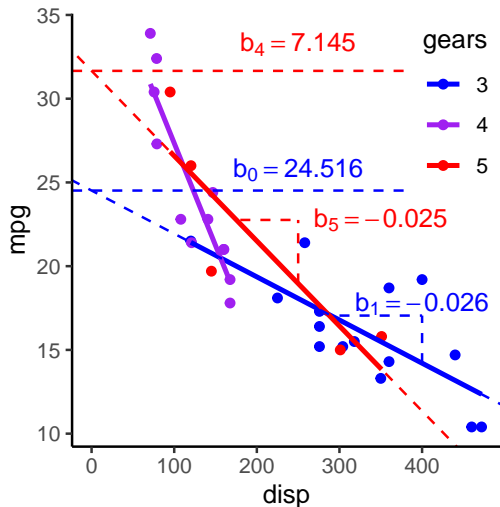
Interactions



$$\begin{aligned} \hat{mpg} = & b_0 + b_1 disp \\ & + b_2 gears_4 + b_3 gears_5 \\ & + b_4 (disp \times gears_4) \\ & + b_5 (disp \times gears_5) \\ mpg \sim & Normal(\hat{mpg}, \sigma) \end{aligned}$$

- Interactions occur when predictors are *multiplied*

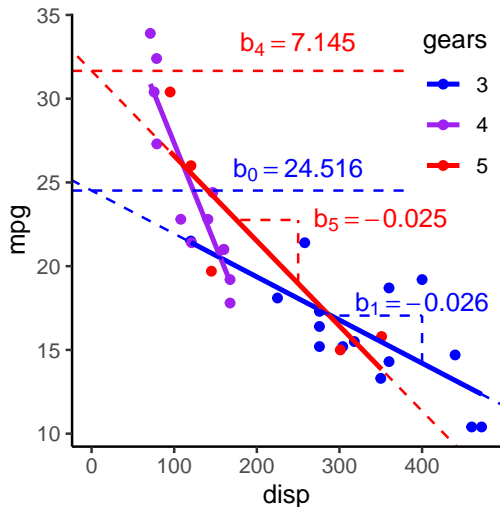
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- Interactions occur when predictors are *multiplied*
- In this case, *disp* is multiplied by *gears₄* and *gears₅*
- *gears* now changes the intercept and the slope of the relationship between *mpg* and *disp*

How do I get R to fit this model?

```
#mpg depends on disp interacted (*) with gears
mod2 <- lm(mpg ~ disp*factor(gear), data = mtcars)
summary(mod2)
```

```
##
## Call:
## lm(formula = mpg ~ disp * factor(gear), data = mtcars)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -4.5986 -1.5990 -0.0143  1.6329  4.9926
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    24.51556     2.462431   9.956 2.32e-10 ***
## disp          -0.025770     0.007265  -3.547 0.001505 **
## factor(gear)4    15.051963     3.558043   4.230 0.000256 ***
## factor(gear)5     7.145380     3.535913   2.021 0.053711 .
## disp:factor(gear)4 -0.096442     0.021261  -4.536 0.000114 ***
## disp:factor(gear)5 -0.025005     0.013320  -1.877 0.071742 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.579 on 26 degrees of freedom
## Multiple R-squared:  0.8465, Adjusted R-squared:  0.817
## F-statistic: 28.67 on 5 and 26 DF,  p-value: 8.452e-10
```

Beware of fitting too many interactions, or else the *Bilbo effect* occurs!

Dummy variables

```
mod2Matrix <- model.matrix(mod2) #Get model matrix (columns used to predict mpg)  
colnames(mod2Matrix) <- gsub('(factor\\(|\\|\\))', '', colnames(mod2Matrix)) #Shrink column headers  
head(mod2Matrix, 20) #Show first 20 rows of model matrix
```

Third challenge

- Make some plots of your model results using `ggeffects`

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 - `lm (y ~ X1 * X2 * X3)`: Full model (everything interacts)
 - `lm (y ~ X1 + X2 + X3 + X2:X3)`: interaction only between X2 and X3

Part 3: Models behaving badly

Motivation

Are my model results reliable?

- Residual checks

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- Transformations

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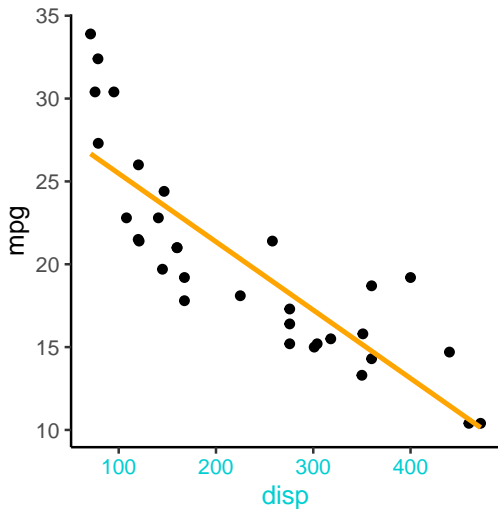
- Residual checks
- Transformations
- Collinearity

Motivation

Are my model results reliable?

- Residual checks
- Transformations
- Collinearity
- How much stuff should I put into my model?

