

Exploratory and Confirmatory Data Analysis

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Learning Objectives

- Some exploratory situations
 - I have a hypothesis but I'm not quite sure how to test it with the variables I have
 - I think some variables could be relevant to a DV but I'm not sure which ones
- Working through an example

The Issues:

- We're often interested in the relationship between variables but don't have clear predictions about how they're related
 - For example, I might be interested in why some tweets go viral and others don't
- The number of possible predictors related to this question is huge and it's not obvious which ones will be most important
 - Includes a photo? Humor? Many, many possible predictors

The Issues:

- Sometimes I might have a hypothesis, but it could be tested in multiple ways and I'm not sure how best to test it.
- For example, I think a tweet including a photo will be retweeted more. What kind of photo though? Any photo? Happy photos?

Exploratory Analyses

- The context I've describe above is a case of exploratory analysis.
- Exploratory analyses can take many forms, but they share in common the fact that you, the researcher, don't have extremely specific predictions about the relationship between your independent variable and your dependent variable
- The exploratory phase of data analysis is a great way to learn a lot about your data, but you also need to be on-guard that you don't think you've detected signal when you've actually detected noise.
 - More on this in a bit

Exploratory analyses done wrong

- "You cannot find your starting hypothesis in your final results. It makes the stats go all wonky." - Ben Goldacre
- If you treat an exploratory result as if you had that hypothesis from the start, then it can cause problems. You will trick yourself.

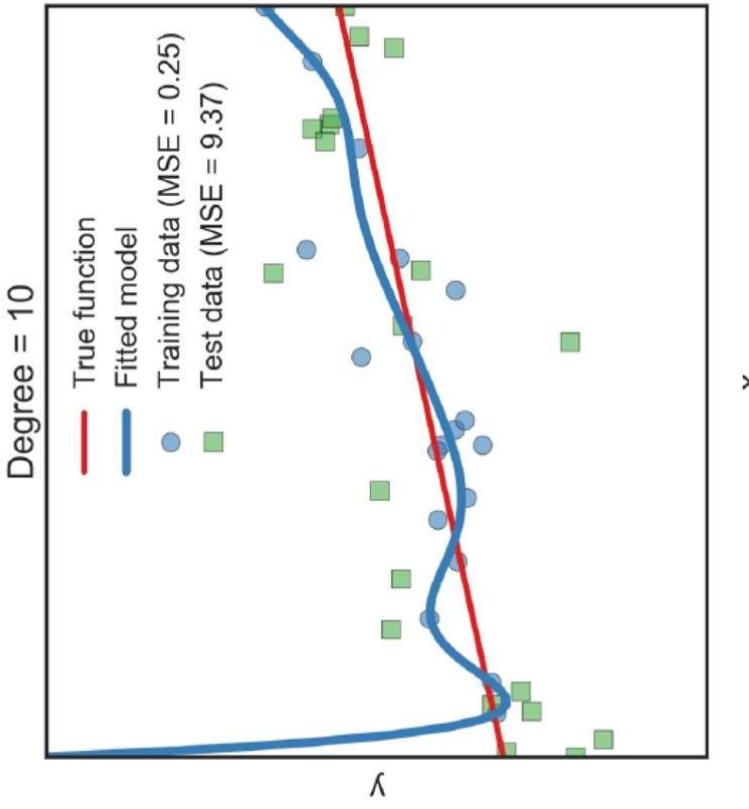
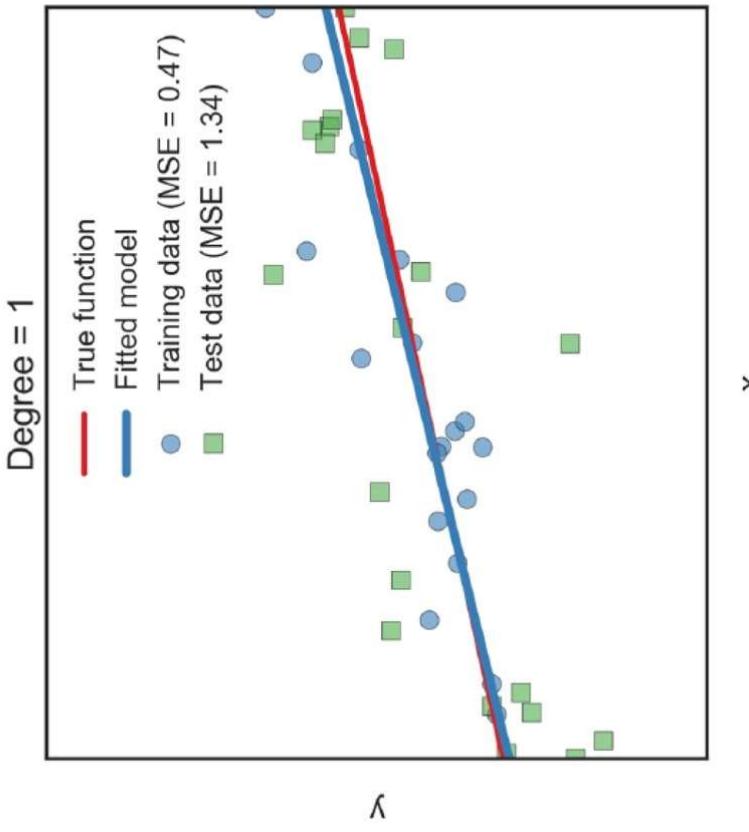
Exploratory analyses done wrong

- Exploratory analyses done poorly: Measure many variables (gender, personality characteristics, age, etc.) and only report those that yield a statistically significant result (stargazing)
 - Include in your paper only those experiments that produced the desired outcome,
 - Treat experiments or initial analyses that didn't turn out favorably as "pilots"
- Why is this problematic?
- In a word, this will lead to model overfitting.

Overfitting

- Overfitting is the tendency for statistical models to mistakenly fit sample-specific noise as if it were signal
 - In a sample of $N = 50$ with 20 uncorrelated predictors, each correlated 0.1 with the DV, the observed (and overfitted) R^2 value will, on average, be 0.45
 - Gives the impression that one could predict values of the DV rather successfully.
 - True R^2 in this situation is only 0.07. Even worse, the average out-of-sample test value of R^2 is only 0.02!

