Lecture 13 Quantitative Political Science

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Agenda

- 1. Associations
- 2. Describing relationships
- 3. Modeling relationships

Associations between two (or more) variables

- Thus far, description & inference for either one variable or two variables drawn from different units
- Now, what you are here for! Relationships between two variables from the *same* units!
- Bivariate:
 - $\circ~X$ (independent / predictor / explanatory / RHS) and Y (dependent / outcome)
 - Implies a causal intuition, but does NOT buy it!
- 4 approaches (more like steps)
 - 1. Displaying relationships
 - 2. Summarizing non-parametrically
 - 3. Summarizing parametrically
 - 4. Making inferences about the relationship in a population from a sample

ullet Crosstab(ulation)s: Y typically in rows, X in columns

```
require(tidyverse)
dat <- read_rds('https://github.com/jbisbee1/PSCI_8356/raw/main/Lectures/Data/sc_debt.Rds')
t <- table(dat$preddeg,dat$control)
t</pre>
```

```
##
## Private Public
## Associate 127 694
## Bachelor's 1193 532
```

ullet Crosstab(ulation)s: Y typically in rows, X in columns

```
prop.table(t,margin = 1) # Rows
##
                  Private
                            Public
     Associate 0.1546894 0.8453106
     Bachelor's 0.6915942 0.3084058
prop.table(t,margin = 2) # Columns
##
##
                               Public
                   Private
##
     Associate 0.09621212 0.56606852
##
     Bachelor's 0.90378788 0.43393148
```

ullet Crosstab(ulation)s: Y typically in rows, X in columns...BAD FOR MANY CATEGORIES OR CONTINUOUS!

```
table(dat$md_earn_wne_p6,dat$sat_avg)
```

```
##
                          851 854 855 861 865 875 876 877 880
     10600
                                                       0
##
     11000
                                                       0
##
     11800
     11900
                                                       0
##
     12200
##
     12800
     12900
                                                       0
     13000
                                                       0
     13400
                                                       0
     13700
     14000
     14100
     14200
     14300
     14400
     14500
```

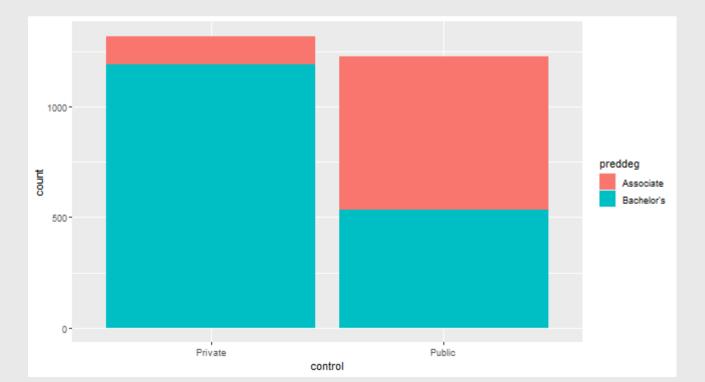
Use bins first

```
##
##
                      (737,1053] (1053,1119] (1119,1205]
##
     (10600, 26100]
                           20.33
                                         3.68
                                                      0.66
##
     (26100,31500]
                           34.00
                                        27.76
                                                     19.87
##
     (31500,37400)
                          31.33
                                        50.17
                                                     39.40
##
     (37400,120400)
                          14.33
                                        18.39
                                                     40.07
##
##
                      (1205, 1557)
##
     (10600, 26100)
                             2.03
##
     (26100,31500]
                             5.07
##
      (31500,37400)
                            22.97
##
      (37400,120400]
                            69.93
```

- Plotting
 - \circ Barplots (geom_bar()): X and Y are both categorical (including binary)
 - \circ Densities / histograms (geom_density() / geom_histogram()): X is binary and Y is continuous
 - \circ Boxplots / violin plots (geom_boxplot() / geom_violin()): X is categorical and Y is continuous
 - \circ Scatterplots (geom_point()): X and Y are both continuous

