

My basic Stata companion

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Research data and statistics, AaUH

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Section 1

Start

Getting started

Start by seeing the following videos:

- The user interface
 - ▶ Stata youtube channel: Tour of the Stata user interface (3:39m)
 - ▶ Amy Penn: Introduction to Stata - Getting started (5:01m)
 - ▶ Amy Penn: Opening a dataset (1:11m)
- Using Stata
 - ▶ Amy Penn: Introduction to Stata - Thinking like Stata (14:58m)
 - ★ The link for the example dataset does not work. Try the example dataset used later in this presentation
 - ▶ Amy Penn: Introduction to Stata - Data Cleaning using the Codebook and Sort Commands (3:10m)
 - ▶ Amy Penn: Introduction to Stata - Generating variables using the generate, replace, and label commands (8:30m)
- Saving commands and logging output
 - ▶ Amy Penn: Introduction to Stata - Do Files & Log Files (5:10m)

help, the most important command

help|h [command|keywords]

- Reading syntax
 - ▷ **first|second|third**: choose one of the three
 - ▷ **[]**: Within brackets means optional
 - ▷ **if**: a qualifier limiting the scope of the command, eg if male == 1
 - ▷ **in first/last**: select only rows with row numbers between first and last

Section 2

Example data and setting up a project

Example project

Infant mortality rates and birth defect rates are very high for low birth weight babies.

Hence, low birth weight is an outcome that has been of concern to physicians for years.

The aim is to see if a set of variables has an effect on birth weights

```
cd [""] [directory] [""]
```

cd changes the current working directory to the specified drive and directory.

Although optional, it is recommended always to use quotations marks ("")

Examples:

- Getting (my) current directory in the Result window

```
cd
```

```
C:\Users\sttp\Documents\nhb\STATA\Presentations\2020-02 RN Course material
```

- Setting (my) current directory in the Result window

```
cd "C:\Users\sttp\Documents\nhb\STATA\Presentations\2020-02 RN Course material"
```

```
C:\Users\sttp\Documents\nhb\STATA\Presentations\2020-02 RN Course material
```

`mkdir` and `dir`

Examples:

- Create a sub directory

```
mkdir "Smoking effect on low birth weight"
```

- Check if sub directory is created

```
dir  
<dir> 10/19/20 13:07 .  
<dir> 10/19/20 13:07 ..  
133.0k 10/19/20 13:05 bwt_by_m_age.png  
93.5k 10/19/20 13:05 bwt_hist.png  
9.5k 9/14/20 8:22 Course plan.xlsx  
<dir> 9/26/20 18:23 data  
<dir> 10/19/20 13:07 do-files  
<dir> 10/09/20 18:08 documents  
5.2k 10/19/20 13:05 my lbt.dta  
<dir> 10/19/20 13:05 output  
<dir> 9/26/20 18:23 RawData  
<dir> 10/19/20 13:07 Smoking effect on low birth weight
```

- Change to sub directory

```
cd "Smoking effect on low birth weight"
```

```
C:\Users\sttp\Documents\nhb\STATA\Presentations\2020-02 RN Course material\Smoking effect on low birth weight
```

Section 3

Important Stata commands

use

```
use ["filename"] [, clear]
```

Load dataset *filename* into the data editor.

Option **clear** is needed to empty the data editor.

Although optional, it is recommended always to use quotations marks ("")

Examples:

- Retrieving a dataset for analysing low birth weight

```
use "https://www.stata-press.com/data/r16/lbw", clear  
(Hosmer & Lemeshow data)
```

keep|drop

keep|drop [varlist|if]

keep or drop variables or rows satisfying if-expression

Examples:

- Keep the necessary variables *bwt*, *low*, *smoke* *age* and *race*

```
keep bwt low smoke age race
```

save

```
save [""]filename[""] [, replace]
```

Saves dataset *filename* into current directory.

Option **replace** means replacing/overwriting a possible existing dataset.

Although optional, it is recommended always to use quotations marks ("")

Examples:

- Saving my dataset (names always in quotes)

```
save "my lbt.dta", replace  
file my lbt.dta saved
```

codebook 1/2

`codebook [varlist] [, compact]`

codebook examines the variable names, labels, and data to produce a codebook describing the dataset.
Option **compact** makes the description short and in one table

Examples:

- Seeing all variables and some of their characteristics

`codebook, compact`

Variable	Obs	Unique	Mean	Min	Max	Label
low	189	2	.3121693	0	1	birthweight<2500g
age	189	24	23.2381	14	45	age of mother
race	189	3	1.846561	1	3	race
smoke	189	2	.3915344	0	1	smoked during pregnancy
bwt	189	133	2944.286	709	4990	birthweight (grams)

codebook 2/2

Examples:

- Seeing some of their characteristics for variable *race*

```
codebook race
```

```
race
```

```
type: numeric (byte)
label: race

range: [1,3]                      units: 1
unique values: 3                  missing .: 0/189

tabulation: Freq.    Numeric   Label
            96        1  white
            26        2  black
            67        3  other
```

An overview of logical expressions

Symbol	Meaning
>	Greater than
<	Lesser than
>=	Greater than or equal
<=	Lesser than or equal
==	Equal to
!=	Not equal to
&	Logical and
	Logical or
!	Logical not
inlist(value1, value2, ...)	Value1 is equal to one of the following values

Example code	Meaning
... if age >= 20	Mothers of age 20 and above
... if inlist(race, 1, 2)	Mothers of race white (1) or black (2)

generate

```
generate|egen new_varname =exp [if]
```

generate|replace creates a new variable. The values of the variable are specified by =exp and possibly [if].

