Stata Cheat Sheet

Large Scale Data - Part 2

1. Basic Setup & Data Import

Standard Setup

// Clear memory and set options
clear all
set more off

// Set working directory
cd "path/to/folder"

// Start logging
capture log close
log using "analysis.log", replace

Import Data

// Import CSV
import delimited "file.csv", clear

// Import Excel
import excel "file.xlsx", ///
sheet("Sheet!") firstrow clear

// Import SAS XPT
import sasxport5 "file.xpt", clear

// Load Stata file
use "file.dta", clear

// Save data
save "filename.dta", replace

A Warning

Always use ${\tt clear}$ or , ${\tt clear}$ option when loading new data to avoid "no; data in memory would be lost" error.

2. Data Exploration

Basic Viewing

// View first 10 rows
list var1 var2 var3 in 1/10

// View specific observations
list if condition

// Browse data (opens viewer)
browse

// Describe variables
describe
describe var1 var2

// View variable type
codebook varname

Summary Statistics

// Basic summary
Summ varname
// Detailed summary (with percentiles)
summ varname, d
// Shows: p1, p5, p10, p25, p50, p75,
// p90, p95, p99
// Multiple variables
summ varl var2 var3, d

Frequency Tables

// Frequency table (include missing)
tab varname, m

// Show numeric codes (not labels)
tab varname, nolabel

// Two-way table with row %
tab var1 var2, row m

// Two-way table with column %
tab var1 var2, col m

// Both row and column %
tab var1 var2, row col

Check Data Size

// Number of observations
display _N
// Number of variables
describe, short

3. Variable Creation Generate New Variable

// Create empty variable gen newvar = . // Create with value gen age_squared = age^2 // Create constant gen constant = 1

Replace Values

// Replace all values
replace varname = new_value
// Conditional replace
replace var = value if condition

A Warning

CRITICAL: Always protect missing values!
Use: if var != . in conditions
Missing (.) is treated as infinity in Stata.

Binary Variables (0/1 Flags)

// Method 1: Standard approach
gen flag = 1 if condition
replace flag = 0 if !condition

// Method 2: One-liner
gen flag = (condition) if var != .

// Example: Age >= 18
gen adult = 1 if age >= 18 & age != .
replace adult = 0 if age < 18

// Example: High cost (>\$50,000)
gen expensive = 1 if cost > 50000 ///
& cost != .
replace expensive = 0 if cost <= 50000

// VERIFY binary variable
summ flag
// mean should be 0-1, min=0, max=1

Categorical Variables

// Method 1: Manual creation
gen category = 1 if condition1
replace category = 2 if condition2
replace category = 3 if condition3

// Add value labels
label define cat_lbl 1 "Low" ///
2 "Medium" 3 "High"
label values category cat_lbl

// Method 2: recode
recode age (0/29=1) (30/49=2) ///
(50/max=3), gen(age_group)

// Method 3: encode string variable
encode string_var, gen(numeric_var)

Top-coding (Capping Outliers)

// Create clean version
gen cost_clean = cost

// Top-code at \$1,000,000
replace cost_clean = 1000000 ///
 if cost > 1000000 & cost != .

// Verify
summ cost, d
summ cost_clean, d
// Check: max of clean version = cap

${\bf Logarithmic\ Transformation}$

// Handle right-skewed data
// Step 1: Add small value to avoid log(0)
gen cost_plus1 = cost + 1

// Step 2: Take log
gen log_cost = log(cost_plus1)

// Alternative: only for positive values
gen log_cost = log(cost) if cost > 0

EGEN - Extended Generation

// Row operations
egen total = rowtotal(var1 var2 var3)
egen mean = rowmean(var1 var2 var3)
egen max = rowmax(var1 var2 var3)
egen min = rowmin(var1 var2 var3)

// Grouped statistics
egen mean_by_group = mean(var), ///
by(group_var)

// Example: County-level averages
egen avg_income_county = mean(income), ///
by(county_name)

// Count non-missing
egen count_nonmiss = count(var), ///
by(group)

// String concatenation
egen fullname = concat(first last), ///
punct("")

4. Missing Values Identify Missing Values

// Count missing
count if varname == .