R Squared is very optimistic



Overfitting continued

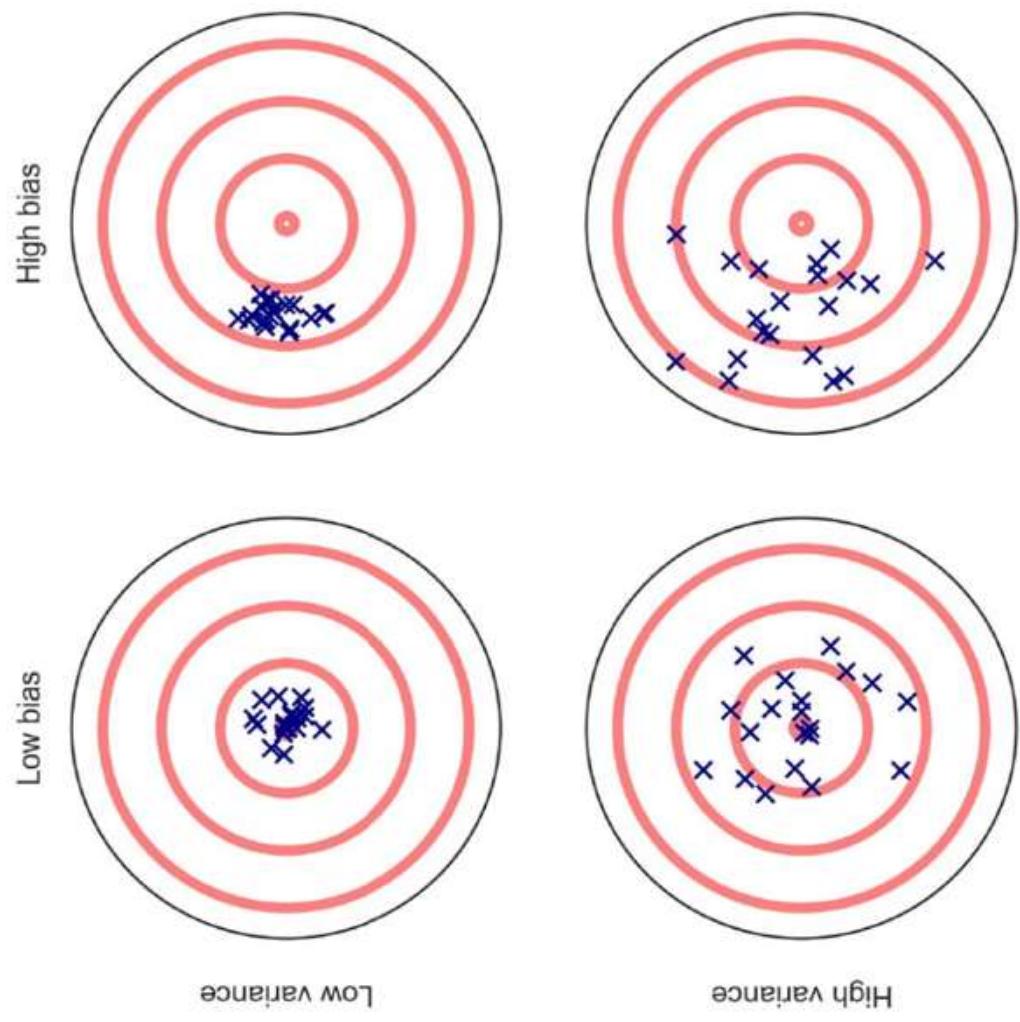
- Don't trust estimates of model performance if those estimates are obtained by "testing" the model on the same data on which it was originally trained
- We need a method for doing exploratory analyses without tricking ourselves.

An aside: The link between p-hacking and overfitting

- p-hacking is a special case of overfitting. Specifically, it is procedural overfitting (Yarkoni and Westfall, 2018). It takes place prior to (or in parallel with) model estimation
 - For example, during data cleaning, model selection, or choosing which analyses to report

We often want to explore though. How do we do that in a principled way?

- First, need to distinguish bias and variance
 - Bias: the tendency for a model to consistently produce answers that are wrong in a particular direction (e.g., estimates that are consistently too high).
 - Variance: the extent to which a model's fitted parameters will tend to deviate from their central tendency across different datasets.



Bias-Variance Tradeoff

- Liberal, flexible data analysis is a low-bias but high-variance approach
- Almost any pattern in data can potentially be detected, at the cost of a high rate of spurious identifications
 - This is exploratory data analysis
- An approach that favors strict adherence to a fixed set of procedures as a high bias, low-variance approach
 - Only a limited range of patterns can be identified, but the risk of pattern hallucination is low
- This is confirmatory data analysis

What to do? Consider lots of possibilities but focus on minimizing prediction error (no stargazing!)

- What's required to do exploratory data analysis that gives you *information* on which you can do confirmatory research?
 - Datasets large enough to support training models
 - Accurately estimate prediction error to assess performance and improve model
 - Exert control over the bias-variance tradeoff when appropriate

Cross-validation

- All of these are directly related to cross-validation and replication
 - To assess our models, we need to quantify out-of-sample prediction error
 - Cross-validation: various techniques involved in training and testing a model on different samples of data

Cross-validation

- Canonical cross-validation

- The classical replication study, where a model is trained on one dataset and then tested on a completely independent dataset. Most typical of experimental research. Less common in correlational research.

