Lab 5 & 6: Stata, graphics, and data

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Reading and saving data in Stata

```
//remove any existing data and variables
clear
// Print working directory (check what the current working directory is)
pwd
// Set the working directory you want -- use your own here, not mine
cd "/Users/robinmejia/Dropbox/Long Data Fall 2015/Assignments/Assignment3"
// read in data using insheet comand
insheet using strength.csv
// Do things to your data if you want...
// Save a file called strength.dta (Stata format) in your working directory
save strength, replace
```

Schizophrenia Data Set

- There is a file called schizophrenia.dta in bCourses.
 Please download it and double click to open it in Stata
- 5 variables:
 - id: 437 people, measured 2 to 4 times each
 - week: discrete scale (0, ..., 6)
 - gender: female/male (0, 1)
 - drug: untreated/treated (0, 1)
 - severity: discrete scale (1, ..., 7)
- Taken from Robert Weiss's website, a good source for longitudinal datasets:
 - https://faculty.biostat.ucla.edu/robweiss/book-data-sets

Goals Today

 Perform basic exploratory data analysis on the relationship between drug status and severity of episodes.

 There is another binary variable in this dataset, gender. Please feel free to explore the code here by adapting it to the relationship between gender and severity of episodes.

One Dimensional Summaries

- How many unique individuals exist here?
- Of those, how many are being treated and how many are not?
- How many females and males?
- What is the distribution of observations over weeks?

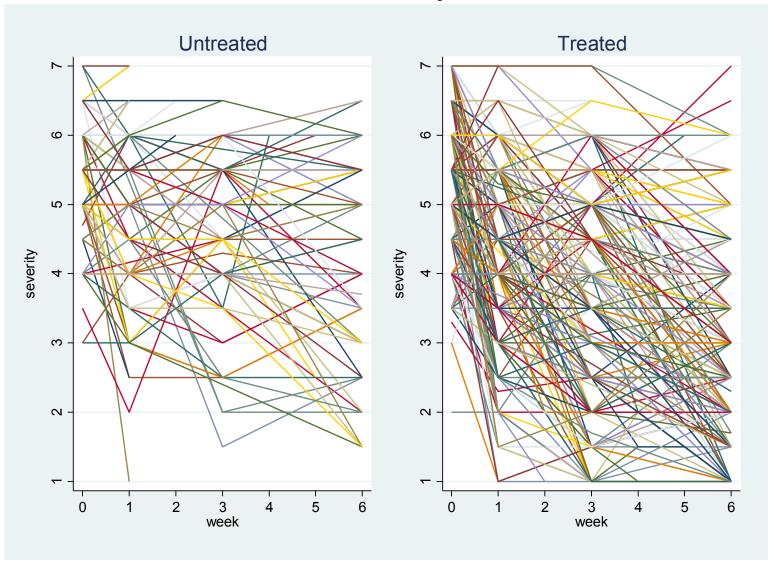
One Dimensional Summaries

- xtset id week
- xtdescribe

Plot severity vs week for 2 groups

```
// This code wil plot severity by week for the treated and untreated.
// Try it and play with the settings to see what happens
xtline severity if drug==0, i(id) overlay t(week) xlab(0(1)6) ylab(1(1)7)
title(Untreated) legend(off)
graph save "untreated.gph", replace
xtline severity if drug==1, i(id) overlay t(week) xlab(0(1)6) ylab(1(1)7)
title(Treated) legend(off)
graph save "treated.gph", replace
gr combine "untreated.gph" "treated.gph"
graph save "untreated_treated.gph", replace
```

Scatterplot



Observations from this plot

- There are fewer untreated than treated people (108 vs. 329)
- Untreated people seem to have a random trajectory – in fact, it seems almost horizontal.
- There might be a downward trend over time in the treated people – it's hard to tell
- This isn't so nice to look at...