// Summary shows N
summ varname
// Total N us variable N

// Show missing in table
tab varname, m

Recode Missing Values

// Set specific values to missing
replace var = . if var == 99
replace var = . if var == 999
replace var = . if var < 0

// Example from BRFSS
replace height = . if height == 7777
replace height = . if height == 9999

▲ Warning

Common Mistake:

Common Mistake:

replace var = 100 if var > 100

This will set missing to 100!

Correct:

replace var = 100 if var > 100 & var != .

Create Missing Indicator

```
// Flag for missing
gen miss_flag = (varname == .)
// Or explicitly
gen miss_flag = 0
replace miss_flag = 1 if varname == .
```

5. String Variables String to Numeric

```
// Basic conversion (force ignores errors)
destring string_var, gen(num_var) force

// Check what couldn't be converted
tab string_var if num_var == .

// Example: Handle "120 +"
destring lengthofstay, gen(los_num) force
replace los_num = 120 //
if lengthofstay == "120 +"
```

Numeric to String

```
// Convert number to string
gen str_var = string(numeric_var)

// With formatting
gen str_var = string(num_var, "%9.2f")
```

String to Categorical (encode)

```
// Create numeric with labels
encode string_var, gen(categorical_var)

// Example: Admission type
encode typeofadmission, ///
gen(admission_type_num)

// Verify
tab typeofadmission admission_type_num
```

String Manipulation

▲ Warning

substr() position starts at 1! substr(str, 1, 2) = first 2 chars substr(str, 2, 1) = 2nd char only

6. Data Merging

Merge Types

```
// 1:1 - Both datasets: 1 row per ID
merge 1:1 id_var using "file.dta"

// 1:M - Master: 1 row/ID, Using: many rows/ID
merge 1:m id_var using "file.dta"

// M:1 - Master: many rows/ID, Using: 1 row/ID
merge m:1 id_var using "file.dta"

// M:M - Both: many rows/ID (rare, avoid)

// Use joinby instead if needed
```

M:1 Merge Example (Most Common)

```
// Merge county data to individual records
// Main: Individual hospitalizations
// (many records per county)
// Using: County characteristics
// (one record per county)

use "hospital_data.dta", clear
// Rename if needed to match
rename hospitalcounty County_Name
// Perform M:1 merge
merge m:1 County_Name ///
using "county_data.dta"
// CHECK merge results
tab _merge
/*
_merge values:
i = master only (hospital records
```

```
with no county match)

2 = using only (counties with no hospital records)

3 = matched successfully

*/

// Keep what you want
keep if _merge == 1 | _merge == 3

// Keeps all hospital records

// Clean up
drop _merge
```

Merge Workflow

```
// Step 1: Check merge variable exists
describe merge_var

// Step 2: Check if unique (if "1" side)
duplicates report merge_var

// Should show 0 duplicates

// Step 3: Ensure variable names match

// If not, rename in one dataset
rename old_name new_name

// Step 4: Check variable types match
describe merge_var

// Both should be numeric or string

// Step 5: Perform merge
merge type merge_var using "file.dta"

// Step 6: Always check _merge!
tab _merge

// Step 7: Keep desired records
keep if _merge == 1 | _merge == 3

// Step 8: Drop _merge
drop _merge
```

7 Tip

Quick merge type decision:

Master(M): Using(1) \rightarrow M:1 Master(1): Using(M) \rightarrow 1:M Master(1): Using(M) \rightarrow 1:M Master(1): Using(1) \rightarrow 1:1

Append (Stack Datasets)

```
// Combine datasets with same structure
use "data2020.dta", clear
append using "data2021.dta"
append using "data2022.dta"

// All rows are kept, stacked vertically
```

7. Data Reshaping Wide to Long

```
// Wide format:
// id bp1 bp2 bp3 hr1 hr2 hr3
// 1 120 118 115 72 70 68

reshape long bp hr, i(id) j(round)

// Long format:
// id round bp hr
// 1 1 120 72
// 1 2 118 70
// 1 3 115 68
```

Long to Wide

```
// Long format:
// id round bp hr
// il 1 120 72
// l 2 118 70

reshape wide bp hr, i(id) j(round)

// Wide format:
// id bp1 bp2 hr1 hr2
// i 120 118 72 70
```

▲ Warning

reshape permanently changes data.
Always save before reshaping!