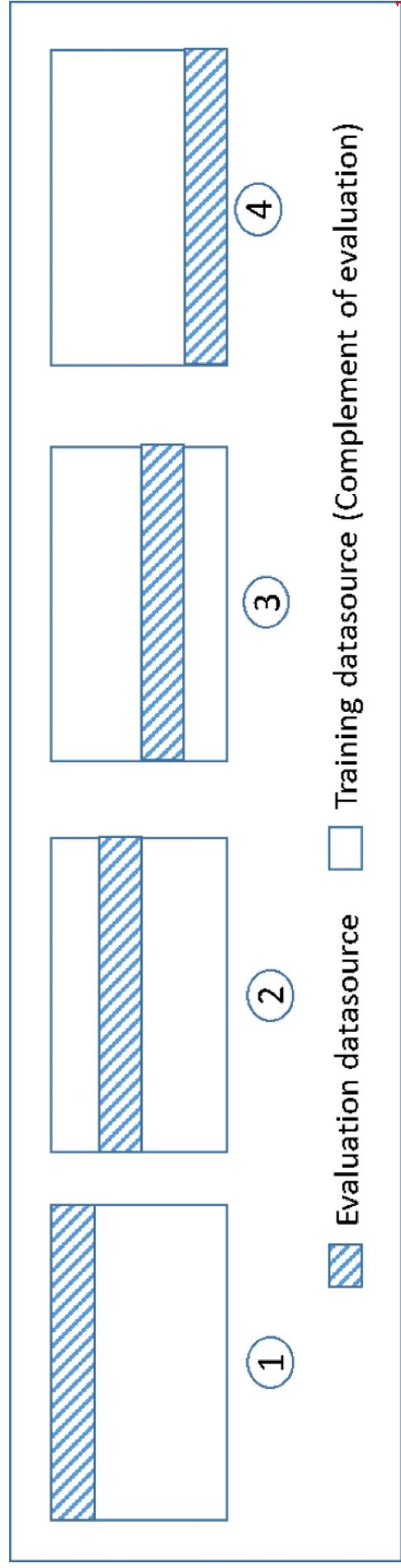


Cross-validation

- Sometimes you can't collect more data though
 - One giant study you want to analyze was run once
 - There is a limited population
 - Limited funds to collect more data

Recycle your dataset.

- Don't assign each observation exclusively to either the training or the test datasets - do both!
- Known as K-Folding where K is the number of folds
- In one "fold" (essentially a subset of your data), one half of the data is used for training and the other half is for testing
- In a second fold, the datasets are reversed, and the training set and test sets exchange roles.
- Typical number of folds is 10