Examples:

- Generate variable *bwtlt1500* being 1 if children have a birth weight less than 1500 and zero otherwise if a value for birth weight exists

```
generate bwlt1500 = bwt < 1500 if !missing(bwt)
```

rename

rename

rename renames one or more variables.

Examples:

- Rename variable *low* to *bwt2500* (Birth Weight Less Than 2500)

```
rename low bwt2500
```

labels

```
labels define|values|variable
```

labels adds or modifies variable and value labels.

Labels can have different values at different times. In datasets they contain information on content, but they can be modified when variables are used in tables or graphs.

Examples:

- Adding labels to variables *bwlt1500* and *bwlt2500*

```
label variable bwlt2500 "Birth weight < 2500g"  
label variable bwlt1500 "Birth weight < 1500g"
```

- Defining a value label and attach it to variables *smoke*, *bwlt1500* and *bwlt2500*
 - ▶ *bwlt** means all variables starting with “bwlt”

```
label define no_yes 0 "no" 1 "yes"  
label values smoke bwlt* no_yes
```

summarize

summarize [varlist] [, detail]

summarize calculates and displays a variety of univariate summary statistics.

Option **detail** gives a detailed summary of the variables in *varlist*

Examples:

- A detailed summary over variable *bwt*

summarize bwt, detail			
birthweight (grams)			
		Percentiles	Smallest
1%	1021		709
5%	1790		1021
10%	1970	1135	Obs 189
25%	2414	1330	Sum of Wgt. 189
50%	2977		Mean 2944.286
		Largest	Std. Dev. 729.016
75%	3475	4174	
90%	3884	4238	Variance 531464.4
95%	3997	4593	Skewness -.2069782
99%	4593	4990	Kurtosis 2.888821

tabulate 1/3

`tabulate|tab1 varname [if] [, sort]`

One-way count table for variable

Option **sort** sort rows in descending order based on frequency

`tabulate|tab2 varname1 varname2 [if] [, chi2 exact]`

Two-way table for pairwise count combinations

Option **chi2** adds Pearson's chisquare test for row and column dependency

Option **exact** adds Fisher's exact test for row and column dependency

tabulate 2/3

Examples:

- Oneway sorted (highest frequency first) count table for variable *race*

```
tabulate race, sort
```

race	Freq.	Percent	Cum.
white	96	50.79	50.79
other	67	35.45	86.24
black	26	13.76	100.00
Total	189	100.00	

tabulate 3/3

Examples:

- Pearson's chisquare and Fisher's exact test for dependence between variables *smoke* and *race*

```
tabulate race smoke, chi2 exact
```

Enumerating sample-space combinations:

stage 3: enumerations = 1

stage 2: enumerations = 21

stage 1: enumerations = 0

race	smoked during pregnancy		Total
	no	yes	
white	44	52	96
black	16	10	26
other	55	12	67
Total	115	74	189

```
Pearson chi2(2) = 21.7790  Pr = 0.000  
Fisher's exact = 0.000
```

ttest

`ttest varname [if] [, by(groupvar)]`

Two-sample t test by a binary group variable

Examples:

- Testing equal mean birth weights between smokers and non-smokers

```
ttest bwt, by(smoke)
```

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]
no	115	3054.957	70.1625	752.409	2915.965 - 3193.948
yes	74	2772.297	76.70106	659.8075	2619.432 - 2925.162
combined	189	2944.286	53.02811	729.016	2839.679 - 3048.892
diff		282.6592	106.9544		71.66693 - 493.6515
diff = mean(no) - mean(yes)			t = 2.6428		
Ho: diff = 0			degrees of freedom = 187		
Ha: diff < 0		Ha: diff != 0		Ha: diff > 0	
Pr(T < t) = 0.9955		Pr(T > t) = 0.0089		Pr(T > t) = 0.0045	

We reject the Ho-hypothesis of no evidence of a weight difference against the two-sided alternative, P-value = 0.01.
The expected difference in birthweight is 282.659kg and the 95% confidence interval is [71.667; 493.652] (kg).

`cs outcomevar exposurevar [if] [, or exact]`

cs is used with cohort study data with equal follow-up time per subject and sometimes with cross-sectional data. Risk is then the proportion of subjects who become cases.

Option **or** makes the command return odds ratio instead of relative risk

Option **exact** adds Fisher's exact test for equal outcome rates between the exposure groups.