8. Regression Analysis Linear Regression

```
// Simple regression
reg outcome predictor

// Multiple regression
reg y x1 x2 x3

// With robust standard errors
reg y x1 x2, robust
// or
reg y x1 x2, r
```

Categorical Variables in Regression

```
// Use i. prefix for categorical
reg outcome continuous_var i.category

// Example
reg totalcosts lengthofstay ///
    i.agegroup i.admission_type

// Stata automatically:
// - Creates dummy variables
// - Omits first category (reference)
// - Shows each category coefficient
```

Continuous Variables

// Default: continuous
reg y x1 x2
// Explicit: c. prefix (optional)
reg y c.x1 c.x2

Interaction Terms

// Categorical x Categorical
reg y i.var1##i.var2
// ## includes main effects + interaction
// Categorical x Continuous
// IMPORTANT: Use c. for continuous!
reg y i.category##c.continuous
// Example: Does income effect vary by sex?
reg health i.sex##c.income age
// Only interaction (no main effects)
reg y i.var1#i.var2

A Warning

Interaction with continuous:
MUST use c. prefix!
Wrong: i.sex##age
Right: i.sex##c.age

Logistic Regression

// Binary outcome (0/1)
logistic binary_y x1 x2 i.category
// Reports Odds Ratios (OR)

// Alternative: logit (reports log-odds)
logit binary_y x1 x2 i.category

// Example
logistic expensive_stay ///
County_Income lengthofstay ///
i.agegroup i.ED_flag

Linear vs Logistic Interpretation

// LINEAR regression on binary outcome reg expensive_stay County_Income, r // Coefficient: Percentage point difference // Example: coef = 0.04 // Interpretation: "County income increase // of \$1000 associated with 4 percentage // point increase in probability of // expensive stay (e.g., 10% to 14%)" // LOGISTIC regression logistic expensive_stay County_Income // Coefficient: Odds Ratio // Example: OR = 1.02 // Interpretation: "County income increase // of \$1000 associated with 2% increase // in the odds of expensive stay"

Survey Weights

// Set survey design
svyset [pweight = weight_var]

// Weighted regression
svy: reg y x1 x2 i.category

// Weighted logistic
svy: logistic binary_y x1 x2

// Why use weights?

// - Make results representative
// - Account for survey design
// - Adjust for non-response

Post-Regression Commands

// Test joint significance
reg y x1 x2 x3
test x1 x2
// Tests: x1 = x2 = 0
// Predicted values
predict yhat
// Residuals
predict resid, residuals
// Margins (adjusted predictions)
reg y x1 i.group
margins group
// Shows predicted y for each group

Individual Fixed Effects

// For panel/longitudinal data
// Controls for all time-invariant
// individual characteristics

// Set panel structure
xtset person_id time_var

// Fixed effects regression
xtreg y x1 x2, fe

// Why use FE?
// - Within-person analysis
// - Control for unmeasured confounders
// - Stronger causal inference

9. Verification & Validation Verify Binary Variables

// Create binary flag
gen flag = 1 if cost > 50000 & cost != .
replace flag = 0 if cost <= 50000</pre>

// CHECK 1: Summary statistics
summ flag
// mean: 0-1, min: 0, max: 1, N correct?

// CHECK 2: Cross-tabulation
tab flag, m
// Should show: 0, 1, and . only

// CHECK 3: Verify cutoff
summ cost if flag == 1
// min should be > 50000
summ cost if flag == 0
// max should be <= 50000

Verify Categorical Variables

// After encoding or recoding
tab old_var new_var
// Check mapping is correct
// With percentages
tab old_var new_var, row col

Verify Continuous Variables

// After transformation
summ original_var, d
summ clean_var, d
// Compare: mean, min, max, N
// Grouped summary
bysort group: summ var
// or
summ var if group == 1
summ var if group == 0

Check for Missing

// Count missing
count if var == .

// Identify observations with missing
list id var if var == .