Read data into R

```
Exploreredf <- read_csv("df1.csv")
```

```
## New names:
##   Rows: 5000 Columns: 7
##   -- Column specification
##   -- 
##     (3): screen_name, media_type, text dbl (4): ...1, ...2, favorite_count,
##     retweet_count
##     i Use `spec()` to retrieve the full column specification for this data. i
##     Specify the column types or set `show_col_types = FALSE` to quiet this message.
##     ` ` -> ` ...1 `
##     * ` ...1 ` -> ` ...2 `
```

```
head(Exploreredf)
```

```
## # A tibble: 6 × 7
##   ⋅·1 ⋅·2 screen_name favorite_count retweet_count media_type text
##   <dbl> <chr>        <dbl> <chr>           <dbl> <chr>
## 1 1    DogsTrust      78462           41   Photo          "With our #Car~
## 2 2    DogsTrust      78731           28   Photo          "Running acro~
## 3 3    Greenpeace     48668           466  Nophoto       "When billion~
## 4 4    AnimalAid       80101           70   Nophoto       "AnimalAid ~
```

```
Exploredf1 <- Exploredf%>%
  mutate(Postnumber = 1:n()%>%
    select(-c(...1)))
sentiment <- Exploredf1 %>%
  unnest_tokens(output = "word", input = "text")
```

```
sentiment_dictionary1 <- get_sentiments("bing")  
head(sentiment_dictionary1)
```

```
## # A tibble: 6 × 2  
##   word      sentiment  
##   <chr>    <chr>  
## 1 2-faces  negative  
## 2 abnormal negative  
## 3 abolish  negative  
## 4 abominable negative  
## 5 abominably negative  
## 6 abominate negative
```

```
sentiment_dictionary2 <- get_sentiments("afinn")
head(sentiment_dictionary2)
```

```
## # A tibble: 6 × 2
##   word      value
##   <chr>    <dbl>
## 1 abandon   -2
## 2 abandoned -2
## 3 abandons -2
## 4 abducted  -2
## 5 abduction -2
## 6 abductions -2
```

```
sentiment_dictionary3 <- get_sentiments("nrc")
head(sentiment_dictionary3)
```

```
## # A tibble: 6 × 2
##   word      sentiment
##   <chr>    <chr>
## 1 abacus   trust
## 2 abandon  fear
## 3 abandon  negative
## 4 abandon  sadness
## 5 abandoned anger
## 6 abandoned fear
```

```
sentiment1df <- merge(sentiment, sentiment_dictionary1, by = "word")  
head(sentiment1df)
```

```
## # 1 word      •••2 screen_name favorite_count retweet_count media_type  
## 1 abdominally    72496      peta          78            34      Nophoto  
## 2 absence       85636      WWF           220           70      Nophoto  
## 3 abundance     122417 AWF_Official    92            24      Photo  
## 4 abundance     73701      peta          0             0      Nophoto  
## 5 abundance     93624      Defenders    53            23      Nophoto  
## 6 abundance     1507       oceana        126           23      Photo  
## # Postnumber sentiment  
## 1            3665 negative  
## 2            2455 negative  
## 3            2377 positive  
## 4            2664 positive  
## 5            1056 positive  
## 6            407  positive
```

```
library(summarytools)

## ## Attaching package: 'summarytools'

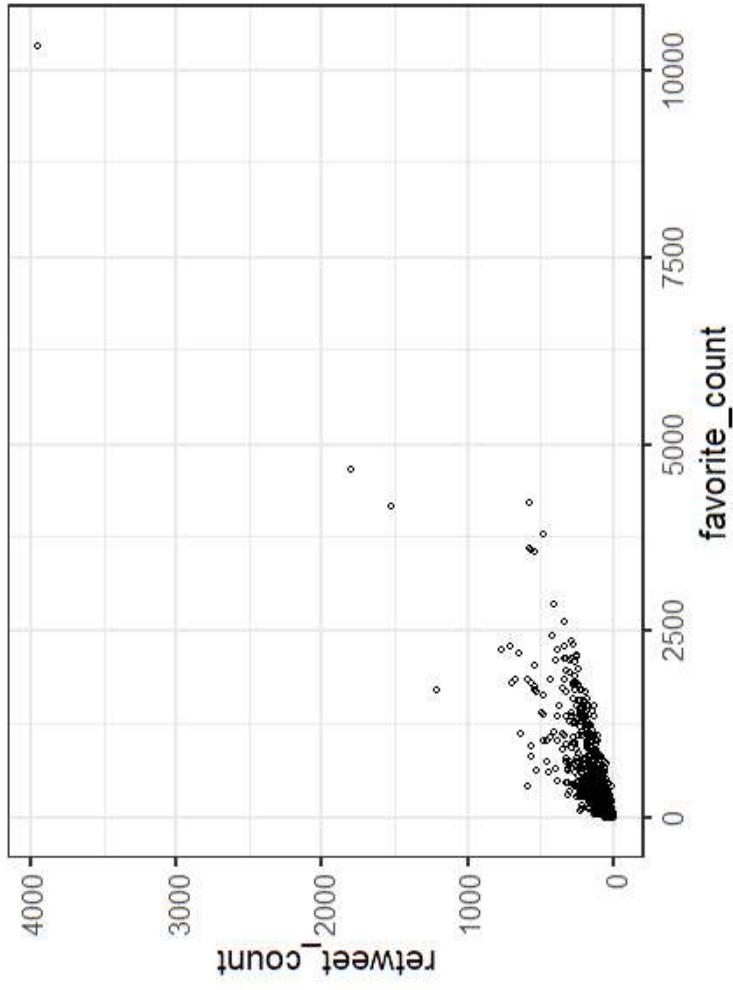
## The following object is masked from 'package:tibble':
##   view

view(dfSummary(sentiment1df))

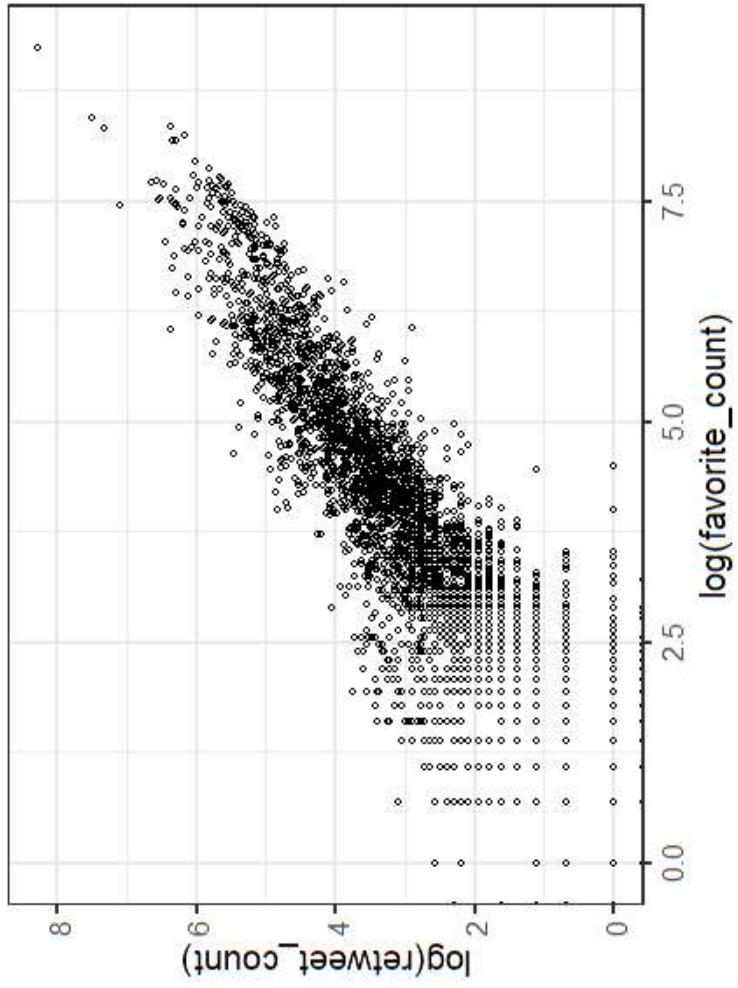
## Switching method to 'browser'

## Output file written: C:\Users\zachs\AppData\Local\Temp\RtmpcBRvG5\file17f0762944b1.html
```

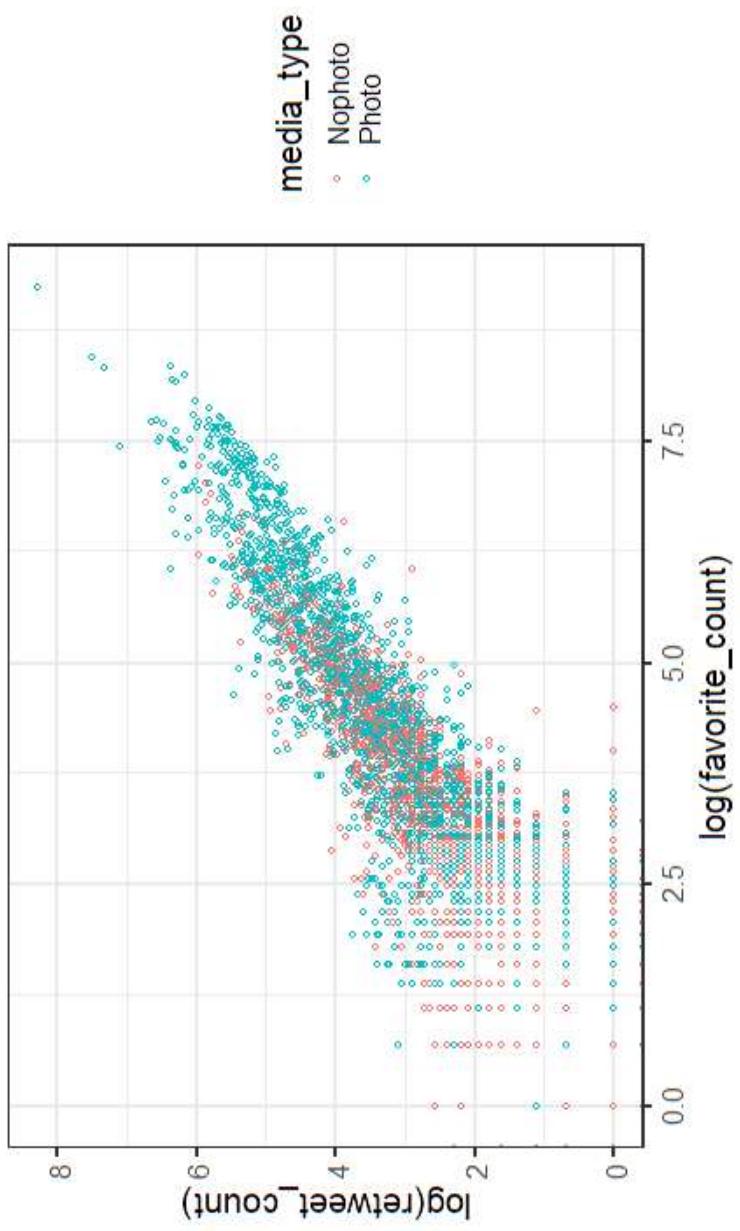
```
ggplot(Exploredf1)+  
  geom_point(aes(y=retweet_count, x = favorite_count), shape=1) +  
  theme_bw(20)
```



