Biostatistics (MATH11230)Parametric estimators of the survival function

Vanda Inácio

University of Edinburgh



Semester 1, 2022/2023

General context

- → In a parametric setup we instead choose a (parametric) distribution that we believe to appropriately describe the distribution of the survival times.
- → As we have discussed last week, popular approaches are the Weibull, lognormal, and gamma distributions. But there are many more (e.g., Rayleigh, Pareto, etc).
- The exponential distribution, although analytically tractable, has the caveat of leading to a constant hazard function with respect to time and so it may be too simplistic in most situations.

General context

- \hookrightarrow Suppose we have a random sample of *n* individuals from a specific population whose true survival/event times are T_1, \ldots, T_n .
- However, due to right censoring (study period ends, dropouts/losses to follow up, competing risks, etc) we do not always have the opportunity of observing those survival times.
- \hookrightarrow Let us denote by C the censoring process and by C_1, \ldots, C_n the (potential) censoring times for each individual.
- → If a subject is not censored we have observed their event time (in this case, we may not observe the censoring time for this individual).
- → On the other hand, if the subject is censored, we observe their censoring time (true event time is larger than censoring time).

Likelihood construction

 \hookrightarrow Let

$$Y_i = \min\{T_i, C_i\}, \quad \text{and} \quad \Delta_i = I(T_i \leq C_i).$$

- \hookrightarrow The observed data are then composed of couples (y_i, δ_i) , for $i = 1, \dots, n$.
- \hookrightarrow If y_i is not censored, observation i contributes with a factor to the likelihood equal to the density function of T evaluated at y_i , $f_T(y_i)$.
- \hookrightarrow If y_i instead represents a censored time, it is only known that t_i (true event time) exceeds y_i (observed event time) and the contribution to the likelihood is the probability that $T_i > c_i$ (equal to $\Pr(T_i > y_i)$) and this probability is $S_T(y_i)$, where S is the survival function of T.

Likelihood construction

 $\,\hookrightarrow\,$ Assuming independent censoring and independent observations, we have that the likelihood is given by

$$L(\theta) = \prod_{i=1}^{n} \left\{ f(y_i; \theta)^{\delta_i} S(y_i; \theta)^{1-\delta_i} \right\}$$

$$= \prod_{i=1}^{n} \left\{ [h(y_i; \theta) S(y_i; \theta)]^{\delta_i} S(y_i; \theta)^{1-\delta_i} \right\}$$

$$= \prod_{i=1}^{n} \left\{ h(y_i; \theta)^{\delta_i} S(y_i; \theta) \right\}.$$

 \hookrightarrow Based on this likelihood, we can then determine the maximum likelihood estimate of θ (which may be a scalar or a vector) and having this we have all we need to estimate the survival, hazard, and cumulative hazard functions.

Example: exponential distribution

- \hookrightarrow Let us assume that the true survival times T_1, \ldots, T_n follow an exponential distribution.

$$f(t;\theta) = \theta e^{-\theta t}, \quad S(t;\theta) = e^{-\theta t}, \quad h(t;\theta) = \theta, \quad \theta > 0.$$

 \hookrightarrow According to the result in the previous slide we have that

$$L(\theta) = \prod_{i=1}^{n} h(y_i; \theta)^{\delta_i} S(y_i; \theta)$$
$$= \prod_{i=1}^{n} \theta^{\delta_i} e^{-\theta y_i}$$
$$= \theta^{\sum_{i=1}^{n} \delta_i} e^{-\theta \sum_{i=1}^{n} y_i}$$

Example: exponential distribution

$$\log L(\theta) = \log \theta \sum_{i=1}^{n} \delta_{i} - \theta \sum_{i=1}^{n} y_{i},$$

and the derivative with respect to θ is

$$\frac{\mathsf{d}}{\mathsf{d}\theta}\log L(\theta) = \frac{1}{\theta}\sum_{i=1}^n \delta_i - \sum_{i=1}^n y_i.$$

 \hookrightarrow Setting the derivative to zero and solving for θ :

$$\frac{d}{d\theta} \log L(\theta) = 0 \Rightarrow \frac{1}{\theta} \sum_{i=1}^{n} \delta_{i} - \sum_{i=1}^{n} y_{i} = 0$$
$$\Rightarrow \hat{\theta} = \frac{\sum_{i=1}^{n} \delta_{i}}{\sum_{i=1}^{n} y_{i}}$$



