Biostatistics (MATH11230)

Nonparametric estimators of the survival function

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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

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

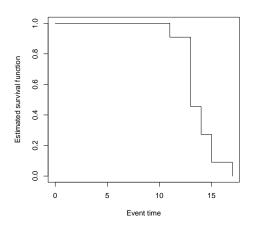
$$\widehat{S}(13) = \frac{\#\{j: t_j > 13\}}{11} = \frac{5}{11},$$

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

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

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

Estimating the survival function: noncensored observations



Estimating the survival function: noncensored observations

- → The empirical estimator of the survival function cannot be used when there are censored observations as it actually disregards the information provided by censored 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 on the available information (observed event times and censoring status).

- \hookrightarrow Let $0 = t_0 < t_1 < t_2 < \ldots < t_J < t_{J+1} = \infty$ denote the unique uncensored 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

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

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

$$\widehat{S}^{\text{KM}}(t) = \prod_{j:t_j \le t} \frac{n_j - d_j}{n_j}, \quad \text{ for } t_j \le t < t_{j+1},$$

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

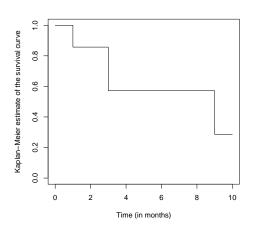
Patient	1	2	3	4	5	6	7
Event time (in months)	1	3	3	6	8	9	10
Censoring status	1	1	1	0	0	1	0

- → 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

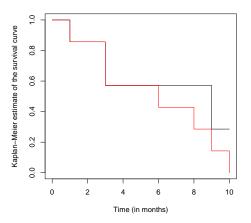
$$\widehat{S}^{KM}(t) = \begin{cases} 1, & t < 1 \\ \frac{6}{7}, & 1 \le t < 3 \\ \frac{4}{7}, & 3 \le t < 9 \\ \frac{2}{7}, & 9 \le t < 10 \end{cases}$$

- \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 $t \geq t_J$.



Estimating the survival function: Kaplan-Meier estimator

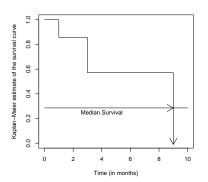
→ Below in red is the empirical estimate of the survival function pretending that the censored event times were the true/observed ones



- → 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 $\widehat{S}^{\text{KM}}(t)=1/2$.

- → For the toy example in slides 12/13, the median survival time is 9, as this is the smallest time for which the survival curve is (equal) or below 0.5.
- → We can get this easily in R as part of the output of the function that fits the Kaplan-Meier estimator (see more in the Supplementary Materials).



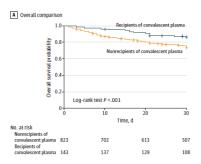
Kaplan-Meier estimator in recent scientific publications

Research

JAMA Oncology | Original Investigation

Association of Convalescent Plasma Therapy With Survival in Patients With Hematologic Cancers and COVID-19

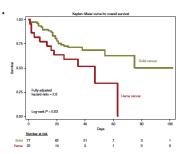
Michard A. Thompson, MD, PhD, Jeffey P. Handsonn, MD, PhD, Paniel K. Shah, MD, MSH-K. Samuel M. Rabinstin, MD, Michael J. Joyner, MD. Torois K. Obuselin, MD, Duriel B. Hora, MD, Parem D. Estabeth A. Grifforth, MD, Anthony P. Golda, MD, Cluir Ahreng, MD, Volenin S. Koshilin, MD, Esperance B. Papodopoulous, MD, Estabeth M. Polioteth, MD, MPH, Christopher T. Su, MD, MMH—Estabeth M. Wall Brundfold, MD, Zhouser Ke, MD, MS; Peter Paral N, MD, Sarigy Michael, SP, PD, Swathon W. Swenfeld, PHD, Drony P. Shah, MD, PHD, Lenemy L. Warner, MD, MS; for the COVID-19 and Carner Concordium.



Kaplan-Meier estimator in recent scientific publications



CD8+ T cells contribute to survival in patients with COVID-19 and hematologic cancer

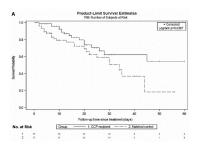


Kaplan-Meier estimator in recent scientific publications

RESEARCH ARTICLE

Early but not late convalescent plasma is associated with better survival in moderate-to-severe COVID-19

Neima Briggs©^{1‡}, Michael V. Gormally^{1‡}, Fangyong Li², Sabrina L. Browning©², Miriam M. Treggiario⁴, Alyssa Morrison⁵, Maudry Laurent-Rolle⁶, Yanhong Deng², Jeanne E. Hendrickson¹², Christopher A. Tormey^{*}, Mahalia S. Desruisseaux⁶ *



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)

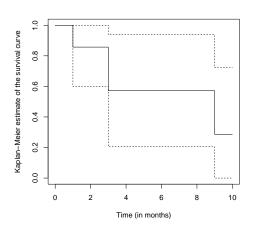
$$\widehat{\text{var}}(\widehat{S}^{\text{KM}}(t)) = \left[\widehat{S}^{\text{KM}}(t)\right]^2 \sum_{j: t_j \leq t} \frac{d_j}{n_j(n_j - d_j)}.$$

→ 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.

$$\widehat{S}^{NA}(t) = \exp\{-\widehat{H}(t)\}\$$

$$= \exp\left\{-\sum_{j:t_j \le t} \frac{d_j}{n_j}\right\}\$$

$$= \prod_{j:t_j \le t} \exp\left\{-\frac{d_j}{n_j}\right\}.$$

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.

