Laboratory 1-2 - Notes Longitudinal Data Analysis Biostatistics 140.655

Topics:

- Reading/Loading Data Plotting Data and Exploring Correlation
- **Ordinary and Weighted Least Squares**

Reading/Loading Data

- Stata: infile (from text file), insheet (from spreadsheet), cut and paste
- SAS: infile (from text file)

data dataname; input varnames; infile 'filename';

first row of the text files, then use the input option. (see handout). To infile for SAS one should delete the variable names from the Data sets are available at network neighborhood in both Stata and Text formats

Examining Data

- Stata: use the editor; table or tabulate give frequencies
- Sas: PROC Print, PROC Freq

See also: Sas and Stata intro handouts.

Plotting Data / Exploring Correlation

- Strength
- Time Dependence
- **Modification**) **Inference Concerning Interaction (Effect**

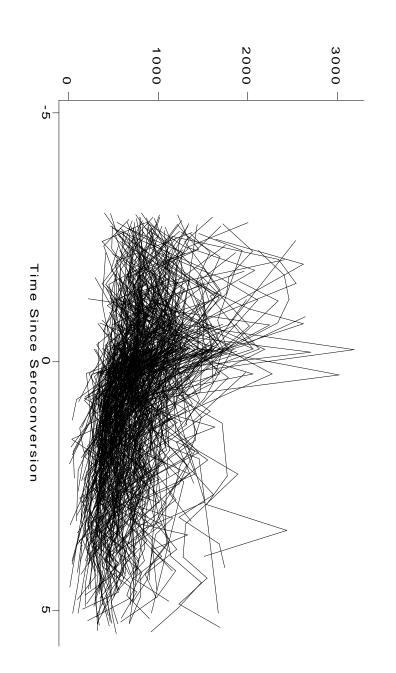
Creating a Longitudinal Plot (Stata)

- Graph command (see Stata Handout)
- Sort Data Points within Sampling Unit, then by time
- Connect Points within sampling Unit

Syntax:

form: command varlist, options

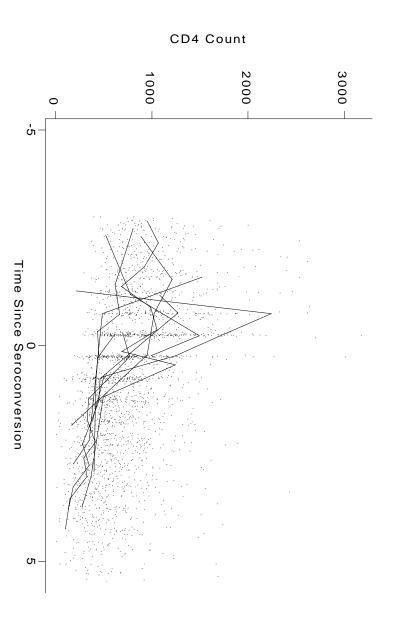
Label var time "Time Since Seroconversion" (labels time variable)



sort id time graph count time, connect(L) symbol(i) xlab ylab

The graph above may not be very useful - it's too noisy.





all individuals) Syntax: (this is a complicated instance - the time points are not the same for

sort id time * sorts by time inside individuals

observation gen first = id!=id[_n-1] *is it a new individual? Pick out each person's first

sort first count order people according to CD4 count *group all 'first' observations together; within this group,

gen decile = first & (int(_n/37)==_n/37)

only be the same as _n/37 if _n/37 is a whole number. (there are 369 "first" observations) The "int" function rounds the result of first observation AND the observation number (_n) is an even multiple of 37 what's in the parenthesis (n/37) to the nearest integer. Hence int(n/37) will * this is the toughest part - an observation is a "decile" obs. if it's a person's

our knowledge of the number of individuals. This statement is strongly dependent on the way we have ordered the data and

sort id time * re-order the data by individual

decile people, mark all of their observations that way. replace decile=decile[_n-1] if id==id[_n-1] *now that we know who are our

gen count2 = count if decile *make a variable of only the decile observations

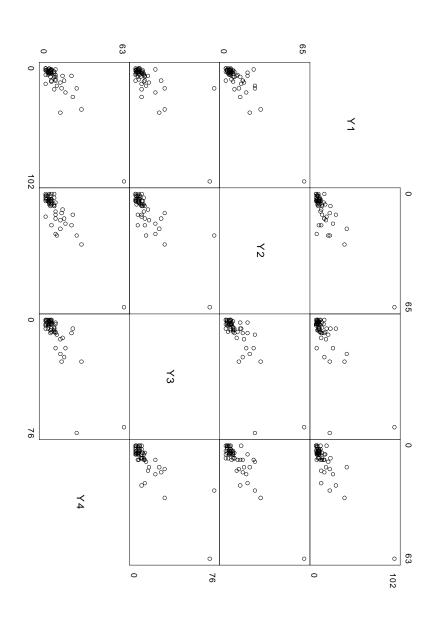
graph count count2 time, s(.i) c(.L) xlab ylab *make the plot

connect the decile points and not the others. Notice that we had to create an extra variable (count 2) so that we could

WHEW!!

SAS: similar difficulties. Explore PROC GPLOT.





To construct this plot, you need the data in WIDE format (see data handout).

syntax:

graph varlist, matrix

in this case:

graph Y1 Y2 Y3 Y4, matrix

Further exploration: Autocorrelation function.

function (see handout). Stata function autocor.ado calculates and plots autocorrelation

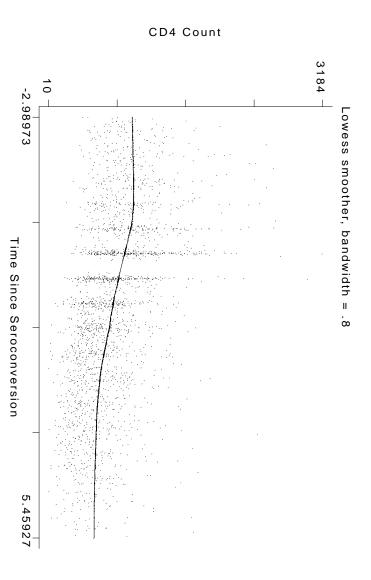
Residuals Plotting:

residuals. For some estimation commands, use PREDICT to construct

Then plot residuals vs. fitted values, predictors.

Graph Smoothing

- Various methods
- Stata: ksm statement: moving average, lowess smoothing
- SAS: PROC GPLOT, Symbol and smoothing options.



Least Squares Estimation

squares can be accomplished using vwls in Stata. most estimation commands in SAS; specifically variance weighted least Simple linear regression commands utilize OLS, including regress, fit (Stata) and PROC REG (SAS). Weighted least squares is an option in