Example: exponential distribution

$$\frac{\mathrm{d}^2}{\mathrm{d}\theta^2}\log L(\theta) = -\frac{1}{\theta^2}\sum_{i=1}^n \delta_i.$$

- \hookrightarrow Evaluating this second derivative at $\widehat{\theta}$ we conclude that it is negative and therefore $\widehat{\theta}$ is the maximum likelihood estimate.
- \hookrightarrow Note that in the case where all observations are uncensored, $\sum_{i=1}^{n} \delta_i = n$, and we recover the 'usual' maximum likelihood estimate, $\widehat{\theta} = n / \sum_{i=1}^{n} t_i$.

Example: Exponential distribution

 \hookrightarrow As an example let us consider the data we have used yesterday to manually compute the Kaplan–Meier estimate:

- \hookrightarrow We have four uncensored observations and so we know that $\sum_{i=1}^{7} \delta_i = 4$.
- \hookrightarrow We further have that $\sum_{i=1}^{7} y_i = 1 + 3 + 3 + 6 + 8 + 9 + 10 = 40$.
- \hookrightarrow Therefore, $\widehat{\theta} = 4/40 = 0.1$.

Example: Weibull distribution

- → A more flexible alternative to the exponential distribution is the Weibull distribution.
- → The defining feature of the Weibull distribution is that the corresponding hazard function is monotonic. It can be monotonically increasing, monotonically decreasing or constant with respect to time.
- → It is fair to mention that the assumption of ever increasing or ever decreasing hazards may be unrealistic in certain situations as well.
- Nevertheless, the Weibull distribution is ubiquitous in survival analysis and we should consider it.

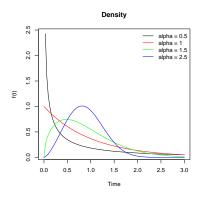
Example: Weibull distribution

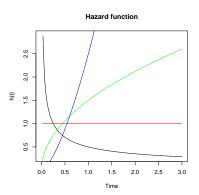
$$f(t;\lambda,\alpha) = \lambda \alpha t^{\alpha-1} e^{-\lambda t^{\alpha}}, \quad S(t;\lambda,\alpha) = e^{-\lambda t^{\alpha}}, \quad h(t;\lambda,\alpha) = \lambda \alpha t^{\alpha-1}, \quad \lambda > 0, \quad \alpha > 0.$$

- \hookrightarrow Here λ is the scale parameter and α is the shape parameter. For $\alpha<$ 1, the hazard is decreasing, $\alpha=$ 1 the hazard is constant (see below), and $\alpha>$ 1 the hazard is increasing. The parameter scale λ stretches or squeezes the distribution.
- \hookrightarrow Note that the exponential distribution is a particular case of the Weibull distribution when the shape parameter α is equal to 1.

Example: Weibull distribution

 \hookrightarrow In the plots below $\lambda = 1$.





Example: Weibull distribution

 \hookrightarrow The likelihood function of $\theta = (\lambda, \alpha)$ is given by

$$L(\theta) = \prod_{i=1}^{n} \left\{ h(y_i; \theta)^{\delta_i} S(y_i; \theta) \right\}$$
$$= \prod_{i=1}^{n} \left\{ [\lambda \alpha y_i^{\alpha - 1}]^{\delta_i} e^{-\lambda y_i^{\alpha}} \right\}.$$

- \hookrightarrow However, there is no closed form for the MLEs of $\theta = (\lambda, \alpha)$.
- → We can use, for instance, optim to maximise the above (log) likelihood function. We can also use, and it is indeed what we will do, the function survreg from the package survival.
- $\hookrightarrow \ \ \text{Please see the Supplementary Materials file for more details about the fitting of the model}.$

- → Before fitting a parametric model, we should check that it fits appropriately to the data at hand.
- → For example, we could plot the Kaplan-Meier estimate of the survival function and the estimated survival model under an exponential or Weibull model.
- → Personally, I prefer the Kaplan-Meier estimator which makes no assumptions regarding the shape of the distribution of the event times.
- \hookrightarrow In most medical applications, the default is to go with the Kaplan-Meier estimator.