

Applied Epidemiology I: Data clearance

A review of using Stata

Enoch Yi-Tung Chen

Department of Medical Epidemiology and Biostatistics, Karolinska Institutet

November 26, 2020

Acknowledgements

This course material is based on my learning from Anastasia Lam's teachings in last year's Applied Epidemiology I lab sessions, and readings from *Epidemiology* by Gordis [?], *A First Course in Probability and Statistics* by Goldsman and Goldsman [?], *Principles of Biostatistics* by Pagano and Gauvreau [?], and *Biostatistics I* by Gabriel and Frumento [?].

I especially want to thank Marlene Stratmann for reviewing the slides and Prof. Paul Dickman for providing me with suggestions to improving the teaching.

Outline

Set up working directory

- Working directory is the folder where all your files are stored, and should be set each time you start.

Set up working directory

- Working directory is the folder where all your files are stored, and should be set each time you start.

- Where is it?

```
. cd
```

```
/Users/Desktop
```

```
. pwd
```

```
/Users/Desktop
```

Set up working directory

- Working directory is the folder where all your files are stored, and should be set each time you start.

- Where is it?

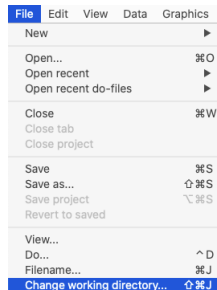
```
. cd
```

```
/Users/Desktop
```

```
. pwd
```

```
/Users/Desktop
```

- Change working directory
 - `cd "/Users/Download"`
 - Click File - Change Working Directory



Import and save data: Import

- **Excel (.xls or .xlsx)**

```
import excel filename, clear firstrow
```

- **Delimited (.csv) or text (.txt)**

```
import delimited filename, clear  
infile filename, clear
```

- **Stata (.dta)**

```
use filename, clear
```

- **SAS (.xpt)**

```
fdause filename, clear
```

- Save your dataset as a Stata file .dta
- The replace option lets you overwrite the existing dataset.
`save "filename", replace`

Manage datasets: Merge

merge adds new variables from a second dataset to your existing dataset.
(Make the dataset wider)

```
. sysuse cancer, clear  
(Patient Survival in Drug Trial)
```

```
. gen id = _n
```

```
. keep id
```

```
. merge 1:1 id using cancer
```

Result	# of obs.
not matched	0
matched	48 (_merge==3)

Manage datasets: Append

append adds new observations to existing variables in your current dataset.
(Make the dataset longer)

```
. use cancer_drug12, clear
```

(Patient Survival in Drug Trial)

```
. append using cancer_drug3.dta // append patients using drug 3
```

Get to know the data: Summarize

`summarize` gives summaries for all your variables, such as number of observations, mean, standard deviation, etc.

```
. sysuse cancer, clear  
(Patient Survival in Drug Trial)  
  
. keep if drug ==1 | drug == 2  
(14 observations deleted)  
  
. summarize age // One variable only (age)
```

Variable	Obs	Mean	Std. Dev.	Min	Max
age	34	56.41176	6.010686	47	67

Get to know the data: Describe

`describe` gives descriptions for all your variables, such as storage type and labels.

```
. describe age
```

variable name	storage type	display format	value label
age	byte	%8.0g	Patient's age at start of exp.

Get to know the data: Codebook

codebook is a combination of summarize and describe and will give a detailed summary of all your variables, including mean, sd, range, percentiles, missing, frequency, etc.

```
. codebook age
```

age	Patient's age at start of exp.
-----	--------------------------------

```
      type:  numeric (byte)
      range:  [47,67]
unique values: 15
      units:  1
      missing.: 0/34
      mean:   56.4118
      std. dev: 6.01069
percentiles:  10%      25%      50%      75%      90%
               49       51       56       61       65
```

Get to know the data: List

`list` lists the observations of specified variables.

```
. list      age if age < 50
```

```
    +-----+  
    |  age  |  
    +-----+  
12. |   49  |  
15. |   49  |  
18. |   49  |  
25. |   49  |  
33. |   47  |  
    +-----+
```

Manage variables: Numeric and string

Numeric: byte, integer, long, float, double – all types of numeric variables that just differ based on min and max length

String: character variables with a certain length (*str#*)

Manage variables: Drop/Keep

- drop is used to delete variables or observations.
- keep is used to keep variables or observations.





```
. sysuse cancer, clear  
(Patient Survival in Drug Trial)  
. drop if drug ==1 | drug == 2  
(34 observations deleted)
```

```
. sysuse cancer, clear  
(Patient Survival in Drug Trial)  
. keep if drug ==1 | drug == 2 // So drug == 3 will be dropped  
(14 observations deleted)
```


Manage variables: Label

- label helps you keep track of your dataset and variables, and helps others understand your data.

```
. // Label a dataset  
. label data "cancerdata"
```

Properties    	
► Variables	
▼ Data	
Frame	default
► Filename	cancer.dta
Label	cancerdata
Notes	

```
. // Label variable in the "Variables" window  
. label variable drug "1=placebo, 2=mild, 3=strong"
```

Variables		
	Name	Label
<input checked="" type="checkbox"/>	studytime	Months to death or end of exp.
<input checked="" type="checkbox"/>	died	1 if patient died
<input checked="" type="checkbox"/>	drug	1=placebo, 2=mild, 3=strong

Manage variables: Label

```
. // Label define claims the value label  
. label define drug_label 1 "placebo" 2 "mild" 3 "strong"  
  
. // Label value then assigns the label to the variables  
. label values drug drug_label
```

drug[1]			1
	studytime	died	drug
1	1	1	placebo
2	1	1	placebo

Manage variables: Rename, recode, generate, replace

- `rename` changes the name of a variable.
`. rename died death`
- `recode` changes variable values.
`. recode drug (3=4)`
- `generate` creates a new variable.
`. generate placebo = 1 if drug == 1`
- `replace` replaces existing variables (or variable values).
`. replace placebo = 0 if drug != 1`

Manage variables: Sort, by, if, in

- sort orders observations in ascending order.
`. sort death`
- by executes a command within a specified variable (e.g. by age group), but data should be sorted first.
`. by death: summarize`
- bysort combines the by and sort commands into one.
`. bysort death: summarize // by death, sort: summarize`
- if is used to select by a condition.
`. list age if death == 1`
- in is used to select by observations.
`. gen id = _n`
`. list id 1/10`

Manage variables: Operators

Operator	Purpose	Example
=	Sets equal operator	generate sex = 1
==	Tests for equality	summarize if sex==1
~= or !=	Indicates 'not equal'	summarize if sex!=0
<, <=	Less than (equal to) or greater	summarize if age<35
>, >=	than (equal to)	
&	Indicates 'and'	summarize outcome if sex==1 & age>=60
	Indicates 'or'	gen x=1 if a==1 & (b==1 c==1)

References