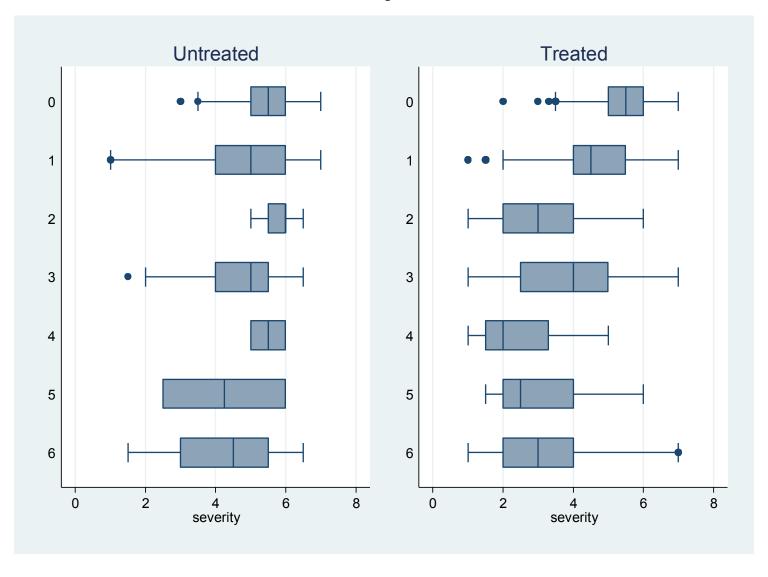
Boxplots of severity by week, Treated vs. Untreated

```
graph hbox severity if drug==0, over(week)
title(Untreated)
graph save "untreated_box.gph", replace
```

```
graph hbox severity if drug==1, over(week) title(Treated) graph save "treated_box.gph", replace
```

```
gr combine "untreated_box.gph" "treated_box.gph" graph save "treated_untreated_box.gph", replace
```

Boxplot



Plot regression lines by id

Another way to look for trends is to smooth the data

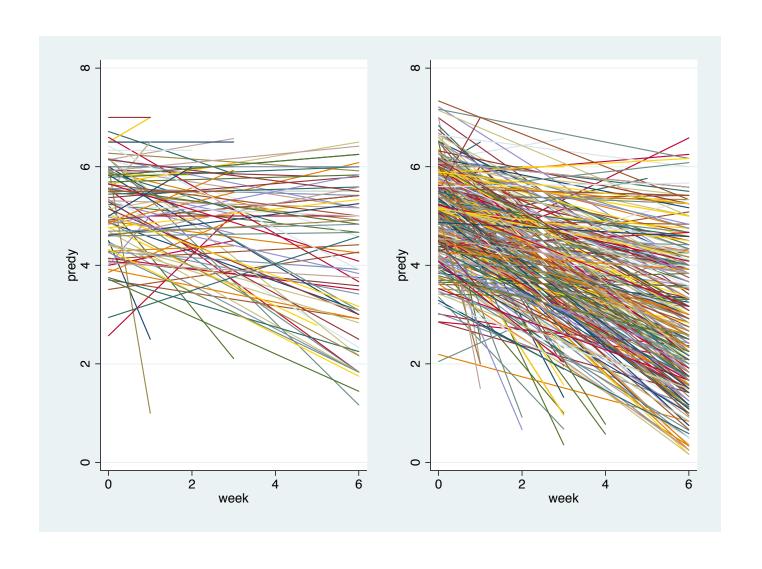
On the Bcourses site, there's a dofile to generate regression lines by id and capture the slope and intercept, with some tips for plotting

Plot regression lines by id

Another way to look for trends is to smooth the data

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Plot regression lines by id



Test the effectiveness of the drug on the severity of the episode

- First, we need to define what "effectiveness" is.
- One way could be to look at the difference between the severity of the attack in the 0th week vs. the 6th week.
- How do we best achieve that? Look at the data in "wide" format

