

Basic Concepts

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Biostatistics

Biostatistics (a combination of biology and statistics; sometimes referred to as **biometry** or **biometrics**) is the application of [statistics](#) to a wide range of topics in [biology](#). The science of biostatistics encompasses the design of biological [experiments](#), especially in [medicine](#), [pharmacy](#), [agriculture](#) and [fishery](#); the collection, summarization, and analysis of data from those experiments; and the interpretation of, and inference from, the results.

Survival time

- Time from a **starting point** until an **event of interest** occurs.
For example,
 - Time from birth to death
 - Duration of marriage
 - Time from treatment to relapse of a specific disease
- The starting point is sometimes harder to define than the endpoint
 - When is the actual beginning of a disease? Using time of diagnosis instead?
- Endpoint not necessarily something negative, such as illness or death
 - Can be, e.g., recovery from a disease

Survival time, cont.

- Also known as **time-to-event** data
- Rarely useful to calculate mean survival time
 - Requires that endpoint actually occurs and is observed for all subjects
 - Survival data rarely normally distributed
- Survival data analyzed by using special methods

Notation

- We denote by a **capital T** the random variable for a person's survival time.
- Since T denotes time, its possible values include all nonnegative numbers; that is, **T** can be any number equal to or greater than zero.

$T = \text{survival time } (T \geq 0)$

random variable



Notation

- We denote by a **small letter t** any specific value of interest for the random variable capital **T** .
- For example, if we are interested in evaluating whether a person survives for more than 5 years after undergoing cancer therapy, small **t** equals 5; we then ask whether capital **T** exceeds 5.

Survives > 5 years?
 $T > t = 5$

Notation

- We denote the **small letter d** to define a **(0,1)** random variable indicating either failure or censorship.
- That is, **d=1** for **failure** if the event occurs during the study period, or **d=0** if the survival time is **censored** by the end of the study period.