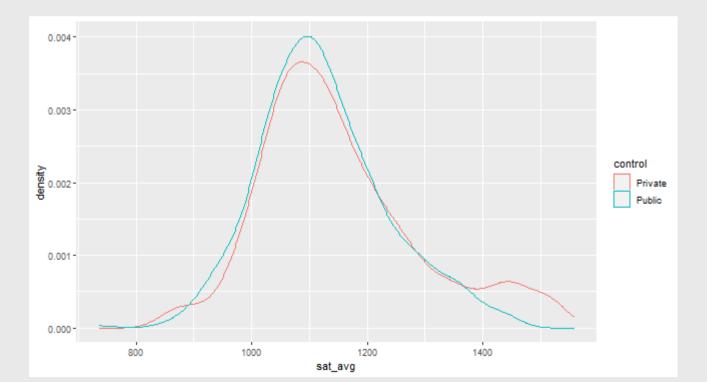
• Barplots ($geom_bar()$): X and Y are both categorical (including binary)

```
dat %>%
  ggplot(aes(x = control,fill = preddeg)) +
  geom_bar()
```



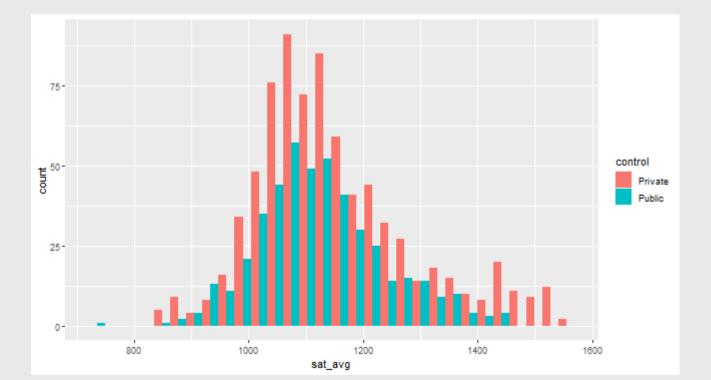
• Densities / histograms ($geom_density()$ / $geom_histogram()$): X is binary and Y is continuous

```
dat %>%
  ggplot(aes(x = sat_avg,color = control)) +
  geom_density()
```



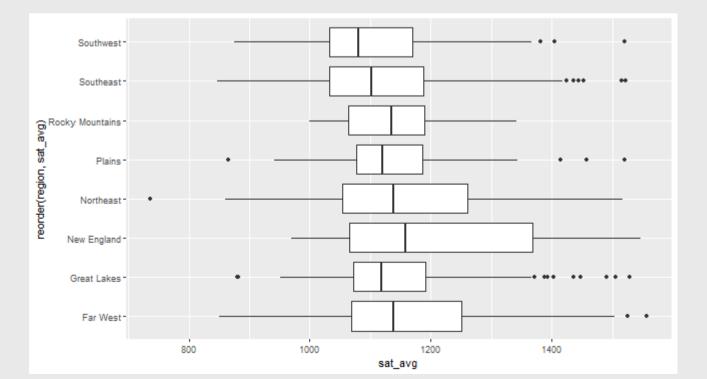
• Densities / histograms ($geom_density()$ / $geom_histogram()$): X is binary and Y is continuous

```
dat %>%
  ggplot(aes(x = sat_avg,fill = control)) +
  geom_histogram(position = 'dodge')
```



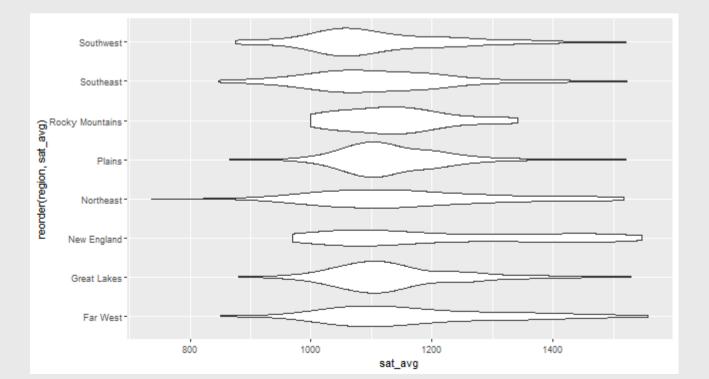
• Boxplots / violin plots (geom_boxplot() / geom_violin()): X is categorical and Y is continuous

```
dat %>%
  ggplot(aes(x = sat_avg,y = reorder(region,sat_avg))) +
  geom_boxplot()
```