Assumptions of linear regression

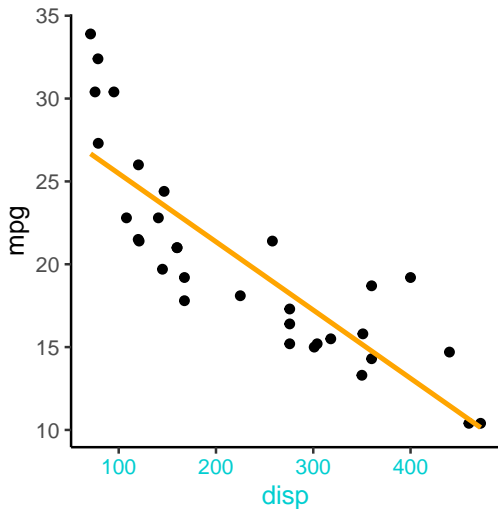


$$\hat{mpg} = b_0 + b_1 disp$$

$$mpg \sim Normal(\hat{mpg}, \sigma)$$

There are 3 main assumptions to this model:

Assumptions of linear regression



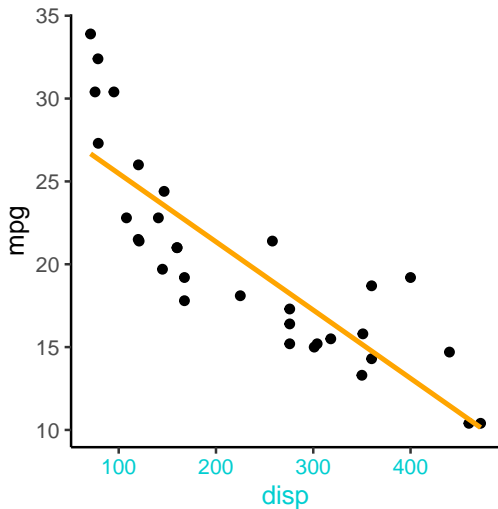
$$\hat{mpg} = b_0 + b_1 disp$$

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- 1 The relationship between *disp* and *mpg* is **linear**

Assumptions of linear regression



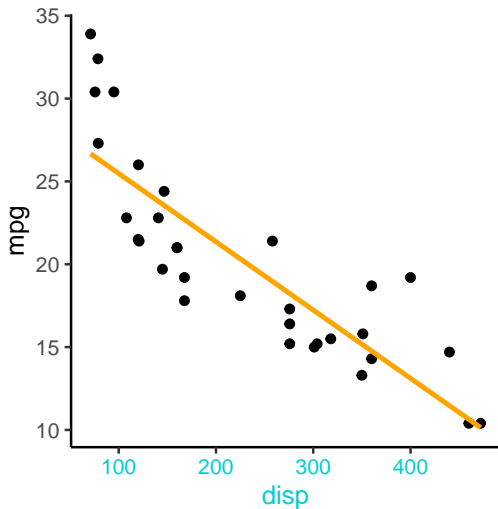
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- ② *mpg* (the data) is **Normally distributed** around \hat{mpg} (the line)

Assumptions of linear regression



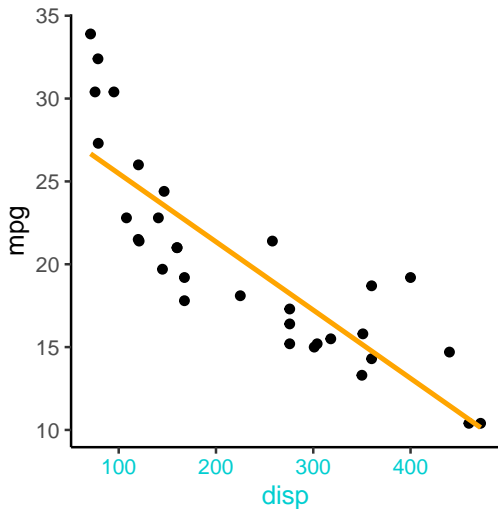
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Assumptions of linear regression



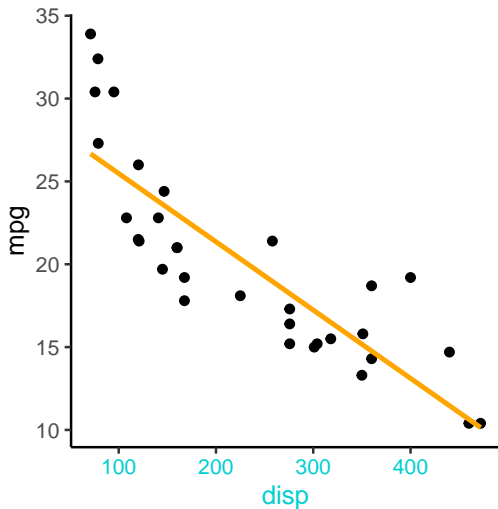
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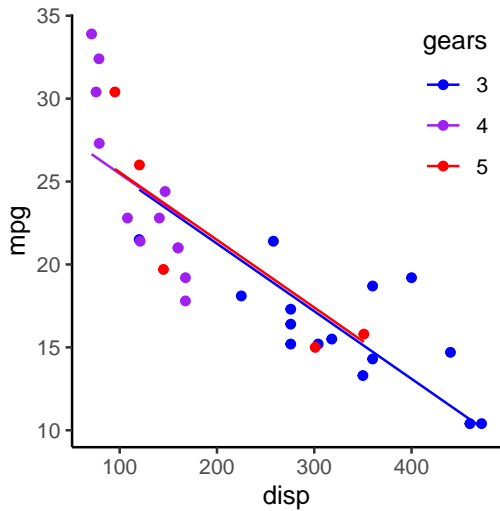
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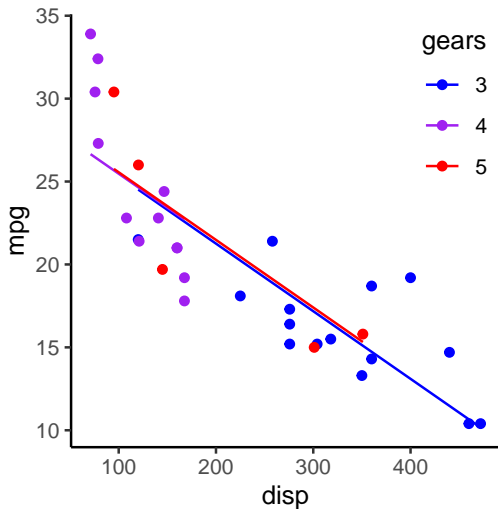
This is pretty easy to see if you only have 1 variable, but...