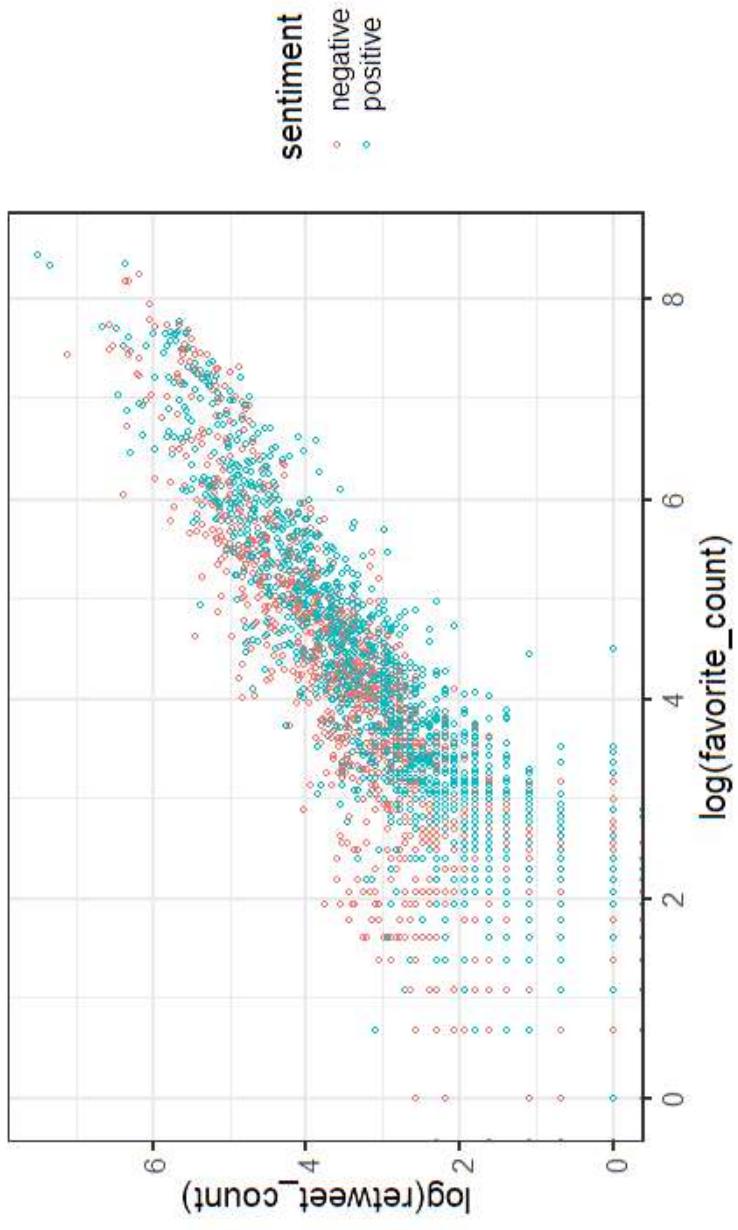
```
ggplot(Exploredf1)+  
  geom_point(aes(y=log(retweet_count) , x = log(favorite_count)) , shape=1) +  
  theme_bw(20)
```



```
ggplot(Exploredf1)+  
  geom_point(aes(y=log(retweet_count), x = log(favorite_count), colour=media_type), shape=1) +  
  theme_bw(20)
```



```
ggplot(sentiment1df) +  
  geom_point(aes(y=log(retweet_count), x = log(favorite_count), colour=sentiment), shape=1) +  
  theme_bw(20)
```



```
m1 <- lm(retweet_count ~ media_type, data = sentiment1df)
summary(m1)

## 
## Call:
## lm(formula = retweet_count ~ media_type, data = sentiment1df)
## 
## Residuals:
##    ##   Min     1Q   Median     3Q    Max
##    ## -68.95 -42.95 -18.95  4.24 1731.05
## 
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) 19.762   1.443   13.70  <2e-16 ***
## media_typePhoto 49.191   1.988   24.75  <2e-16 ***
## ---      
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## 
## Residual standard error: 89.9 on 8202 degrees of freedom
## Multiple R-squared:  0.06947, Adjusted R-squared:  0.06936 
## F-statistic: 612.4 on 1 and 8202 DF, p-value: < 2.2e-16
```

```
m2 <- lm(retweet_count ~ media_type + sentiment, data=sentiment1df)
summary(m2)

## 
## Call:
## lm(formula = retweet_count ~ media_type + sentiment, data = sentiment1df)

## Residuals:
##    Min     1Q   Median     3Q    Max 
## -78.32 -43.76 -13.77  4.67 1736.24 

## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) 28.333    1.873   15.12   < 2e-16 ***
## media_typePhoto 49.984    1.985   25.18   < 2e-16 ***
## sentimentpositive -14.560    2.039  -7.14 1.01e-12 ***
## ---      
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 

## 
## Residual standard error: 89.62 on 8201 degrees of freedom
## Multiple R-squared:  0.07522, Adjusted R-squared:  0.075 
## F-statistic: 333.5 on 2 and 8201 DF, p-value: < 2.2e-16
```

```
library(purrr)
library(modelr)
```

```
## Warning: package 'modelr' was built under R version 4.1.3
```

```
cv <- crossv_kfold(sentiment1df, k = 10)
```

```
cv
```

```
## # A tibble: 10 × 3
##   train          test
##   <named list> <named list>
## 1 <resample [7, 383 × 8]> <resample [821 × 8]> 01
## 2 <resample [7, 383 × 8]> <resample [821 × 8]> 02
## 3 <resample [7, 383 × 8]> <resample [821 × 8]> 03
## 4 <resample [7, 383 × 8]> <resample [821 × 8]> 04
## 5 <resample [7, 384 × 8]> <resample [820 × 8]> 05
## 6 <resample [7, 384 × 8]> <resample [820 × 8]> 06
## 7 <resample [7, 384 × 8]> <resample [820 × 8]> 07
## 8 <resample [7, 384 × 8]> <resample [820 × 8]> 08
## 9 <resample [7, 384 × 8]> <resample [820 × 8]> 09
## 10 <resample [7, 384 × 8]> <resample [820 × 8]> 10
```

```
models0 <- map(cv$train, ~lm(retweet_count ~ 1, data = .))
models1 <- map(cv$train, ~lm(retweet_count ~ media_type, data = .))
models2 <- map(cv$train, ~lm(retweet_count ~ media_type + sentiment, data = .))
```

