

st_is — Survival analysis subroutines for programmers

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Syntax

Verify that data in memory are survival-time data

```
st_is 2 {full | analysis}
```

Display or do not display summary of survival-time variables

```
st_show [noshow]
```

Risk-group summaries

```
st_ct "[byvars]" -> newtvar newpopvar newfailvar [newcensvar [newentvar ]]
```

You must have `stset` your data before using `st_is`, `st_show`, and `st_ct`; see [\[ST\] stset](#).

Description

These commands are provided for programmers wishing to write new `st` commands.

`st_is` verifies that the data in memory are survival-time (`st`) data. If not, it issues the error message “data not st”, `r(119)`.

`st` is currently “release 2”, meaning that this is the second design of the system. Programs written for the previous release continue to work. (The previous release of `st` corresponds to Stata 5.)

Modern programs code `st_is 2 full` or `st_is 2 analysis`. `st_is 2` verifies that the dataset in memory is in release 2 format; if it is in the earlier format, it is converted to release 2 format. (Older programs simply code `st_is`. This verifies that no new features are `stset` about the data that would cause the old program to break.)

The `full` and `analysis` parts indicate whether the dataset may include past, future, or past and future data. Code `st_is 2 full` if the command is suitable for running on the analysis sample and the past and future data (many data management commands fall into this category). Code `st_is 2 analysis` if the command is suitable for use only with the analysis sample (most statistical commands fall into this category). See [\[ST\] stset](#) for the definitions of past and future.

`st_show` displays the summary of the survival-time variables or does nothing, depending on what you specify when `stsetting` the data. `noshow` requests that `st_show` display nothing.

`st_ct` is a low-level utility that provides risk-group summaries from survival-time data.

Remarks and examples

Remarks are presented under the following headings:

Definitions of characteristics and st variables
Outline of an st command
Using the st_ct utility
Comparison of st_ct with sttoct
Verifying data
Converting data

Definitions of characteristics and st variables

From a programmer's perspective, `st` is a set of conventions that specify where certain pieces of information are stored and how that information should be interpreted, together with a few subroutines that make it easier to follow the conventions.

At the lowest level, `st` is nothing more than a set of Stata characteristics that programmers may access:

<code>char _dta[_dta]</code>	<code>st</code> (marks that the data are <code>st</code>)
<code>char _dta[st_ver]</code>	2 (version number)
<code>char _dta[st_id]</code>	<i>varname</i> or nothing; <code>id()</code> variable
<code>char _dta[st_bt0]</code>	<i>varname</i> or nothing; <code>t0()</code> variable
<code>char _dta[st_bt]</code>	<i>varname</i> ; <i>t</i> variable from <code>stset t, ...</code>
<code>char _dta[st_bd]</code>	<i>varname</i> or nothing; <code>failure()</code> variable
<code>char _dta[st_ev]</code>	list of numbers or nothing; <i>numlist</i> from <code>failure(varname[==numlist])</code>
<code>char _dta[st_enter]</code>	contents of <code>enter()</code> or nothing; <i>numlist</i> expanded
<code>char _dta[st_exit]</code>	contents of <code>exit()</code> or nothing; <i>numlist</i> expanded
<code>char _dta[st_orig]</code>	contents of <code>origin()</code> or nothing; <i>numlist</i> expanded
<code>char _dta[st_bs]</code>	# or 1; <code>scale()</code> value
<code>char _dta[st_o]</code>	<code>_origin</code> or #
<code>char _dta[st_s]</code>	<code>_scale</code> or #
<code>char _dta[st_ifexp]</code>	<i>exp</i> or nothing; from <code>stset ... if exp ...</code>
<code>char _dta[st_if]</code>	<i>exp</i> or nothing; contents of <code>if()</code>
<code>char _dta[st_ever]</code>	<i>exp</i> or nothing; contents of <code>ever()</code>
<code>char _dta[st_never]</code>	<i>exp</i> or nothing; contents of <code>never()</code>
<code>char _dta[st_after]</code>	<i>exp</i> or nothing; contents of <code>after()</code>
<code>char _dta[st_befor]</code>	<i>exp</i> or nothing; contents of <code>before()</code>
<code>char _dta[st_wt]</code>	weight type or nothing; user-specified weight
<code>char _dta[st_wv]</code>	<i>varname</i> or nothing; user-specified weighting variable
<code>char _dta[st_w]</code>	<code>[weighttype=weightvar]</code> or nothing
<code>char _dta[st_show]</code>	<code>noshow</code> or nothing
<code>char _dta[st_t]</code>	<code>_t</code> (for compatibility with release 1)
<code>char _dta[st_t0]</code>	<code>_t0</code> (for compatibility with release 1)
<code>char _dta[st_d]</code>	<code>_d</code> (for compatibility with release 1)
<code>char _dta[st_n0]</code>	# or nothing; number of <code>st</code> notes
<code>char _dta[st_n1]</code>	text of first note or nothing
<code>char _dta[st_n2]</code>	text of second note or nothing
<code>char _dta[st_set]</code>	text or nothing. If filled in, <code>streset</code> (see [ST] stset) will refuse to execute and present this text as the reason

All st datasets also have the following four variables:

```

    _t0  time of entry (in t units) into risk pool
    _t   time of exit (in t units) from risk pool
    _d   contains 1 if failure, 0 if censoring
    _st  contains 1 if observation is to be used and 0 otherwise

```

Thus, in a program, you might code

```

display "the failure/censoring base time variable is _t"
display "and its mean in the uncensored subsample is"
summarize _t if _d

```

No matter how simple or complicated the data, these four variables exist and are filled in. For instance, in simple data, `_t0` might contain 0 for every observation, and `_d` might always contain 1.