What if I have many variables?



- Difficult to see if the assumptions are met

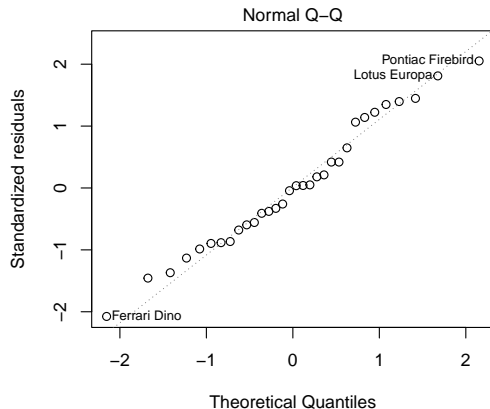
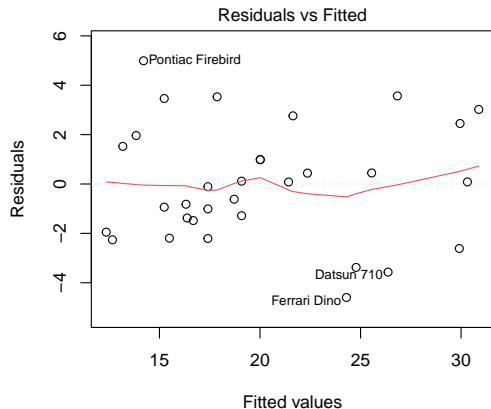
What if I have many variables?



- Difficult to see if the assumptions are met
- In general, we use **residual plots** or **simulation** to assess whether model assumptions are met

Solution: residual checks

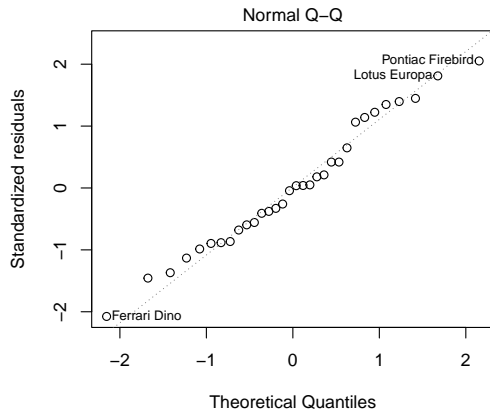
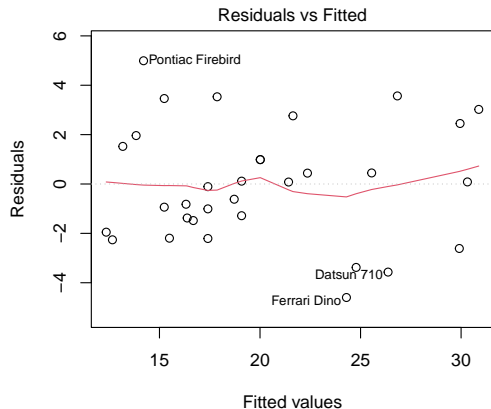
Some common ways of checking the assumptions: **residual plots**



- Points in Plot 1 should show *no pattern* (shotgun blast)

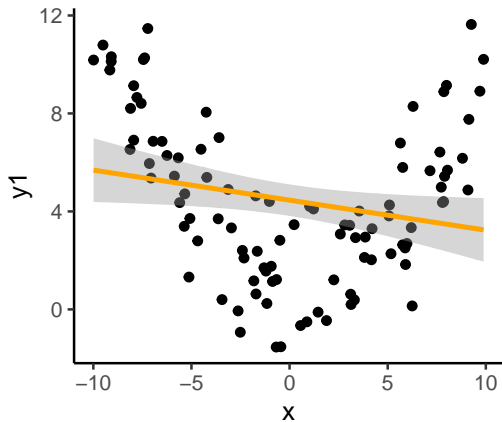
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Some common ways of checking the assumptions: **residual plots**



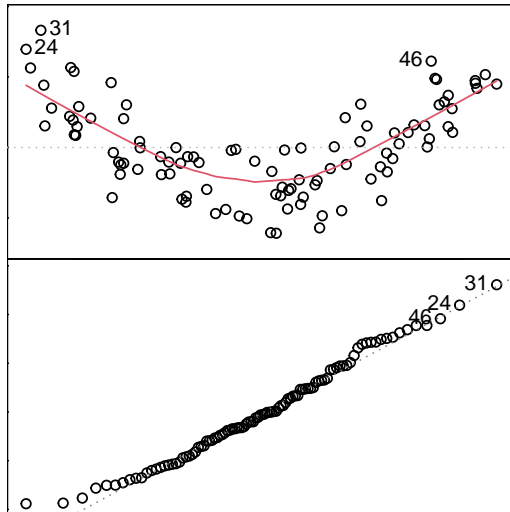
- Points in Plot 1 should show *no pattern* (shotgun blast)
- Points in Plot 2 should be *roughly* on top of the 1:1 line

