Multiple failure-time data

Multiple failure-time data or multivariate survival data are frequently encountered in biomedical and other investigations. These data arise from time-to-event studies when either of two or more events (failures) occur for the same subject, or from identical events occurring to related subjects. In these studies, failure times are correlated within subject, violating the independence of failure times assumption required in traditional survival analysis.

We follow Therneau's (1997) suggestion that for analysis purposes, failure events should be classified according to

- Whether they have a natural order
- Whether they are recurrences of the same types of events.

The counting process approach to survival analysis

A general approach to survival analysis was introduced by Andersen & Gill (1982) where each subject is considered as a counting process (counting events)

- N_i^(k)(t) is the total number of events of type k for each subject
 i up to time t
- $Y_i^{(k)}(t)$ is an indicator function with $Y_{ik}(t) = 1$ if subject *i* is at risk at time *t* for event of type *k*

In this formulation the hazard is considered as an "intensity" process such that

$$\lambda_i^{(k)}(t) = Y_i^{(k)}(t)\lambda_0^{(k)}(t)\exp\{\beta' Z_i\}$$

By judicious choice of the various components of the process as defined above, the counting process approach can handle all kinds of survival data including

- Time updated covariates $Z_i(t)$
- Discontinuous risk sets
- Multiple failures of the different type (competing risks)
- Multiple failures of the same type (both ordered and unordered)

Unordered failures

Failures of the same type include, for example, repeated lung infections with pseudomonas in children with cystic fibrosis, or the development of breast cancer in genetically predisposed families.

Failures of different types include adverse reactions to therapy in cancer patients on a particular treatment protocol, or the development of connective tissue disease symptoms in a group of third graders exposed to hazardous waste.

Ordered failures

Ordered events may result from a study that records the time to first myocardial infarction (MI), second MI, and so on. These are ordered events in the sense that the second event cannot occur before the first event. Unordered events, on the other hand, can occur in any sequence. For example, in a study of liver disease patients, a panel of seven liver function laboratory tests can become abnormal in a specific order for one patient and in a different order for another patient. The order in which the tests become abnormal (fail) is random. Two main approaches to modeling these data have gained popularity over the last few years:

• The frailty model method.

In these models the association between failure times is explicitly modeled as a random-effect term, called the <u>frailty</u> shared by all members of the cluster and assumed to follow a known statistical distribution (often the gamma distribution), with mean equal to one and unknown variance.

• Variance-corrected models.

In this approach the dependencies between failure times are not included in the models. Instead, the covariance matrix of the estimators is adjusted to account for the additional correlation. These models are easily estimated in Stata.

In this lecture we illustrate the main ideas for estimating these models using the Cox proportional hazard model.

Brief mathematical detail and definitions

Let $T_i^{(k)}$ and $U_i^{(k)}$ be the failure and censoring time of the kth failure type $(k = 1, \dots, K)$ in the ith subject $(i = 1, \dots, m)$, and let $\mathbf{Z}_i^{(k)}$ be a *p*-vector of possibly time-dependent covariates, for the ith subject with respect to the kth failure type.

"Failure type" is used here to mean both failures of different types and failures of the same type. Assume that $T_i^{(k)}$ and $U_i^{(k)}$ are independent, conditional on the covariate vector $(\mathbf{Z}_i^{(k)})$.

Define $X_i^{(k)} = \min(T_i^{(k)}, U_i^{(k)})$ and $\delta_{ij} = I(T_i^{(j)} \leq U_i^{(j)})$ where I(.) is the indicator function, and let β be a *p*-vector of unknown regression coefficients. Under the proportional hazard assumption, the hazard function of the ith subject for the kth failure type is

$$\lambda^{(k)}(t; \mathbf{Z}_i^{(k)}) = \lambda_0(t) e^{\mathbf{Z}_i^{(k)} \beta}$$

if the baseline hazard function is assumed to be equal for every failure type, or

$$\lambda^{(k)}(t; \mathbf{Z}_i^{(k)}) = \lambda_{0k}(t) e^{\mathbf{Z}_i^{(k)} \beta}$$

if the baseline hazard function is allowed to differ by failure type (Lin 1994).

Maximum likelihood estimates of for the above models are obtained from the Cox's partial likelihood function, $L(\beta)$, assuming independence of failure times. The estimator $\hat{\beta}$ has been shown to be a consistent estimator for β and is asymptotically normal as long as the marginal models are correctly specified (Lin 1994).