How does map function work?

```
my_list <- list(  
  c(1, 2, 6),  
  c(4, 7, 1),  
  c(9, 1, 5)  
)
```

```
my_list
```

```
## [[1]]  
## [1] 1 2 6  
## [[2]]  
## [1] 4 7 1  
## [[3]]  
## [1] 9 1 5
```

```
my_list[[1]]
```

```
## [1] 1 2 6
```

```
my_list[[1]] %>% mean()
```

```
## [1] 3
```

```
my_list[[2]] %>% mean()
```

```
## [1] 4
```

```
my_list[[3]] %>% mean()
```

- -

```
my_list
```

```
## [[1]]  
## [1] 1 2 6  
## [[2]]  
## [1] 4 7 1  
## [[3]]  
## [1] 9 1 5
```

```
my_list %>% map(mean)
```

```
## [[1]]  
## [1] 3  
## [[2]]  
## [1] 4  
## [[3]]  
## [1] 5
```

```
map(my_list, mean)
```

```
## [[1]]  
## [1] 3  
## ...
```

```
# gives a vector of dbl  
my_list %>% map_dbl(mean)  
  
## [1] 3 4 5  
  
# anonymous function  
# ~ creation function with no specific name in tidyverse  
# . every element in the list  
  
my_list %>% map(~ . * 2)  
  
## [[1]]  
## [1] 2 4 12  
## [[2]]  
## [1] 8 14 2  
## [[3]]  
## [1] 18 2 10
```

```
get_pred <- function(model, test_data){  
  data <- as.data.frame(test_data)  
  pred <- add_predictions(data, model)  
  return(pred)  
}  
  
pred0 <- map2_df(models0, cv$test, get_pred, .id = "Run")  
pred1 <- map2_df(models1, cv$test, get_pred, .id = "Run")  
pred2 <- map2_df(models2, cv$test, get_pred, .id = "Run")
```

Mean Squared Error to assess model fit

$$\text{MSE} = \frac{1}{n} \sum_{i=1}^n (Y_i - \hat{Y}_i)^2$$

MSE = mean squared error

n = number of data points

Y_i = observed values

\hat{Y}_i = predicted values

- To find the MSE, take the observed value, subtract the predicted value, and square that difference. Repeat that for all observations. Then, sum all of those squared values and divide by the number of observations.

Interpreting Mean Squared Error

- An MSE of zero, meaning that the estimator $\hat{\theta}$ predicts observations of the parameter θ with perfect accuracy.
- Two or more statistical models may be compared using their MSEs as a measure of how well they explain a given set of observations.
- Mean squared error has the disadvantage of heavily weighting outliers. This property, undesirable in many applications, has led researchers to use alternatives such as the mean absolute error.

```
MSE0 <- pred0 %>% group_by(Run) %>%
  summarise(MSE = mean((retweet_count - pred)^2))
```

```
MSE0
```

```
## # A tibble: 10 × 2
##   Run     MSE
##   <chr>  <dbl>
## 1 1      5371.
## 2 10     15987.
## 3 2      5232.
## 4 3      9980.
## 5 4      7747.
## 6 5      10138.
## 7 6      8114.
## 8 7      6248.
## 9 8      8788.
## 10 9     9250.
```

```
MSE1 <- pred1 %>% group_by(Run) %>%
  summarise(MSE = mean( (retweet_count - pred)^2))
```

```
MSE1
```

```
## # A tibble: 10 × 2
##   Run     MSE
##   <chr>  <dbl>
## 1 1      4930.
## 2 2     15116.
## 3 2     4795.
## 4 3     9447.
## 5 4     7011.
## 6 5     9380.
## 7 6     7691.
## 8 7     5779.
## 9 8     8238.
## 10 9    8462.
```

```
MSE2 <- pred2 %>% group_by(Run) %>%
  summarise(MSE = mean( (retweet_count - pred)^2))
```

MSE2

```
## # A tibble: 10 × 2
##   Run     MSE
##   <chr>  <dbl>
## 1 1      4898.
## 2 10     14964.
## 3 2      4718.
## 4 3      9438.
## 5 4      7005.
## 6 5      9372.
## 7 6      7670.
## 8 7      5742.
## 9 8      8204.
## 10 9     8363.
```

```
mean(MSE0$MSE)
## [1] 8685.354

mean(MSE1$MSE)
## [1] 8084.726

mean(MSE2$MSE)
## [1] 8037.398
```

```
ConfirmRmdf <- read_csv("df2.csv")
```

```
## New names:
##   Rows: 20000 Columns: 7
##   -- Column specification
##   -
##     (3): screen_name, media_type, text dbl (4): ...1, ...2, favorite_count,
##     retweet_count
##     i Use `spec()` to retrieve the full column specification for this data. i
##     Specify the column types or set `show_col_types = FALSE` to quiet this message.
##     `` -> `...1`
##     * `...1` -> `...2`
```

```
head(ConfirmRmdf)
```