Problem 1: Non-linear relationship

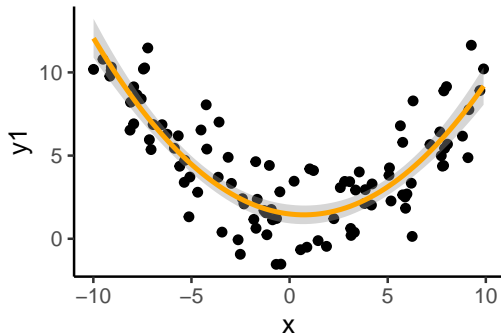


Model: `lm(y1~x,data=d1)`

- y_1 clearly follows a U-shaped relationship, not a linear one

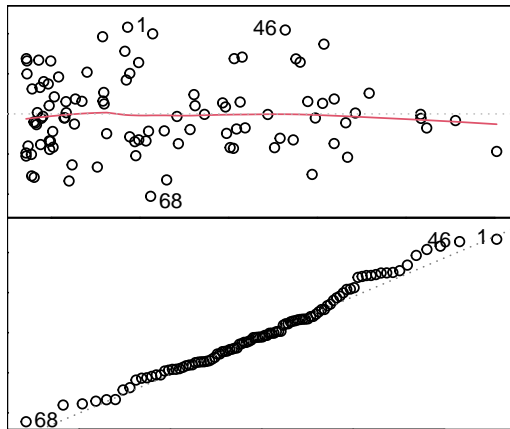


Solution: transform predictors



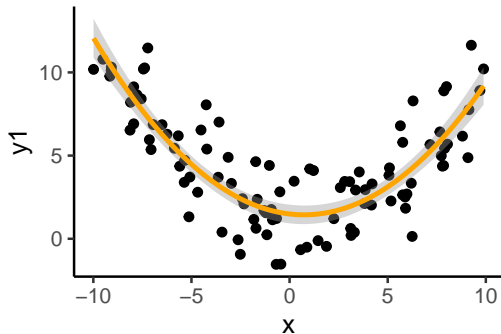
Model: `lm(y1~poly(x,2),data=d1)`

- *log* and *square-root* transformations are common



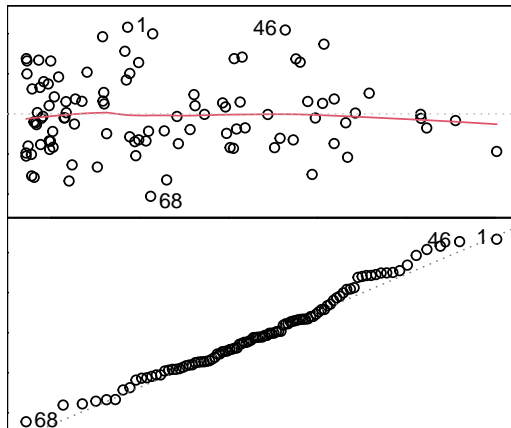
- Warning: Polynomials can do weird things; consider whether this is biologically reasonable!

Solution: transform predictors



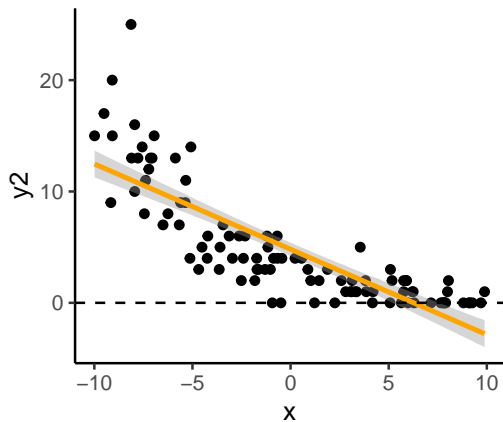
Model: `lm(y1~poly(x,2),data=d1)`

- *log* and *square-root* transformations are common
- Can also use *additive* (wiggly) models



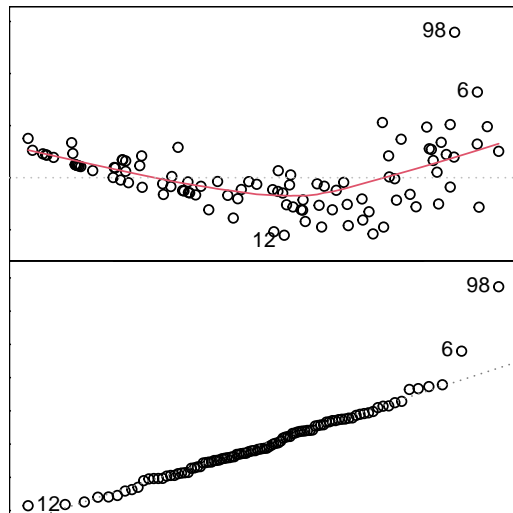
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Problem 2a: Non-normal response

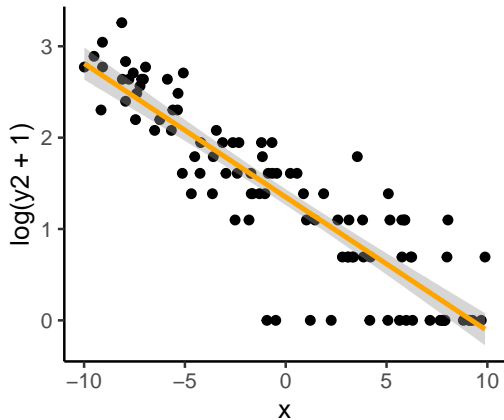


Model: `lm(y2~x,data=d1)`

- y_2 is count data (integers ≥ 0).
Very common in ecological data.