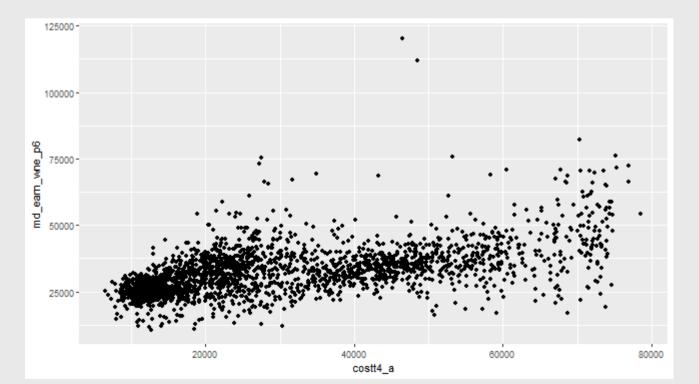
• Boxplots / violin plots (geom_boxplot() / geom_violin()): X is categorical and Y is continuous

```
dat %>%
  ggplot(aes(x = sat_avg,y = reorder(region,sat_avg))) +
  geom_violin()
```



• Scatterplots ($geom_point()$): X and Y are both continuous

```
dat %>%
  ggplot(aes(x = costt4_a,y = md_earn_wne_p6)) +
  geom_point()
```



- Conditional means
 - \circ What is the average value of Y for a given value of X?

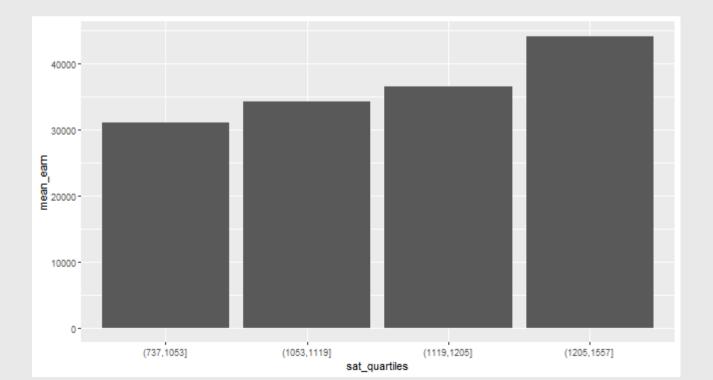
```
dat %>%
  drop_na(sat_quartiles) %>%
  group_by(sat_quartiles) %>%
  summarise(mean_earn = mean(md_earn_wne_p6,na.rm=T))
```

- Conditional means
 - \circ What is the average value of Y for a given value of X?

```
p <- dat %>%
  drop_na(sat_quartiles) %>%
  group_by(sat_quartiles) %>%
  summarise(mean_earn = mean(md_earn_wne_p6,na.rm=T)) %>%
  ggplot(aes(x = sat_quartiles,y = mean_earn)) +
  geom_bar(stat = 'identity')
```

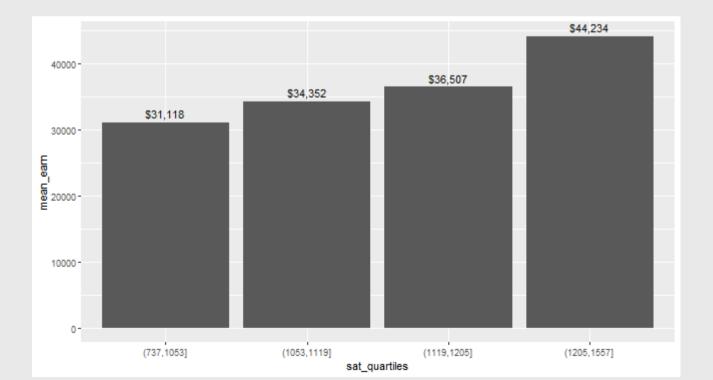
- Conditional means
 - \circ What is the average value of Y for a given value of X?