	# A tibble: 6 × 7	# ⋯.1	# ⋯.2	screen_name	favorite_count	retweet_count	media_type	text
	<code><dbl></code>	<code><dbl></code>	<code><dbl></code>	<code><chr></code>	<code><dbl></code>	<code><dbl></code>	<code><chr></code>	<code><chr></code>
## 1	1	1	539	oceana	438	121	Photo	GOOD NEWS~@Isabella~We<U+0092~
## 2	2	2	22589	sascampaigns	0	0	Nophoto	14 Nophoto
## 3	3	3	62705	ClimateReality	49	32	Photo	Demand ha~Wildfire ~
## 4	4	4	114033	pawtitions	9	0	Nophoto	226 Nophoto
## 5	5	5	38309	therightblue	1			Last mont~
## 6	6	6	62185	ClimateReality	654			

Similar computations as on Explor

df

```
Confirmdf <- Confirmdf%>%
  mutate(Postnumber = 1:n()%>%
    select(-c(...1))  
  
confirmsegment <- Confirmdf %>%
  unnest_tokens(output = "word", input = "text")
```

```
sentiment_dictionary1 <- get_sentiments("bing")  
head(sentiment_dictionary1)
```

```
## # A tibble: 6 × 2  
##   word      sentiment  
##   <chr>    <chr>  
## 1 2-faces  negative  
## 2 abnormal negative  
## 3 abolish  negative  
## 4 abominable negative  
## 5 abominably negative  
## 6 abominate negative
```

```
sentiment_dictionary2 <- get_sentiments("afinn")
head(sentiment_dictionary2)
```

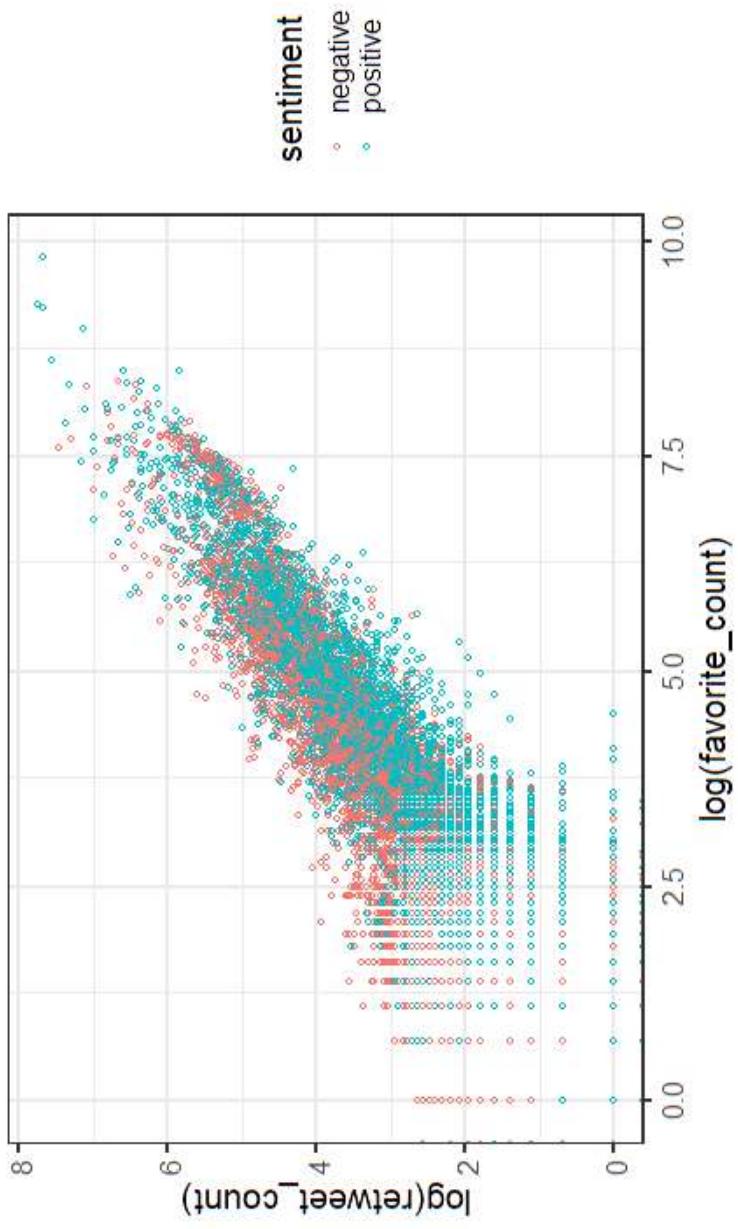
```
## # A tibble: 6 × 2
##   word      value
##   <chr>    <dbl>
## 1 abandon   -2
## 2 abandoned -2
## 3 abandons -2
## 4 abducted  -2
## 5 abduction -2
## 6 abductions -2
```

```
sentiment_dictionary3 <- get_sentiments("nrc")
head(sentiment_dictionary3)
```

```
## # A tibble: 6 × 2
##   word      sentiment
##   <chr>    <chr>
## 1 abacus   trust
## 2 abandon  fear
## 3 abandon  negative
## 4 abandon  sadness
## 5 abandoned anger
## 6 abandoned fear
```

```
confirmsentiment1df <- merge(confirmssentiment, sentiment_dictionary1, by = "word")  
head(confirmsentiment1df)  
  
## # word ...2 screen_name favorite_count retweet_count media_type  
## 1 abnormal 70074 HSIGlobal 420 168 Nophoto  
## 2 abnormal 80025 MoveTheWorld 315 331 Photo  
## 3 abnormal 32344 savingoceans 6 2 Photo  
## 4 abolish 112289 Network4Animals 23 21 Photo  
## 5 abolish 101041 FarmSanctuary 48 10 Photo  
## 6 abound 48231 Greenpeace 46 19 Nophoto  
## Postnumber sentiment  
## 1 12913 negative  
## 2 11822 negative  
## 3 6203 negative  
## 4 9532 negative  
## 5 3559 negative  
## 6 16354 positive
```

```
ggplot(confirmationsentiment1df)+  
  geom_point(aes(y=log(retweet_count), x = log(favorite_count), colour=sentiment), shape=1) +  
  theme_bw(20)
```