Some st datasets also contain the variables

```

    _origin  evaluated value of origin()
    _scale   evaluated value of scale()

```

The `_dta[st_o]` characteristic contains either the name `_origin` or a number, often 0. It contains a number when the origin does not vary across observations. `_dta[st_s]` works the same way with the `scale()` value. Thus the origin and scale are `_dta[st_o]` and `_dta[st_s]`. In fact, these characteristics are seldom used because variables `_t` and `_t0` are already adjusted.

Some st datasets have an `id()` variable that clusters together records on the same subject. The name of the variable varies, and the name can be obtained from the `_dta[st_id]` characteristic. If there is no `id()` variable, the characteristic contains nothing.

Outline of an st command

If you are writing a new st command, place `st_is` near the top of your code to ensure that your command does not execute on inappropriate data. Also place `st_show` following the parsing of your command's syntax to display the key st variables. The minimal outline for an st command is

```

program st name
    version 13
    st_is 2 ...
    ... syntax command ...
    ... determined there are no syntax errors ...
    st_show
    ... guts of program ...
end

```

`st_is 2` appears even before the input is parsed. This is to avoid irritating users when they type a command, get a syntax error, work hard to eliminate the error, and then learn that "data not st".

A fuller outline for an st command, particularly one that performs analysis on the data, is

```

program st name
    version 13
    st_is 2 ...
    syntax ... [, ... noSHow ... ]
    st_show 'show'
    marksample touse
    quietly replace 'touse' = 0 if _st==0
    ... guts of program ...
end

```

All calculations and actions are to be restricted, at the least, to observations for which `_st` \neq 0. Observations with `_st` = 0 are to be ignored.

Using the `st_ct` utility

`st_ct` converts the data in memory to observations containing summaries of risk groups. Consider the code

```
st_is 2 analysis
preserve
st_ct "" -> t pop die
```

Typing this would change the data in memory to contain something akin to count-time data. The transformed data would have observations containing

```
t      time
pop    population at risk at time t
die    number who fail at time t
```

There would be one record per time `t`, and the data would be sorted by `t`. The original data are discarded, which is why you should code `preserve`; see [P] [preserve](#).

The above three lines of code could be used as the basis for calculating the Kaplan–Meier product-limit survivor-function estimate. The rest of the code is

```
keep if die
gen double hazard = die/pop
gen double km     = 1-hazard if _n==1
replace km       = (1-hazard)*km[_n-1] if _n>1
```

`st_ct` can be used to obtain risk groups separately for subgroups of the population. The code

```
st_is 2 analysis
preserve
st_ct "race sex" -> t pop die
```

would change the data in memory to contain

```
race
sex
t      time
pop    population at risk at time t
die    number who fail at time t
```

There would be one observation for each `race–sex–t` combination, and the data would be sorted by `race sex t`.

With this dataset, you could calculate the Kaplan–Meier product-limit survivor-function estimate for each `race–sex` group by coding

```
keep if die
gen double hazard = die/pop
by race sex: gen double km = 1-hazard if _n==1
by race sex: replace km = (1-hazard)*km[_n-1] if _n>1
```

`st_ct` is a convenient subroutine. The above code fragment works regardless of the complexity of the underlying survival-time data. It does not matter whether there is one record per subject, no censoring, and one failure per subject, or multiple records per subject, gaps, and recurring failures for the same subject. `st_ct` forms risk groups that summarize the events recorded by the data.

`st_ct` can provide the number of censored records and the number who enter the risk group. The code

```
st_ct "" -> t pop die cens ent
```

creates records containing

```

t      time
pop    population at risk at time t
die    number who fail at time t
cens   number who are censored at t (after the failures)
ent    number who enter at t (after the censorings)

```

As before,

```
st_ct "race sex" -> t pop die cens ent
```

would create a similar dataset with records for each `race-sex` group.

Comparison of `st_ct` with `sttoct`

`sttoct`—see [ST] [sttoct](#)—is related to `st_ct`, and in fact, `sttoct` is implemented in terms of `st_ct`. The differences between them are that

- `sttoct` creates ct data, meaning that the dataset is marked as being ct. `st_ct` merely creates a useful dataset; it does not `ctset` the data.
- `st_ct` creates a total population at-risk variable—which is useful in programming—but `sttoct` creates no such variable.
- `sttoct` eliminates thrashings—censorings and reentries of the same subject as covariates change—if there are no gaps, strata shifting, etc. `st_ct` does not do this. Thus, at a particular time, `sttoct` might show that there are two lost to censoring and none entered, whereas `st_ct` might show 12 censorings and 10 entries. This makes no difference in calculating the number at risk and the number who fail, which are the major ingredients in survival calculations.
- `st_ct` is faster.

Verifying data

As long as you code `st_is` at the top of your program, you need not verify the consistency of the data. That is, you need not verify that subjects do not fail before they enter, etc.

The dataset is verified when you `stset` it. If you make a substantive change to the data, you must rerun `stset` (which can be done by typing `stset` or `streset` without arguments) to reverify that all is well.

Converting data

If you write a program that converts the data from one form of st data to another, or from st data to something else, be sure to issue the appropriate `stset` command. For instance, a command we have written, `stbase`, converts the data from st to a simple cross-section in one instance. In our program, we coded `stset`, `clear` so that all other st commands would know that these are no longer st data and that making st calculations on them would be inappropriate.

Even if we had forgotten, other st programs would have found many of the key st variables missing and would have ended with a “[such-and-such] not found” error.

Also see

[ST] [stset](#) — Declare data to be survival-time data

[ST] [sttoct](#) — Convert survival-time data to count-time data

[ST] [survival analysis](#) — Introduction to survival analysis & epidemiological tables commands