Code to go from long to wide format

drop nvals

```
//reshape to wide by id, creating multiple week variables reshape wide severity drug gender, i(id) j(week)
```

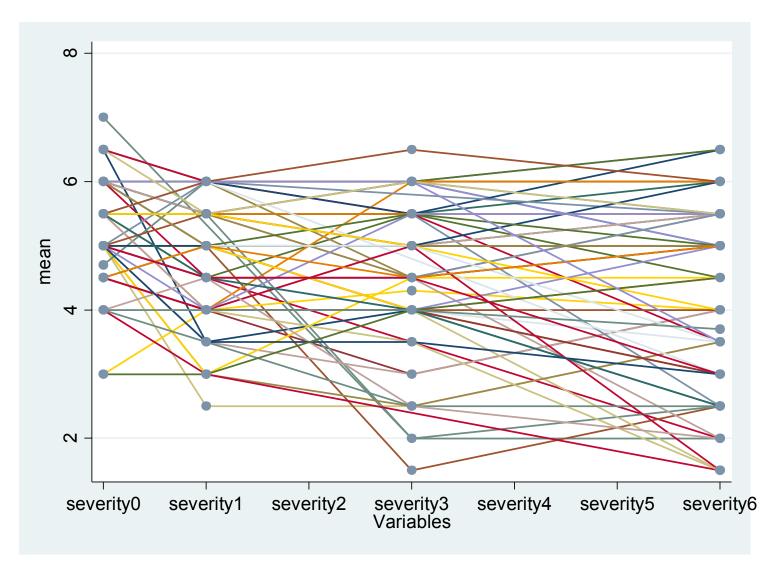
// this keeps individuals for whole we have relevant observations

keep if severity0 != . & severity6 != .

Plot of the new data

- Let's look at our data again. Previously, we made a plot of data in the long format. We can use a Stata user-written function called "profileplot" to get a general idea of what the remaining data looks like in "wide" format.
- To install "profileplot," run findit profileplot in the command window and follow the instructions
- General because profile plot groups similar observations...pros and cons?

profileplot severity0 severity1 severity2 severity3 severity4 severity5 severity6 if drug0==0, by(id) legend(off)



Statistical Analysis

Now that we have visualized our data, we would like some statistical way to answer the following question:

Is there a difference in severity of episodes between the two treatment groups (being on the drug vs. off of it)? I.e. does the drug make a difference?

Two Sample T-test reminder

- Why do you do a 2 sample t-test?
 - Compare responses from two groups (Are women and men different heights? Did a cholesterol reduction drug actually work?)
- Necessary assumptions:
 - Each group is considered to be a sample from two separate populations (pop 1: treated with drug, pop 2: not treated with drug)
 - Responses are independent
 - Distribution of outcome variable is approximately normal (use a histogram to see this)

Two Sample T-test reminder

- H_0 : $\mu_1 = \mu_2$ or equivalently $\mu_1 \mu_2 = 0$ (The two population means are equal)
- Alternate hypothesis can be one-sided or two-sided (≥, ≤, or ≠)
- Test statistic:

$$t = \frac{\chi_1 - \chi_2}{\sqrt{s_1^2 / n_1 + s_2^2 / n_2}}$$

 Compare to a t-distribution with the smaller of n1-1 or n2-1 degrees of freedom (or if n1=n2, use n1+n2-2 degrees of freedom) to get a p-value

Run a t-test

```
keep severity0 drug0 gender0 severity6
```

gen difference = severity6 - severity0

ttest difference, by(drug0)

Two-sample t test with equal variances

Other options

 You have slopes for each of your treated and untreated individuals. What about a t-test to test for a difference in the mean values of the slopes?

 You could also approach this question using the regression models we are covering in lecture. We will discuss this next week, but feel free to start now.