The resulting estimated covariance matrix obtained as the inverse of the information matrix, however, $I^{-1} = -\partial^2 \log L(\beta)/\partial\beta\partial\beta'$ does not take into account the additional correlation in the data, and therefore, it is not appropriate for testing or constructing confidence intervals for multiple failure time data.

Sandwich estimators

Lin and Wei (1989) proposed a modification to this naive estimate, appropriate when the Cox model is misspecified. The resulting robust variance-covariance matrix is estimated as

$$V = I^{-1}U'UI^{-1} = D'D$$

where U is a $n \times p$ matrix of efficient score residuals and D is the $n \times p$ vector of leverage residuals resulting from differences in the estimation of β if each observation *i* is removed from the data set (this is called *dfbeta* by many software packages). The above formula assumes that the *n* observations are independent (i.e., there is a single observation per subject – no clustering).

Sandwich estimators with clustered survival data

When observations are not independent, but can be divided into m independent groups (G_1, G_2, \dots, G_m) , then the robust covariance matrix takes the form

$$V = I^{-1}G'GI^{-1}$$

where G is a $m \times p$ matrix of the group efficient score residuals.

Implementation and examples

Implementation of all variance-adjusted models involves three steps: Setting up the data (mainly correctly specifying the time intervals), correct definition of the risk sets (by setting up $Y^{(k)}(t)$) and care in the estimation method. All of the following models can be handled:

- 1. Unordered failure events
 - (a) Unordered failure events of the same type
 - (b) Unordered failure events of different types (competing risk)
- 2. Ordered failure events
 - (a) The Andersen-Gill model
 - (b) The marginal risk set model
 - (c) The conditional risk set model (time from entry)
 - (d) The conditional risk set model (time from the previous event)

We will focus on the latter kind of models (i.e., ordered failure-time models):

1. The Andersen & Gill approach

The simplest method to implement follows the counting process approach of Andersen and Gill (1982).

The basic assumption is that all failure types are indistinguishable. This is a "conditional model" because the time interval for failure k starts at the conclusion of the interval when failure k - 1 occurred.

A major limitation of this approach is that it does not allow more than one event to occur at a given time. In addition, the A-G model assumes that all failures within the same subject are independent and models any clustering as explicit interactions included in the model. This assumption is usually untenable.

2. The WLW model

A second model, proposed by Wei, Lin, and Weissfeld (1989), is based on the idea of marginal risk sets. For this analysis, the data are treated like a set of unordered failures, so each event has its own stratum and each patient appears in all strata.

The marginal risk set at time t for event k is made up of all subjects under observation at time t regardless of whether they had experienced or not events $1, \dots, k-1$.

3. The PWP model

A third method proposed by Prentice, Williams, and Peterson (1981) is known as the <u>conditional risk set model</u>. The data are set up as for Andersen and Gill's counting processes method, except that the analysis is stratified by failure order. The assumption made is that a subject is not at risk of a second event until the first event has occurred and so on.

Thus, the conditional risk set at time t for event k is made up of all subjects under observation at time t that have had event k-1. There are two variations to this approach: Time from entry and time from previous event (the so-called "gap-time model").

In the first variation, time to each event is measured from entry time, and in the second variation, time to each event is measured from the previous event.

The above three approaches will be illustrated using the bladder cancer data presented by Wei, Lin, and Weissfeld (1989). These data were collected from a study of 85 subjects randomly assigned to either a treatment group receiving the drug thiotepa or to a group receiving a placebo control. For each patient, time for up to four tumor recurrences was recorded in months (r1-r4).

The bladder cancer data

The four models for ordered failures are illustrated by use of the bladder cancer data published in Wei, Lin & Weisfeld (1989).

id 	group	futime	number	size	r1	r2	r3	r4
1	placebo	1	1	3	0	0	0	0
2	placebo	4	2	1	0	0	0	0
3	placebo	7	1	1	0	0	0	0
4	placebo	10	5	1	0	0	0	0
5	placebo	10	4	1	6	0	0	0
6	placebo	14	1	1	0	0	0	0
7	placebo	18	1	1	0	0	0	0
8	placebo	18	1	3	5	0	0	0
9	placebo	18	1	1	12	16	0	0

[.] list in 1/9, noobs

This dataset includes data on 86 subjects with bladder cancer with follow-up between 0 and 64 months. The data for the first subject that had zero follow-up have been excluded leaving data on 85 subjects. The following are the first nine observations in the data.

The id variable identifies the patients, group is the treatment group, futime is the total follow-up time for the patient, number is the number of initial tumors, size is the initial tumor size, number is the number of initial tumors and r1 to r4 are the times to first, second, third, and fourth recurrence of tumors.