// Missing by group
tab group, m
bysort group: count if var == .

Verify Merge Success

// After merge
tab _merge

// List unmatched from master
list id if _merge == 1

// List unmatched from using
list id if _merge == 2

// Check merged variable
summ merged_var if _merge == 3

// Should have valid values

10. Common Workflows Clean Outcome Variable

// Step 1: Explore
codebook outcome
summ outcome, d
tab outcome, m

// Step 2: Identify issues
// - Missing values?
// - Outliers?
// - Correct range?

// Step 3: Create clean version
gen outcome_clean = outcome

// Step 4: Handle missing
replace outcome_clean = . if outcome < 0

// Step 5: Handle outliers (top-code)
replace outcome_clean = 1000000 ///
if outcome > 1000000 & outcome != .

// Step 6: Verify
summ outcome_clean, d
tab outcome outcome_clean, m

Prepare Covariates

// Continuous variable
// - Check range
summ age, d
// - Handle missing
replace age = . if age == 99
// - Create squared term if needed
gen age_squared = age "2

// Categorical variable (numeric)
// - Check values
tab category, m
// - Create labeled version
label define cat_lbl 1 "A" 2 "B" 3 "C"
label values category cat_lbl

// Categorical variable (string)
// - Encode to numeric
encode string_var, gen(category_num)
// - Verfy
tab string_var category_num
// Binary flag
// - Create O/1
gen flag = 1 if condition & var != .
replace flag = 0 if !condition
// - Verfy
summ flag

Complete Analysis Example

11. Important Reminders

Critical Points

▲ Warning

Top 5 Common Mistakes:

- 1. Missing values: Always use & var != .
- 2. substr(): Position starts at 1, not 0
- 3. Interaction: Use ${\tt c.}$ for continuous
- 4. _merge: Always tab _merge after merge
- 5. Binary range: Check summ shows 0-1

Statistical Significance vs Practical

```
// Example: p=0.007, diff=3 minutes
// Statistical: YES (p<0.05)
// Practical: MAYBE
// - 3 min might be too small
// - But >10% relative difference
// - Large sample = "overpowered"
// (can detect tiny differences)
// Always discuss BOTH in interpretation
```

Regression Interpretation

// Linear regression coefficient: // "Each 1-unit increase in X is

```
// associated with beta-unit change in Y,
// controlling for other variables."

// Logistic regression OR:
// "Each I-unit increase in X is
// associated with ORs change in the
// odds of Y, controlling for others."

// Binary outcome + linear regression:
// "Each I-unit increase in X is
// associated with beta percentage point
// change in probability of Y."
```

When to Use What

Data Types:

- Cross-sectional survey: Prevalence, associations
- Longitudinal survey: Within-person changes, causality
- Claims data: Utilization, costs, readmission
- EHR data: Clinical details, single system

Merge Types:

- M:1: Individual records + area-level data
- 1:M: Visits + medications per visit
- 1:1: Same IDs in both datasets

Regression Types:

- Linear: Continuous outcome
- Linear: Binary outcome (percentage points)
- Logistic: Binary outcome (odds ratios)
- Fixed effects: Panel data, within-person

Cheat Sheet Usage Tips

Tip

4

During the exam:

- 1. Start with exploration (summ, tab, describe)
- 2. Create clean versions of variables
- 3. Verify each step before moving on
- 4. Check merge with tab _merge
- 5. Verify binary variables with summ
- 6. Write complete interpretations

Quick Reference

Task	Command
Load data	use "file.dta", clear
Import CSV	import delimited "file.csv"
Summary stats	summ var, d
Frequency	tab var, m
Create var	gen newvar = expr
Binary flag	gen flag = (condition)
Top-code	replace var = cap if var ; cap & var != .
Encode string	encode str, gen(num)
String to num	destring str, gen(num) force
M:1 merge	merge m:1 id using "file.dta"
Check merge	tab _merge
Linear reg	reg y x1 x2, r
With category	reg y x1 i.cat
Interaction	reg y i.cat1##c.cont
Logistic	logistic binary_y x1 x2
Verify binary	summ flag (should be 0-1)
Cross-check	tab oldvar newvar

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