Biostatistics (MATH11230)

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General context

- → An initial step when analysing survival or even times is to provide numerical or graphical summaries of the event times for subjects in a particular group.
- → Such summaries may be of interest in their own right or as a preliminary step before a more detailed analysis of the event times is conducted.
- \hookrightarrow Event times are convenient summarised through estimates of the survival or hazard function.

Estimating the survival function: noncensored observations

In the case of noncensoring, an obvious estimator of the survival function is the empirical estimator, given by

$$\widehat{S}(t) = \frac{\text{number of individuals with event times} > t}{\text{number of individuals in the dataset}} \\ = \frac{\#\{j: t_j > t\}}{n},$$

where t_1, \ldots, t_n are the event times and n is the number of individuals in the dataset.

- \hookrightarrow Note that $\widehat{S}(t) = 1$ for values of t below the smallest event time and $\widehat{S}(t) = 0$ for values of t above the largest event time.
- \hookrightarrow Equivalently, $\widehat{S}(t) = 1 \widehat{F}(t)$, where $\widehat{F}(t)$ is the empirical cumulative distribution function, that is,

$$\widehat{F}(t) = \frac{\#\{j: t_j \leq t\}}{n}.$$



Estimating the survival function: noncensored observations

 \hookrightarrow Let us consider the following event times (say, in months):

11 13 13 13 13 14 14 15 15 17

$$\widehat{S}(11) = \frac{\#\{j: t_j > 11\}}{11} = \frac{10}{11},$$

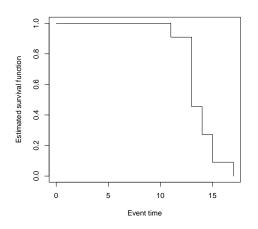
$$\widehat{S}(13) = \frac{\#\{j: t_j > 13\}}{11} = \frac{5}{11},$$

$$\widehat{S}(14) = \frac{\#\{j: t_j > 14\}}{11} = \frac{3}{11},$$

$$\widehat{S}(15) = \frac{\#\{j: t_j > 15\}}{11} = \frac{1}{11},$$

$$\widehat{S}(17) = \frac{\#\{j: t_j > 17\}}{11} = \frac{0}{11}.$$

Estimating the survival function: noncensored observations



Estimating the survival function: noncensored observations

- → Nonparametric estimators of the survival function that take into account the partial information available from the censored observations have been proposed.
- → The two most commonly used nonparametric estimators for right-censored data are the Kaplan–Meier estimator of the survival function and the Nelson–Aalen estimator of the cumulative hazard function.
- → Both estimators allow to make inferences about the distribution of the true event times based o the available information (observed event times and censoring status).

- \hookrightarrow Let us start by introducing some notation.
- \hookrightarrow Let $0 = t_0 < t_1 < t_2 < \ldots < t_J < t_{J+1} = \infty$ denote the unique noncensored event times, with d_1, \ldots, d_J the corresponding number of events at time $j, j = 1, \ldots, J$.
- \hookrightarrow Further, let n_1, \ldots, n_J be the size of the risk set at each event time, i.e., n_j is the number of individuals still event free just before t_j .

Estimating the survival function: Kaplan-Meier estimator

→ By the law of total probability, we have that

$$\Pr(T > t_j) = \Pr(T > t_j \mid T > t_{j-1}) \Pr(T > t_{j-1}) + \Pr(T > t_j \mid T \le t_{j-1}) \Pr(T \le t_{j-1}).$$

 \hookrightarrow The fact that $t_{i-1} < t_i$ implies that

$$Pr(T > t_i \mid T \le t_{i-1}) = 0,$$

as it is impossible for an individual to survive past t_j if he or she did not survive an earlier time t_{j-1} .