Examples:

- Estimation of relative risk of birthweight below 2500g for smokers vs non-smokers with 95% confidence interval.

smoked during pregnancy			
	Exposed	Unexposed	Total
Cases	30	29	59
Noncases	44	86	130
Total	74	115	189
Risk	.4054054	.2521739	.3121693
	Point estimate	[95% Conf. Interval]	
Risk difference	.1532315	.0160718 .2903912	
Risk ratio	1.607642	1.057812 2.443262	
Attr. frac. ex.	.377971	.0546528 .5907112	
Attr. frac. pop	.1921887		
chi2(1) = 4.92 Pr>chi2 = 0.0265			

We reject the Ho-hypothesis of same risk of birth weight below 2500g for smokers and non-smokers against the two-sided alternative, P-value = 0.03.

The expected risk of birth weight below 2500g is 1.61 times higher for smokers than for non-smokers. The 95% confidence interval for the relative risk is [1.06; 2.44].

Section 4

Graphing in Stata

histogram|hist

`histogram varname [, norm]`

`histogram` generates a histogram for a variable named `varname`.

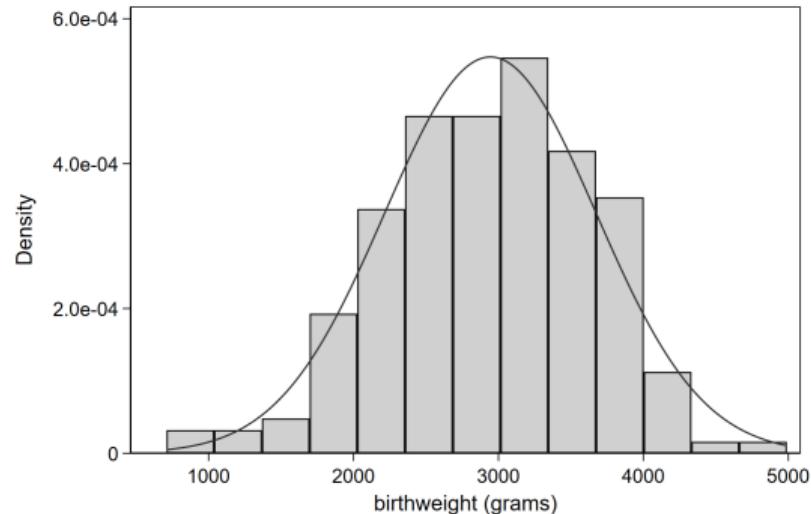
Option **norm** adds best fitting normal curve. For testing if data is normal distributed

Examples:

- Check if birthweight normally distributed

`histogram bwt, norm`

(`bin=13, start=709, width=329.30769`)



scatter

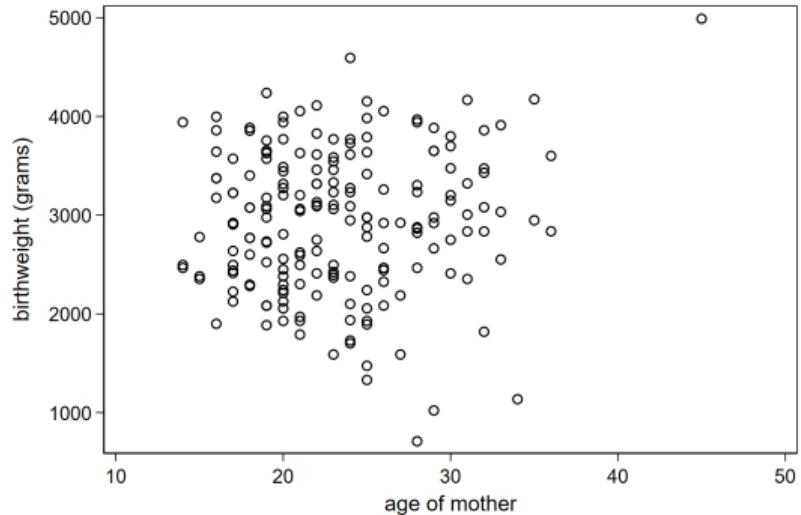
`scatter y-varname x-varname'`

scatter plots pairs of x and y values.

Examples:

- See dependency how mothers age affect birthweight

`scatter bwt age`



Section 5

Where to go now?

Useful stuff

Statistics

- seanharrisonblog.com: Series on Evaluation of Scientific Publications

Stata

- Stata Youtube Channel - Videos on usage
- UCLA IDRE (Institute for Digital Research and Education)
- Survey Design and Analysis Services: Tips on graphics
- Stata cheat sheets

Books

- Kirkwood and Sterne (2003)
- Peacock, Kerry, and Balise (2017)

References

- Kirkwood, B., and J. Sterne. 2003. *Essential Medical Statistics*. Wiley.
- Peacock, Janet L., Sally M. Kerry, and Raymond R. Balise. 2017. *Presenting Medical Statistics from Proposal to Publication*. Oxford, UK: Oxford University Press.
<https://oxfordmedicine.com/view/10.1093/med/9780198779100.001.0001/med-9780198779100>.