p



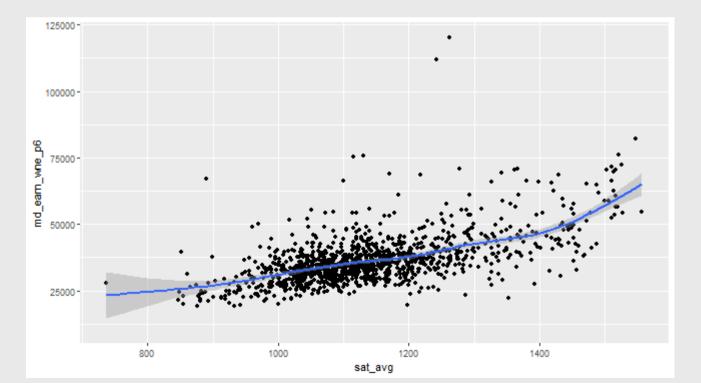
- Conditional means
 - \circ What is the average value of Y for a given value of X?

```
p + geom_text(aes(label = scales::dollar(round(mean_earn))),vjust = -.5)
```



Smoothers

```
dat %>%
  ggplot(aes(x = sat_avg,y = md_earn_wne_p6)) +
  geom_point() + geom_smooth()
```



- Want to use *models* to describe **theoretical** relationships
- Want minimal **assumptions**...thus far?
- ullet For inferences about μ with **large** samples
 - \circ **identicality**: necessary for $ar{Y}$ to be unbiased for μ
 - \circ independence: necessary for $VAR(ar{Y})=rac{\sigma^2}{n}$
- ullet For inferences about μ with **small** samples

$$\circ~Y \sim \mathcal{N}(\mu, \sigma^2)$$

- For inferences about differences in population means with **large** samples
 - Two samples are drawn independently
- For inferences about differences in population means with **large** samples
 - Two samples are drawn independently
 - Two samples have the **same variance**
 - Underlying populations are Normal
- This is a pretty short list!
- Lots more to come with bivariate and multivariate analysis!

- How to describe a bivariate relationship?
- Start with notion of correlation

$$ho = rac{COV(Y_1,Y_2)}{\sigma_1\sigma_2} \ = rac{E[(Y_1-\mu_1)(Y_2-\mu_2)]}{\sigma_1\sigma_2}$$

ullet Translating to bivariate world is easy, just use X and Y

$$ho = rac{E[(X - \mu_X)(Y - \mu_Y)]}{\sigma_X \sigma_Y}$$

- But ρ is a theoretical quantity (a **parameter**)
- What is a good estimator?

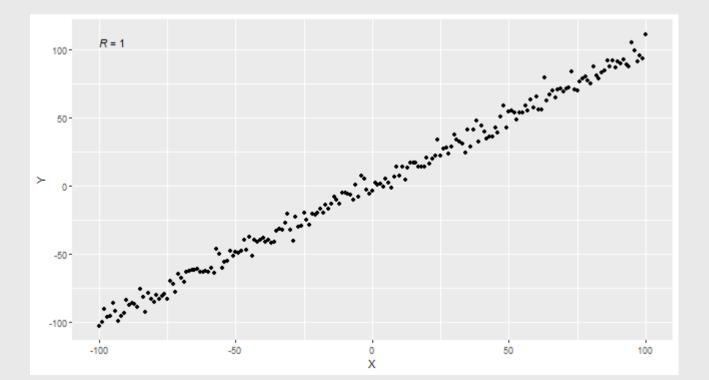
$$r = rac{\sum_{i} (X_{i} - ar{X})(Y_{i} - ar{Y})}{\sqrt{\sum_{i} (X_{i} - ar{X})^{2} \sum_{i} (Y_{i} - ar{Y})^{2}}}$$

- Replace the covariance with the sample covariance $s_{XY}=\sum_i(X_i-\bar{X})(Y_i-\bar{Y})$ and the standard deviations for both variables with their sample analogues $s_X=\sqrt{\sum_i(X_i-\bar{X})^2}$ and $s_Y=\sqrt{\sum_i(Y_i-\bar{Y})^2}$.
- How good is this? It depends on the underlying data

```
X \leftarrow seq(-100,100,by = 1)
```