A recurrence time of zero indicates no tumor.

1. The Andersen-Gill model

To illustrate the bladder cancer data and how each of the four models creates a different data set we consider the data for subject #25 under the four models.

Under the A-G model, the data from this subject are as follows:

_							
id	group	number	size	rec	status	tstart	tstop
1		-					1
25	1	2	1	1	1	0	3
25	1	2	1	2	1	3	6
25	1	2	1	3	1	6	8
25	1	2	1	4	1	8	12
25	1	2	1	5	0	12	30
_							

This subject has experienced four recurrences at times 3, 6, 8 and 12 and was followed until time 30.

2. The Wei, Lin & Weisfeld model Under the WLW model, each patient is simultaneously at risk for all failures (thus the clock starts at time zero).

Once the fourth failure has been experienced the subject is no longer at risk for another failure (unlike the A-G model above) so the data for subject # 25 above become, under the WLW model:

+	group	number	size	rec	status	tstart	tstop
25	1	2	1	1	1	0	3
25	1	2	1	2	1	0	6
25	1	2	1	3	1	0	8
25	1	2	1	4	1	0	12
+							+

- 3. The Prentice, Williams and Peterson model In the time since entry PWP model, data are set up similarly with the A-G model but the ordering of the failure is considered by the model. In addition, time starts from entry for each interval.
 - (a) <u>The total time model</u> Under the PWP total time model the above data will be given as

id	group	number	size	rec	status	tstart	tstop
25	1	2	1	1	1	0	3
25	1	2	1	2	1	0	6
25	1	2	1	3	1	0	8
25	1	2	1	4	1	0	12
25 +	1	2	1 	4	1	0	

(b) The gap-time model

Under the gap-time model the clock starts at the end of the previous failure, so the data for the same subject are given by

_						
id	group	number	size	rec	status	gap
25	1	2	1	1	1	3
25	1	2	1	2	1	3
25	1	2	1	3	1	2
25	1	2	1	4	1	4

Implementing the Andersen-Gill model

To implement the Andersen and Gill model using the results from the bladder cancer study, the data are set up as follows: for each patient there must be one observation per event or time interval.

In general, if a subject has one event, then there will be two observations. The first observation will cover the time from entry until the time of the event, and the second observation the time from the event to the end of follow-up. The data for the nine subjects listed above are

. list if id<=10, noobs

+-	id	group	tstart	tstop	status	number	size
	1	placebo	0	1	0	1	3
I	2	placebo	0	4	0	2	0
	3	placebo	0	7	0	1	0
	4	placebo	0	10	0	5	0
	5	placebo	0	6	1	4	0
-							
I	5	placebo	6	10	0	4	0
I	6	placebo	0	14	0	1	0
I	7	placebo	0	18	0	1	0
	8	placebo	0	5	1	1	3
	8	placebo	5	18	0	1	3
-							
I	9	placebo	0	12	1	1	1
I	9	placebo	12	16	1	1	1
Ι	9	placebo	16	18	0	1	1
 +-	9	ртасеро 		18 		۱ 	⊥

In the original data, subjects 1 through 4 had no tumors recur, thus, each of these 4 patients has only one censored (status==0) observation spanning from tstart=0 to end of follow-up.

Patient 5 (id==5) had one tumor recur at 6 months and was followed until month 14. This patient has two observations in the final dataset; one from tstart to tumor recurrence (tstop==6), ending in an event (status==1), and another from tstart==6 to end of follow-up (tstop==10), ending as censored (status==0). The data are set-up as follows: . stset tstop , fail(status) exit(time .) id(id) enter(tstart) id: id failure event: status != 0 & status < .</pre> obs. time interval: (tstop[_n-1], tstop] enter on or after: time tstart exit on or before: time . 190 total obs. 0 exclusions 190 obs. remaining, representing 85 subjects 112 failures in multiple failure-per-subject data 2711 total analysis time at risk, at risk from t = 0 earliest observed entry t = 0 last observed exit t = 64

A critical component of the data set-up is to specify the start of each interval by the tstart variable.