Issues though...

```
predictedvalues <- predict(m1, data=sentiment1df, interval="prediction")  
  
## Warning in predict.lm(m1, data = sentiment1df, interval = "prediction"): predictions on current data re...  
  
View(predictedvalues)
```

```
v1 <- c(19.76, -156, 196)
v2 <- c(68.95279, -107, 245)
v3 <- c("Nophoto", "Photo")

smallldf <- rbind(v1,v2)

smallldf <- as.data.frame(smallldf) %>%
  rename("meanretweet" = "V1",
         "Lower" = "V2",
         "Upper" = "V3")

smalldf <- cbind(smallldf, v3)

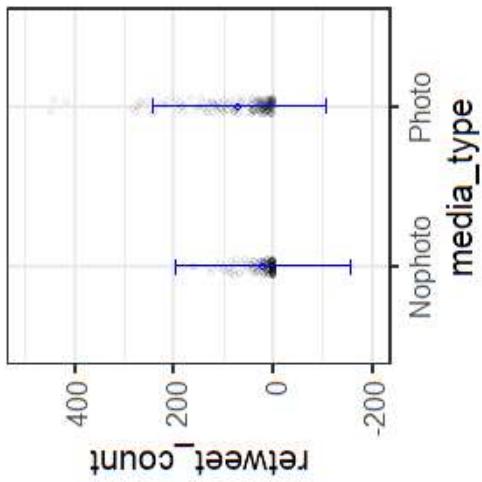
smalldf <- smalldf %>%
  rename("media_type" = "v3")

sampleConfirmsdf <- confirmsentimentdf %>%
  sample_n(400)
```

What's wrong with this picture?

```
ggplot() +  
  geom_jitter(data=sampleConfirmdf, aes(x = media_type, y = retweet_count), width = .05, height=.01)  
  geom_point(data=smallldf, aes(x= media_type, y= meanretweet), shape = 1, colour = "blue") +  
  geom_errorbar(data=smallldf, aes(x= media_type, ymin = lower,ymax=upper), width=.1, colour = "blue")  
  scale_y_continuous(limits = c(-200,500)) +  
  theme_bw(20)
```

Warning: Removed 5 rows containing missing values (`geom_point()``).



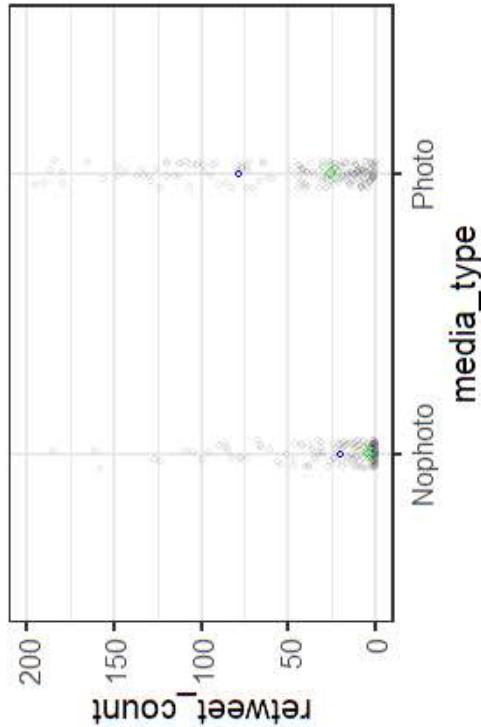
Impossible values in our error bars

- Need a GLM that takes into account the data are counts!

```
m1poss <- glm(retweet_count ~ media_type, data = Confirmdf, family = "poisson")  
summary(m1poss)  
  
##  
## Call:  
## glm(formula = retweet_count ~ media_type, family = "poisson",  
## data = Confirmdf)  
  
## Deviance Residuals:  
##   Min     1Q Median     3Q    Max  
## -10.81  -5.71  -4.60   0.42  428.98  
  
## Coefficients:  
## (Intercept) 2.789954  0.002365 1179.9 <2e-16 ***  
## media_typePhoto 1.277500  0.002737  466.8 <2e-16 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
  
## (Dispersion parameter for poisson family taken to be 1)
```

```
sampleConfirmdf %>%
  group_by(media_type)%>%
  summarise(meanrt = mean(retweet_count),
            medianrt = median(retweet_count))%>%
  ggplot() +
  geom_jitter(data=sampleConfirmdf, aes(x = media_type, y = retweet_count), width = .05, height=.01,
  geom_point(aes(x= media_type, y= meanrt), shape = 1, colour = "blue")+
  geom_point(aes(x= media_type, y= medianrt), shape = 5, colour = "green")+
  scale_y_continuous(limits = c(0,200))+
  theme_bw(20)
```

Warning: Removed 56 rows containing missing values (`geom_point()`).



```

m2poss <- glm(retweet_count ~ media_type + sentiment , data = sentiment1df, family = "poisson")

## Call:
## glm(formula = retweet_count ~ media_type + sentiment, family = "poisson",
##      data = sentiment1df)

## Deviance Residuals:
##    Min      1Q      Median      3Q      Max
## -12.911   -6.852    -4.662     0.719    93.321

## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) 3.155850  0.003983  792.2  <2e-16 ***
## media_typePhoto 1.267198  0.004053  312.7  <2e-16 ***
## sentimentpositive -0.312762  0.003292  -95.0  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## (Dispersion parameter for poisson family taken to be 1)

## Null deviance: 770433 on 8203 degrees of freedom
## Residual deviance: 644619 on 8201 degrees of freedom
## AIC: 677966

## Number of Fisher Scoring iterations: 6

```