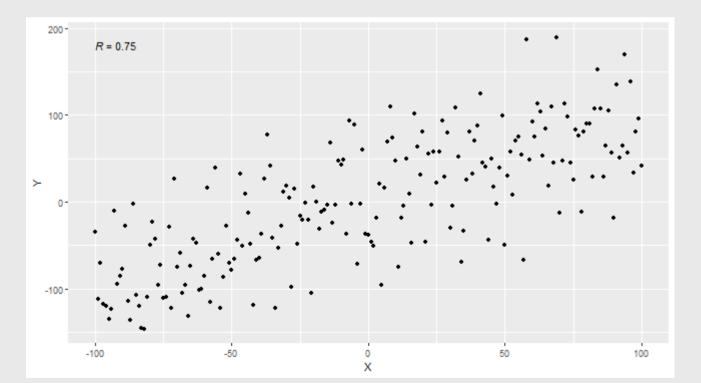
• Works well with linear relationships

```
Y <- X + rnorm(length(X),mean = 0,sd = 5)
data.frame(X = X,Y = Y) %>%
  ggplot(aes(x = X,y = Y)) + geom_point() + stat_cor(p.digits = NA,label.sep = '')
```



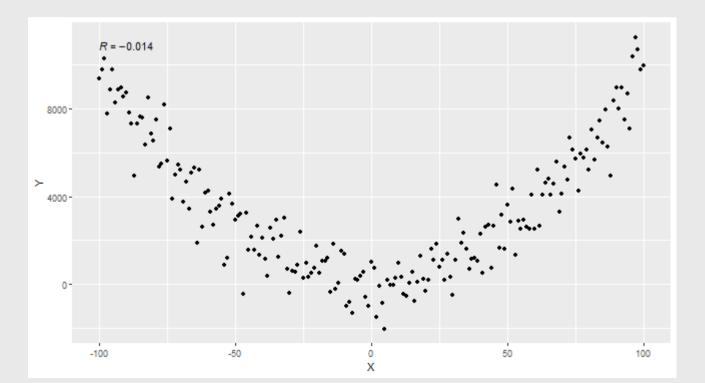
Still works ok with more noise

```
Y <- X + rnorm(length(X),mean = 0,sd = 50)
data.frame(X = X,Y = Y) %>%
  ggplot(aes(x = X,y = Y)) + geom_point() + stat_cor(p.digits = NA,label.sep = '')
```



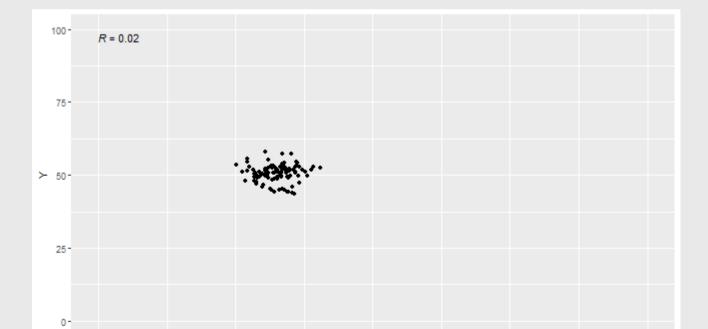
• Doesn't work well with curvelinear relationships

```
Y <- X^2 + rnorm(length(X), mean = 0, sd = 1000)
data.frame(X = X,Y = Y) %>%
  ggplot(aes(x = X,y = Y)) + geom_point() + stat_cor(p.digits = NA, label.sep = '')
```



Very sensitive to outliers

```
X <- rnorm(100, mean = 33, sd = 3)
Y <- rnorm(100, mean = 50, sd = 3)
data.frame(X = X,Y = Y) %>%
    ggplot(aes(x = X,y = Y)) + geom_point() + stat_cor(p.digits = NA, label.sep = '') + lims(x = c(0,100),y)
= c(0,100))
```



Very sensitive to outliers

```
X[75] <- Y[75] <- 75
data.frame(X = X,Y = Y) %>%
  ggplot(aes(x = X,y = Y)) + geom_point() + stat_cor(p.digits = NA,label.sep = '') +
  xlim(c(0,100)) + ylim(c(0,100))
```