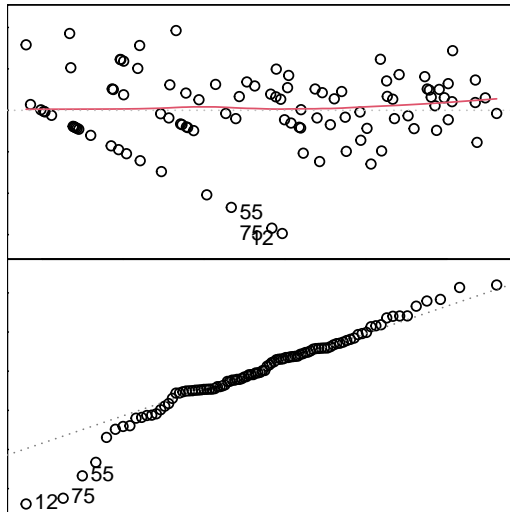


Solution: transform data to meet assumptions

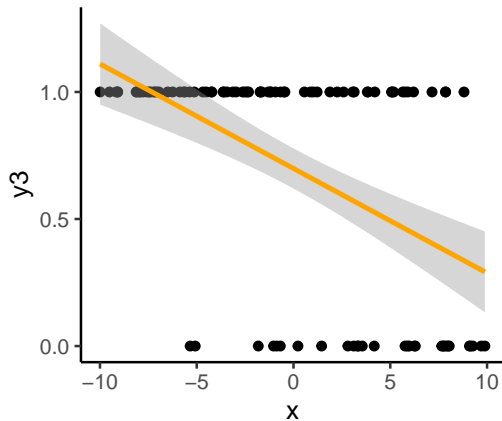


Model: `lm(log(y2+1)~x,data=d1)`

- Square-root transformations are also common

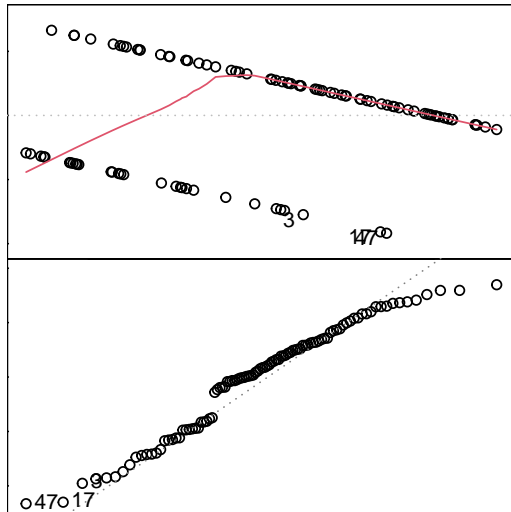


Problem 2b: Non-normal response

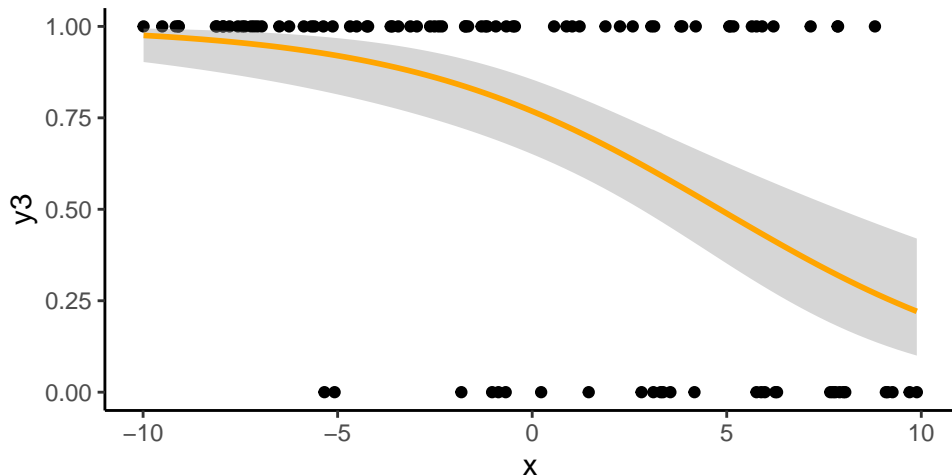


Model: `lm(y3~x,data=d1)`

- y_3 is binomial data (success/failure).
Very common in ecological data.

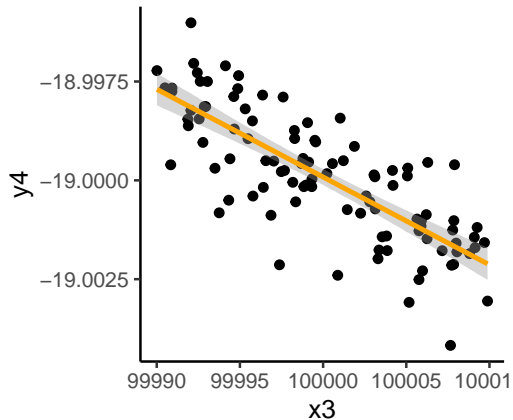


Solution: use a Generalized Linear Model (GLM)



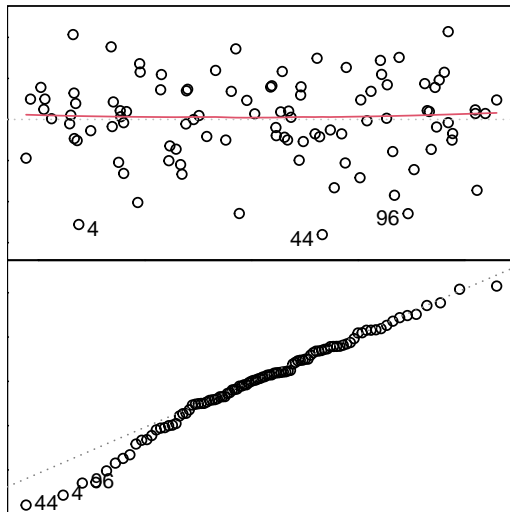
- This is a topic for another lecture. Hold tight!