The Andersen-Gill Cox model is fit as follows:									
. stcox group size number, nohr nolog									
failure _d: status analysis time _t: tstop enter on or after: time tstart exit on or before: time . id: id	<pre>failure _d: status analysis time _t: tstop enter on or after: time tstart exit on or before: time .</pre>								
Cox regression Breslow method for ties									
No. of subjects = 85 No. of failures = 112 Time at risk = 2711	Number of obs = 190								
Log likelihood = -460.07958	LR chi2(3) = 14.05 Prob > chi2 = 0.0028								
_t Coef. Std. Err. z	P> z [95% Conf. Interval]								
group 4070966 .2000726 -2.03 size 0400877 .0702575 -0.57 number .1606478 .0480081 3.35	0.04279923170149615 0.5681777899 .0976146 0.001 .0665536 .2547419								

The marginal risk set model (Wei, Lin, and Weissfeld)

The marginal risk model ignores the ordering of events and treats each failure as different ype of failure $i = 1, \dots, 4$. The resulting data for the first five subjects are given as follows:

- . list if id<=2, noobs
- . list if id <2

id	group	futime	number	size	rec	status
1	1	1	1	3	1	0 0
1	1	1	1	3	2	0
1	1	1	1	3	3	0
1	1	1	1	3	4	0
2	1	4	2	1	1	0
2	1	4	2	1	2	0
2	1	4	2	1	3	0
2	1	4	2	1	4	0

The data are set up as follows:

```
. stset futime, failure(status)
     failure event: status != 0 & status != .
obs. time interval: (0, time]
exit on or before: failure
                           _____
     340 total obs.
        0 exclusions
     340
          obs. remaining, representing
      112
           failures in single record/single failure data
          total analysis time at risk, at risk from t =
    8522
                                                                 0
                             earliest observed entry t =
                                                                 \mathbf{0}
                                  last observed exit t =
                                                                59
```

Conspicuous is the fact that there is no accounting for clustering of these data by subject. Each observation is considered independent of the others.

The Cox model is fitted with the sandwich estimator, clustering on each subject and stratifying on each failure type.									
. stcox group s	size number,	nohr strata	(rec) clu	uster(id)	nolog				
failure _d: status analysis time _t: futime Stratified Cox regr Breslow method for ties									
No. of subjects	; =	340		Numb	er of obs	=	340		
No. of failures	; =	105							
Time at risk	=	8522							
				Wald	chi2(3)	=	15.19		
Log pseudolikel	ihood =	402.74353		Prob	> chi2	=	0.0017		
		(S	td. Err.	adjusted	for 85 cl	uste	ers in id)		
I		Robust							
_t	Coef.	Std. Err.	Z	P> z	[95% Co	nf.	Interval]		
group	5575149	.3096526	-1.80	0.072	-1.16442	3	.0493931		
size	0663418	.0973908	-0.68	0.496	257224	2	.1245407		
number	.2082931	.0660605	3.15	0.002	.078816	9	.3377692		
	Stratified by rec								

The conditional risk set model (time from entry)

As previously mentioned, there are two variations of the conditional risk set model. The first variation in which time to each event is measured from entry is illustrated in this section.

The data are set up as for Andersen and Gill's method, however, a variable indicating the failure order is included. The analysis is then stratified by this variable. The resulting observations for the first five subjects are

. list if id<=5, noobs

+-	 id	group	tstart	tstop	status	number	size	rec
	1	1	0	1	0	1	3	1
I	2	1	0	4	0	2	1	1
	3	1	0	7	0	1	1	1
I	4	1	0	10	0	5	1	1
Ι	5	1	0	6	1	4	1	1
-								
Ι	5	1	0	10	0	4	1	2
+-								+

The resulting dataset is identical to that used to fit Andersen and Gill's model except that the **rec** variable identifies the failure risk group for each time span.

For the first 4 individuals, who have not had a recurrence, the **rec** value is one so that they are at risk for a first recurrence the whole follow-up time. The last individual, id==5, was at risk for a first recurrence for 6 months (**rec==1**) and at risk of a second recurrence (**rec==2**) from 6 months to the end of follow-up at 10 months.

The data are set up as follows:

```
. stset tstop, fail(status) exit(time .) enter(tstart)
```

```
failure event: status != 0 & status < .
obs. time interval: (0, tstop]
 enter on or after: time tstart
 exit on or before: time .
```

183 total obs.

- 0 exclusions
- 183 obs. remaining, representing 112 failures in single record/single failure data total analysis time at risk, at risk from t = 3907 earliest observed entry t = last observed exit t =59

0

0

Note that there is no clustering by subject as the time for all intervals starts at zero.