 \hookrightarrow Therefore,

$$S(t_j) = \Pr(T > t_j) = \Pr(T > t_j \mid T > t_{j-1}) \Pr(T > t_{j-1}).$$

Estimating the survival function: Kaplan-Meier estimator

→ But, by definition of survival function, we have that

$$Pr(T > t_{j-1}) = S(t_{j-1}),$$

and thus

$$S(t_j) = \Pr(T > t_j) = \Pr(T > t_j \mid T > t_{j-1})S(t_{j-1}).$$

 \hookrightarrow It also holds that

$$S(t_{j-1}) = \Pr(T > t_{j-1} \mid T > t_{j-2})S(t_{j-2}),$$

and that

$$S(t_{j-2}) = \Pr(T > t_{j-2} \mid T > t_{j-3})S(t_{j-3}),$$

and that

 \hookrightarrow This implies that

$$S(t_j) = \Pr(T > t_j \mid T > t_{j-1}) \times \Pr(T > t_{j-1} \mid T > t_{j-2}) \times \ldots \times \Pr(T > t_2 \mid T > t_1) S(t_1). \tag{1}$$



Estimating the survival function: Kaplan-Meier estimator

- → We must now simply plug in estimates of each of the terms on the right-hand side of Equation (1).

$$\begin{split} \widehat{\Pr}(T > t_j \mid T > t_{j-1}) &= 1 - \widehat{\Pr}(T \leq t_j \mid T > t_{j-1}) \\ &= 1 - \frac{\# \text{ number of events in } (t_{j-1}, t_j]}{\# \text{ number of individuals at risk at time } t_j} \\ &= 1 - \frac{d_j}{n_j} \\ &= \frac{n_j - d_j}{n_i}. \end{split}$$

→ This leads to the Kaplan–Meier estimator of the survival curve

$$\widehat{\mathcal{S}}^{\mathsf{KM}}(t) = \prod_{j:t_j \leq t} \frac{n_j - d_j}{n_j},$$

with $\widehat{S}^{KM}(t) = 1$ for $t < t_1$.



- → This estimator was originally proposed by Kaplan and Meier in 1958, hence the name Kaplan–Meier estimator.
- → This estimator is also often referred to as the product limit estimator.
- This approach is undeniably the most used one to estimate and summarise survival curves
- → This method is so widespread that the original article is the most highly cited article in the history of statistics.

Estimating the survival function: Kaplan-Meier estimator

 \hookrightarrow To demonstrate the computation of $\widehat{S}^{\text{KM}}(t)$ we consider the following hypothetical dataset

Patient	1	2	3	4	5	6	7
Time (in months)	1	3	3	6	8	9	10
Censoring status	1	1	1	0	0	1	0

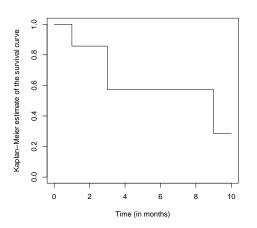
- → Here a censoring status equal to 1 means that the corresponding event time is not censored and 0 that it is censored.
- → Let us construct the Kaplan–Meier estimate of the survival curve:

Estimating the survival function: Kaplan-Meier estimator

→ We thus have that

$$\widehat{S}^{KM}(t) = \begin{cases} 1, & t < 1 \\ \frac{6}{7}, & 1 \le t < 3 \\ \frac{4}{7}, & 3 \le t < 9 \\ \frac{2}{7}, & 9 \le t < 10 \end{cases}$$

- \hookrightarrow Note that the estimate of $\widehat{S}^{\text{KM}}(t)$ is undefined for t > 10 because the largest observation is a censored event time and $\widehat{S}^{\text{KM}}(t)$ cannot be estimated consistently beyond this time.
- \hookrightarrow On the other hand, if the largest event time is an uncensored observation, then $n_J = d_J$, and so $\widehat{S}^{\text{KM}}(t)$ is zero for $t \geq t_J$.



- → As we could notice, the Kaplan–Meier estimate of the survival function is a step function, in which the estimated survival probabilities are constant between adjacent event times and decrease at each event time.
- → If there are no censored observations in the dataset, the Kaplan–Meier estimator reduces
 to the empirical estimator of the survival function that we have seen at the beginning of the
 lecture.