Problem: variables are on different scales

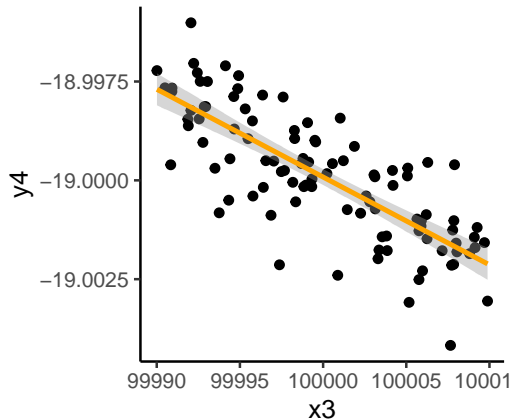


Model: `lm(y4~x3,data=d1)`

- y_4 is tiny, while x_3 is huge

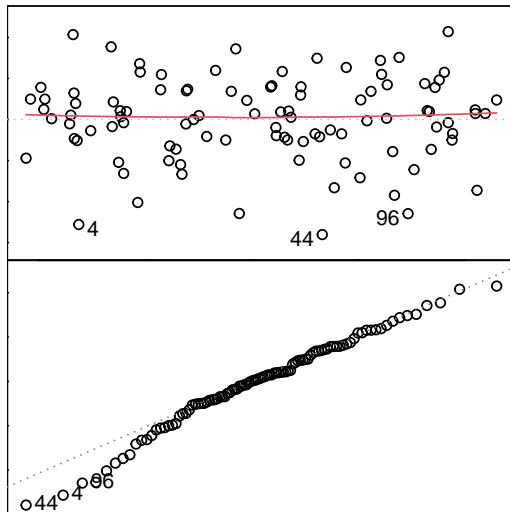


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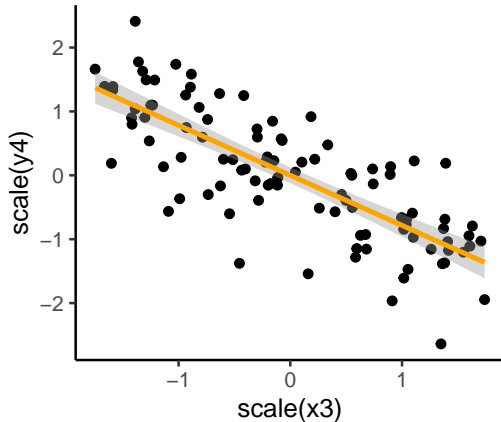


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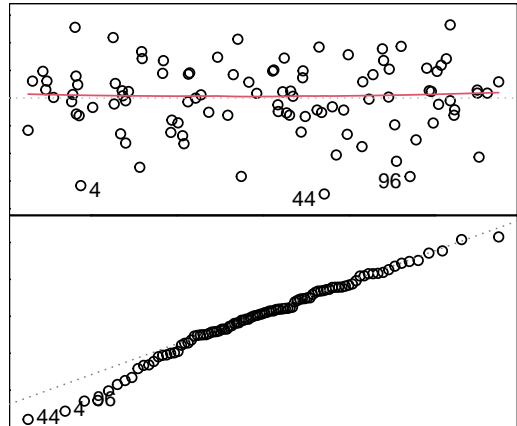
- y_4 is tiny, while x_3 is huge
- OK for now, but can cause problems when fitting other models



Solution: scale data/predictors before fitting

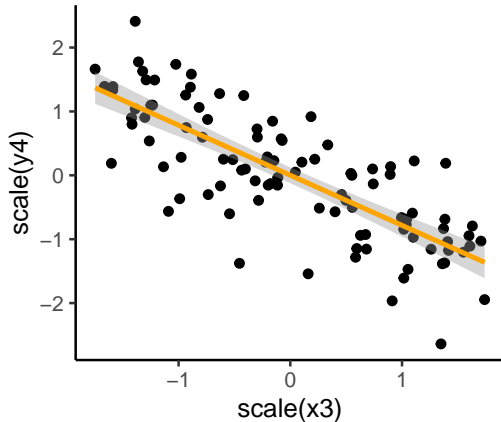


```
#Subtracts mean, divides by SD  
d1$s.y4 <- scale(y4)  
d1$s.x3 <- scale(x3)  
lm(s.y4~s.x3,data=d1) #Refit
```