The total-time PWP model is									
. stcox group size number,	. stcox group size number, nohr nolog strata(rec)								
failure _d: status analysis time _t: tstop enter on or after: time tstart exit on or before: time .									
Stratified Cox regr Breslow method for ties									
No. of subjects = No. of failures = Time at risk = 3	183 112 3907		Numb	er of obs	= 183				
Log likelihood = -367.17	'326		LR chi2(3) = 8.75 Prob > chi2 = 0.0328						
t Coef.	Std. Err.	 Z	P> z	[95% Cont	. Interval]				
group 4897246 size 0377304 number .1102692	.2092469 .0675414 .0510491	-2.34 -0.56 2.16	0.019 0.576 0.031	8998411 1701092 .0102149	0796082 .0946484 .2103235				
Stratified by rec									

A robust (sandwich) estimate of the variance can be added: . stcox group size number, nohr nolog robust strata(rec)

failure _d: status
analysis time _t: tstop
enter on or after: time tstart
exit on or before: time .

Stratified Cox regr. -- Breslow method for ties

No. of subjects	=	183	Number of obs	=	183
No. of failures	=	112			
Time at risk	=	3907			
			Wald chi2(3)	=	9.32
Log pseudolikelihood	d =	-367.17326	Prob > chi2	=	0.0254

| Robust _t | Coef. Std. Err. z P>|z| [95% Conf. Interval]

 +							
group	4897246	.1979229	-2.47	0.013	8776464	1018029	
size	0377304	.0652989	-0.58	0.563	1657138	.090253	
number	.1102692	.0501329	2.20	0.028	.0120105	.2085278	
					Stratified by rec		

Gap time model

The gap time PWP model measures time to each event from the time of the previous event. Time is measured from zero to the gap between each failure.

group	status	number	size	rec	gap
1	0	1	3	1	1
1	0	2	1	1	4
1	0	1	1	1	7
1	0	5	1	1	10
1	0	4	1	2	6 I
1	1	4	1	1	4
	group 1 1 1 1 1 1 1 1	group status 1 0 1 0 1 0 1 0 1 0 1 0 1 0 1 1	group status number 1 0 1 1 0 2 1 0 1 1 0 5 1 0 4 1 1 4	group status number size 1 0 1 3 1 0 2 1 1 0 1 1 1 0 5 1 1 0 4 1 1 1 4 1	group status number size rec 1 0 1 3 1 1 0 2 1 1 1 0 1 1 1 1 0 5 1 1 1 0 4 1 2 1 1 4 1 1

. list if id<=5, noobs

The gap reflects the time between failures. The first four subjects had no recurrences so the gap is the total time of follow-up. Subject 5 experienced a recurrence at 6 months and then was followed up to 10 months, so the gap between the first failure and the end of follow-up is 4 months.

```
The data are set up as follows:
. stset gap status
    failure event: status != 0 & status < .
obs. time interval: (0, gap]
 exit on or before: failure
     183 total obs.
       5 obs. end on or before enter()
                                   _____
     178
         obs. remaining, representing
     112
          failures in single record/single failure data
          total analysis time at risk, at risk from t =
    2480
                                                               0
                            earliest observed entry t =
                                                               0
                                 last observed exit t =
                                                               59
The analysis proceeds as in the case of data with single
```

observations per subject, i.e., we do not include the id() option.

The correspo . stcox group s	onding gap ize number,	o-time mod nohr nolog	del is strata(re	ec)				
failure _d: status analysis time _t: gap								
Stratified Cox regr Breslow method for ties								
No. of subjects No. of failures Time at risk	= = =	178 112 2480		Numb	er of obs =	= 178		
Log likelihood = -363.16022				LR chi2(3) = 8. Prob > chi2 = 0.03				
t	Coef.	Std. Err.	z	P> z	[95% Conf.	Interval]		
group size number	2695213 .0068402 .1535334	.2076622 .0700105 .0521059	-1.30 0.10 2.95	0.194 0.922 0.003	6765318 1303777 .0514077	.1374892 .1440582 .255659		
					Stratif	ied by rec		

Clustering by id and robust variance estimation is done as follows: . stcox group size number, nohr nolog robust strata(rec) cluster(id)							
failure _d: status analysis time _t: gap							
Stratified Cox regr Breslow method for ties							
No. of subjects	3 =	178		Numb	er of obs =	178	
No. of failures	s =	112					
Time at risk	=	2480					
				Wald	chi2(3) =	11.99	
Log pseudolike	Prob	> chi2 =	0.0074				
(Std. Err. adjusted for 85 clusters in id)							
I.		Robust					
_t	Coef.	Std. Err.	Z	P> z	[95% Conf.	Interval]	
group	2695213	.2093108	-1.29	0.198	6797628	.1407203	
size	.0068402	.0625862	0.11	0.913	1158265	.129507	
number	.1535334	.0491803	3.12	0.002	.0571418	.2499249	
Stratified by rec							

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