- \hookrightarrow A key summary statistic of the survival function is the **median survival time**.
- \hookrightarrow The median survival time is defined as the smallest time t such that $S(t) \le 1/2$.
- \hookrightarrow This can be estimated from the Kaplan–Meier plot by finding where the curve intersects the horizontal line $\widehat{S}^{\text{KM}}(t)=1/2$.

Estimating the survival function: Kaplan-Meier estimator

The variance of the Kaplan–Meier estimator can be approximated by the so-called Greenwood formula (see, for example, Collett, 2014, chapter 2)

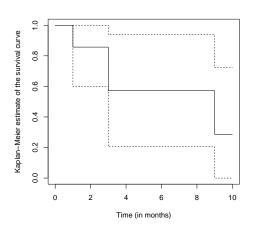
$$\widehat{\text{var}}(\widehat{S}^{\text{KM}}(t)) = \left[\widehat{S}^{\text{KM}}(t)\right]^2 \sum_{j: t_j \leq t} \frac{d_j}{n_j(n_j - d_j)}.$$

→ For large samples, the following result holds

$$\frac{\widehat{S}^{\text{KM}}(t) - S(t)}{\sqrt{\widehat{\text{var}}(\widehat{S}^{\text{KM}}(t))}} \sim \mathsf{N}(0,1).$$

 \hookrightarrow This result can be used to derive a confidence interval for S(t)

$$\left(\widehat{S}^{\text{KM}}(t) - z_{\alpha/2} \sqrt{\widehat{\text{var}}(\widehat{S}^{\text{KM}}(t))}, \widehat{S}^{\text{KM}}(t) + z_{\alpha/2} \sqrt{\widehat{\text{var}}(\widehat{S}^{\text{KM}}(t))}\right).$$



- \hookrightarrow This CI is not accurate (may produce limits beyond the range of zero or one) when $\widehat{S}^{\text{KM}}(t)$ is close to 0 or 1, so often CIs are first calculated for a transformation, for example, $\log(-\log S(t))$.
- → For more details about this, I refer the interested reader to Collett, 2014, chapter 2.
- \hookrightarrow The package survival implements both approaches. See more in the Supplementary Materials file.

Estimating the survival function: Nelson-Aalen estimator

- → An alternative estimator of the survival function is based on the so called Nelson–Aalen estimator of the cumulative hazard function, proposed independently by Nelson and Aalen in the 70s.
- → This estimator is given by

$$\widehat{H}(t) = \sum_{j:t_j \leq t} \frac{d_j}{n_j}.$$

- \hookrightarrow The estimated cumulative hazard up to time t is just the sum of the estimated hazards at all event times up to t.

Estimating the survival function: Nelson-Aalen estimator

→ From this estimator, one can obtain the Nelson–Aalen estimate of the survival function

$$\begin{split} \widehat{S}^{\text{NA}}(t) &= \exp\{-\widehat{H}(t)\} \\ &= \exp\left\{-\sum_{j:t_j \le t} \frac{d_j}{n_j}\right\} \\ &= \prod_{j:t_j \le t} \exp\left\{-\frac{d_j}{n_j}\right\}. \end{split}$$

Estimating the survival function: Nelson-Aalen estimator

- Interestingly, the Kaplan-Meier estimator of the survival function can actually be regarded as a first-order Taylor expansion approximation, around zero, of the Nelson-Aalen estimator.
- → Recall that based on the first order Taylor expansion around zero, we can write

$$f(x) \approx f(0) + (x - 0)f'(0).$$

- \hookrightarrow Letting $f(x) = e^{-x}$, we have that $e^{-x} \approx 1 x$.
- \hookrightarrow Thus,

$$\widehat{S}^{\mathsf{NA}}(t) pprox \prod_{j:t_j \leq t} \left(1 - \frac{d_j}{n_j}\right) = \widehat{S}^{\mathsf{KM}}(t).$$

Estimating the survival function: Nelson-Aalen estimator

→ For the hypothetical dataset in slide 12, we can also compute the Nelson–Aalen estimate of the survival curve.