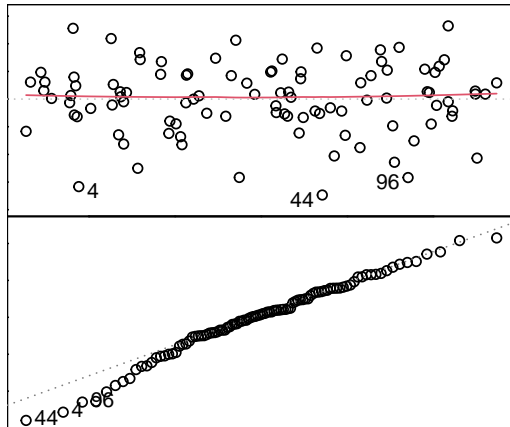


- Residuals are the same as before

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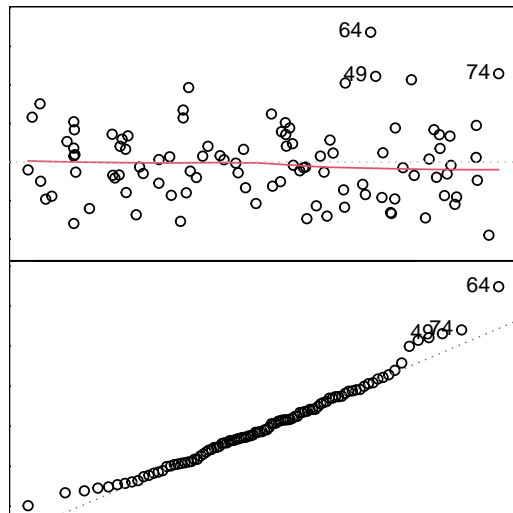


- Residuals are the same as before
- Coefficients are now related to *scaled* data and predictor

But wait... there's more (assumptions)!

One more assumption:

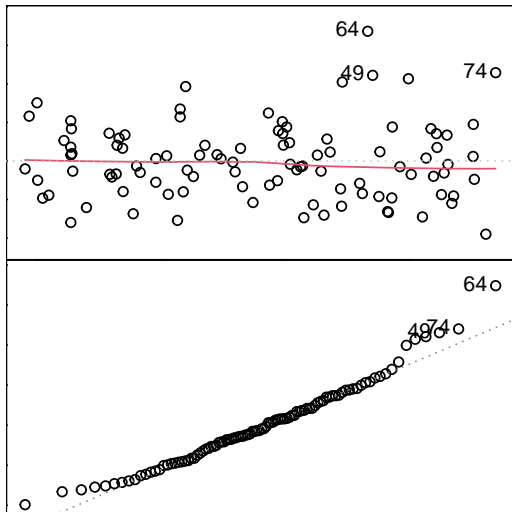
- ④ If you have 2+ predictors in your model, the predictors are not related to each other



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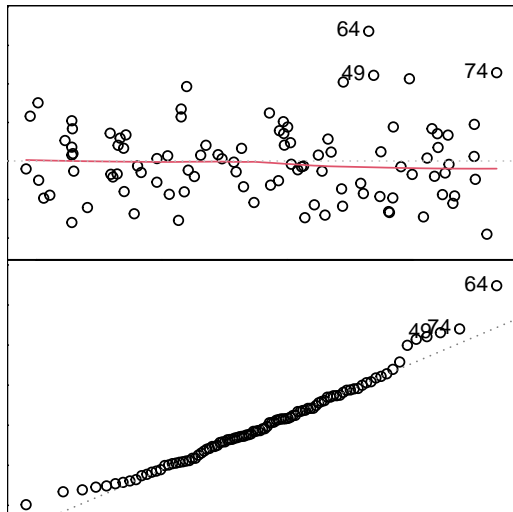
- ④ If you have 2+ predictors in your model, the predictors are not related to each other
- Say we have 2 predictors, x and x_2 :
`lm(y0~x+x2,data=d1)`



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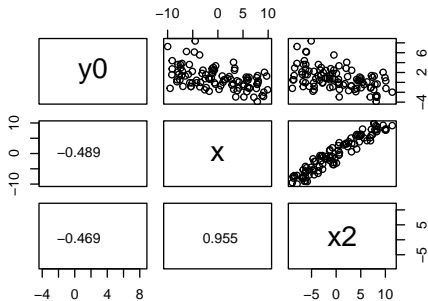
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- ④ If you have 2+ predictors in your model, the predictors are not related to each other
- Say we have 2 predictors, x and x_2 :
`lm(y0~x+x2,data=d1)`
- Model fits, and residuals look OK, but there's trouble ahead!



Uh oh! Collinearity!

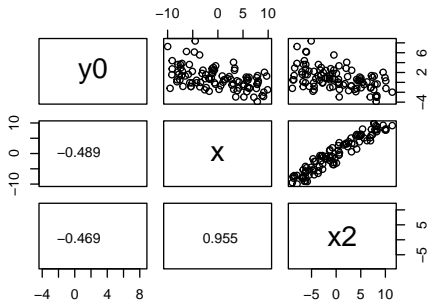
```
#Function to print correlation (r) value  
corText <- function(x,y){  
  text(0.5,0.5,round(cor(x,y),3))}  
#Pairplot of y0, x, and x2  
pairs(d1[,c('y0','x','x2')],  
      lower.panel=corText)
```



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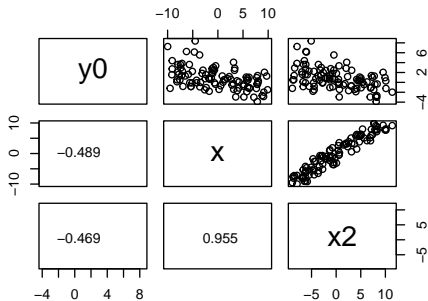
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- x and x^2 mean basically the same thing!