Your turn

We've looked at the data by drug treatment. Now, can you perform a similar analysis looking to see if there's a gender effect? Some ideas

- Create a scatterplot and a boxplot demonstrating the change over time of severity in each individual, conditioning on gender.
- Put the data in wide format and create a profileplot of each of the genders.
- Examine the difference in severity of attacks between weeks 0 and 6 between genders. Do a t-test. What do your results suggest?

Modeling

We've looked at graphics and summary measures, and thought about how to test for differences using summary measures.

Why would you want to model?

Modeling

- We can include other variables
- Examine different assumptions and related variances
- Eventually, different mean models
- What models might you try with this data?

What models might you try on this data?

- Cross sectional model regressing strength on time and treatment
- Cross sectional model with robust standard errors
- Including gender
- Others?

Are these standard errors reasonable?

. regress severity week drug

Source	SS	df	MS		Number of obs F(2, 1600)		1603 302.24
Model Residual	951.170459 2517.62532	2 1600	475.58523 1.57351583		Prob > F R-squared Adj R-squared	=	0.0000 0.2742 0.2733
Total	3468.79578	1602	2.16529075		Root MSE	=	1.2544
severity	Coef.	Std. E	rr. t	P> t	[95% Conf.	Ιn	terval]
week drug _cons	3207993 6397973 5.597435	.01413 .07385 .07134	03 -8.66	0.000	3485266 7846508 5.457505		2930719 4949438 .737365

Is this reasonable?

. regress severity week drug

Source	SS	d f		MS		Number of obs F(2, 1600)		1603 302.24
Model Residual	951.170459 2517.62532	2 1600		.58523 351583		Prob > F R-squared Adj R-squared	=	0.0000 0.2742 0.2733
Total	3468.79578	1602	2.16	529075		Root MSE	=	1.2544
severity	Coef.	Std.	Err.	t	P> t	[95% Conf.	Ιn	terval]
week drug _cons	3207993 6397973 5.597435	.0141 .0738 .0713	503	-22.69 -8.66 78.46	0.000 0.000 0.000	3485266 7846508 5.457505		2930719 4949438 .737365

Is this reasonable?

. regress severity week drug, robust

Linear regression

Number of obs = 1603 F(2, 1600) = 301.24 Prob > F = 0.0000 R-squared = 0.2742 Root MSE = 1.2544

severity	Coef.	Robust Std. Err.	t	P> t	[95% Conf.	. Interval]
week	3207993	.0145419	-22.06	0.000	3493223	2922762
drug	6397973	.070897	-9.02	0.000	7788581	5007366
_cons	5.597435	.0635448	88.09	0.000	5.472796	5.722075

Adding gender

. regress severity week drug gender

Source	SS	df		MS		Number of obs F(3, 1599)		1603 201.69
Model Residual	952.266632 2516.52915	3 1599		422211 381435		Prob > F R-squared Adj R-squared	=	0.0000 0.2745 0.2732
Total	3468.79578	1602	2.16	529075		Root MSE	=	1.2545
severity	Coef.	Std.	Err.	t	P> t	[95% Conf.	Ιn	terval]
week drug gender _cons	3209089 6412884 .0523612 5.573707	.0141 .0738 .0627 .0768	789 403	-22.70 -8.68 0.83 72.57	0.000 0.000 0.404 0.000	3486401 7861981 0707007 5.423062	-: :	2931777 4963787 1754231 .724353

Adding gender

. regress severity week drug gender, robust

Linear regression

Number of obs = 1603 F(3, 1599) = 201.25 Prob > F = 0.0000 R-squared = 0.2745 Root MSE = 1.2545

severity	Coef.	Robust Std. Err.	t	P> t	[95% Conf.	Interval]
week	3209089	.0145409	-22.07	0.000	3494301	2923877
drug	6412884	.0710023	-9.03	0.000	7805559	502021
gender	.0523612	.0626472	0.84	0.403	070518	.1752404
_cons	5.573707	.0702801	79.31	0.000	5.435857	5.711558

Any other ideas?