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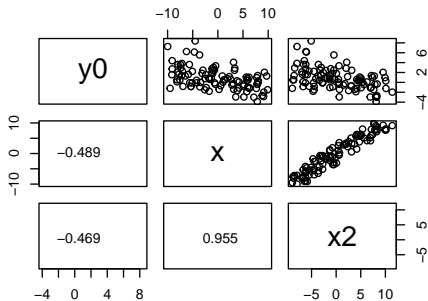
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- Also revealed using variance-inflation factors (VIFs):

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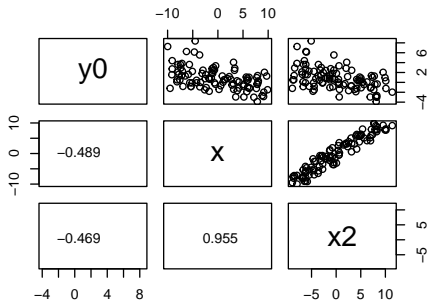
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```



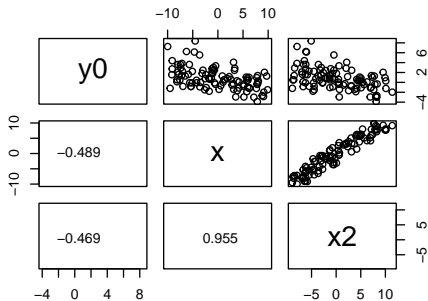
- x and x2 mean basically the same thing!
- Also revealed using variance-inflation factors (VIFs):

```
library(car)
#VIF scores:
# 1 = no problem
# 1-5 = some problems
# 5+ = big problems!
vif(m2)
```

```
##           x           x2
## 11.31812 11.31812
```

Uh oh! Collinearity!

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#Function to print correlation (r) value
corText <- function(x,y){
  text(0.5,0.5,round(cor(x,y),3))}
#Pairplot of y0, x, and x2
pairs(d1[,c('y0','x','x2')],
      lower.panel=corText)
```



- x and $x2$ mean basically the same thing!
- Also revealed using variance-inflation factors (VIFs):

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library(car)
#VIF scores:
# 1 = no problem
# 1-5 = some problems
# 5+ = big problems!
vif(m2)
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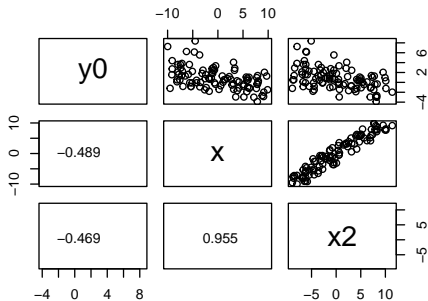
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Causes:

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- Data: other underlying (possibly unmeasured) relationships

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#Correct model

```
m1 <- lm(y0~x,data=d1)
```

	Estimate	Std. Error	Pr(> t)
(Intercept)	0.7851936	0.1943002	0.0001059
x	-0.1900346	0.0342596	0.0000002

#Incorrect model

```
m2 <- lm(y0~x+x2,data=d1)
```

	Estimate	Std. Error	Pr(> t)
(Intercept)	0.7860300	0.1955770	0.0001155
x	-0.1812556	0.1158464	0.1209288
x2	-0.0094931	0.1196074	0.9369028

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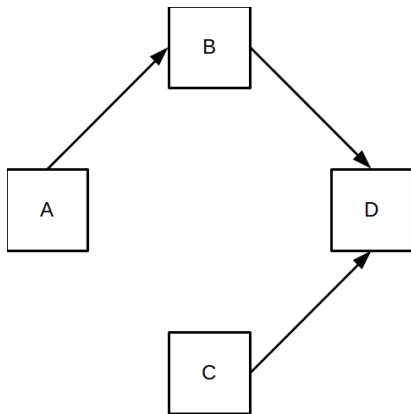
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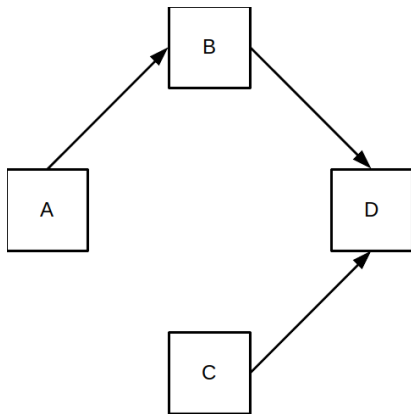
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- Simple graphical model, where the effect of A on D is *mediated* by B.
- “Correct” lm model of D: $\text{lm}(D \sim B + C)$

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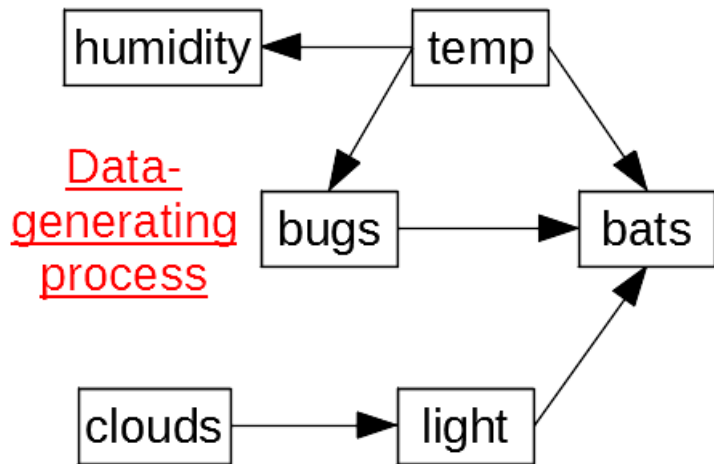
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- Formulate a causal model that seems reasonable
 - Draw it out on paper/in PowerPoint using flow diagrams
- Fit an `lm` model of bats from your causal model, check the assumptions, and update as necessary

Here's the answer



This is the **true** process that generated the data. Model for bats should look like:

```
lm(log(bats+0.1)~poly(temp,2)+light+bugs,data=dat)
```

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 - ③ If a relationship is unclear, use a *double-headed arrow*