$d = (0, 1)$ random variable

$$= \begin{cases} 1 & \text{if failure} \\ 0 & \text{censored} \end{cases}$$

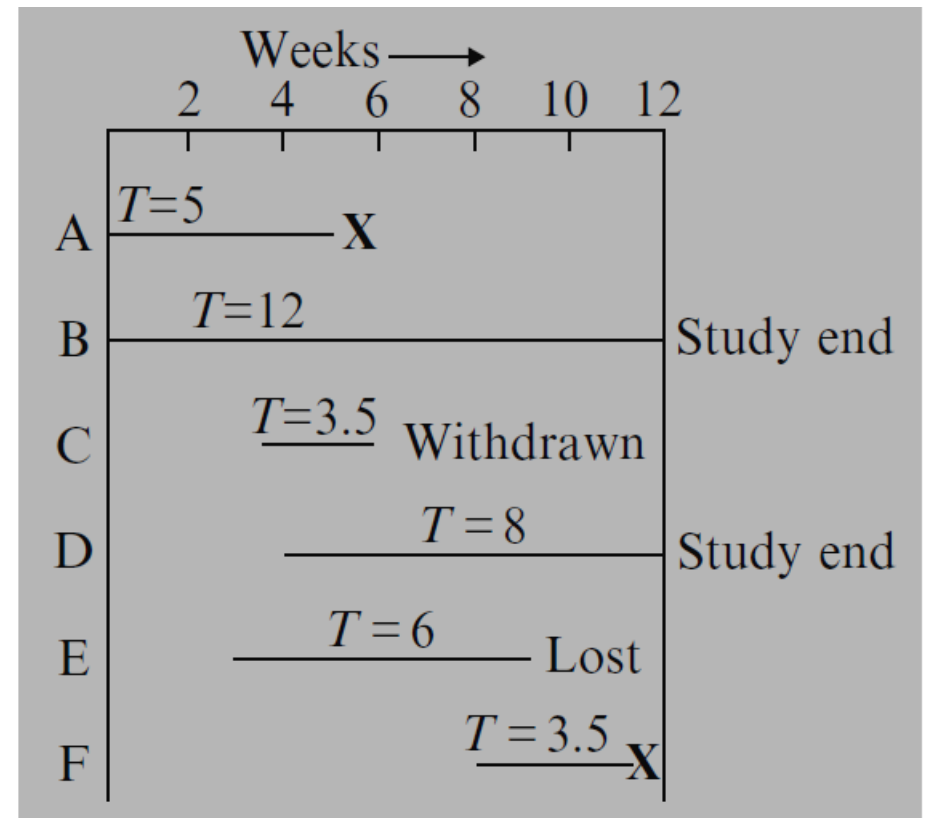
Censoring

- Exact time of the endpoint is not known
 - Event of interest not observed for all subjects at the end of the study
- Survival time is only partly known (e.g., “at least as large as”)
- Often due to data collected during a limited period of time
- Three main types
 - Right-censoring
 - Left-censoring
 - Interval-censoring

Reasons for Censoring

- A person does not experience the event before **the study ends**;
- A person is **lost to follow-up** during the study period;
- A person **withdraws from the study** because of death (if death is not the event of interest) or some other reason (e.g., adverse drug reaction or other competing risk)

Example of Censoring

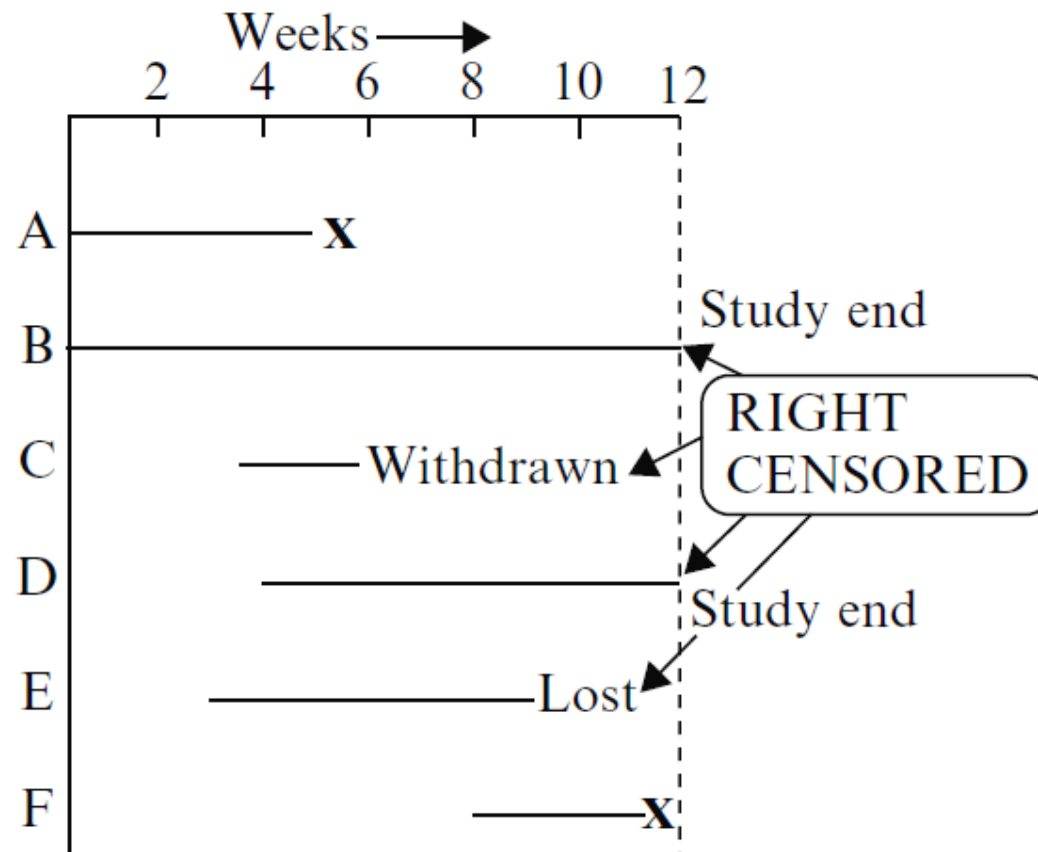


In **summary**, of the six persons observed, two get the event (persons A and F) and four are censored (B, C, D, and E).

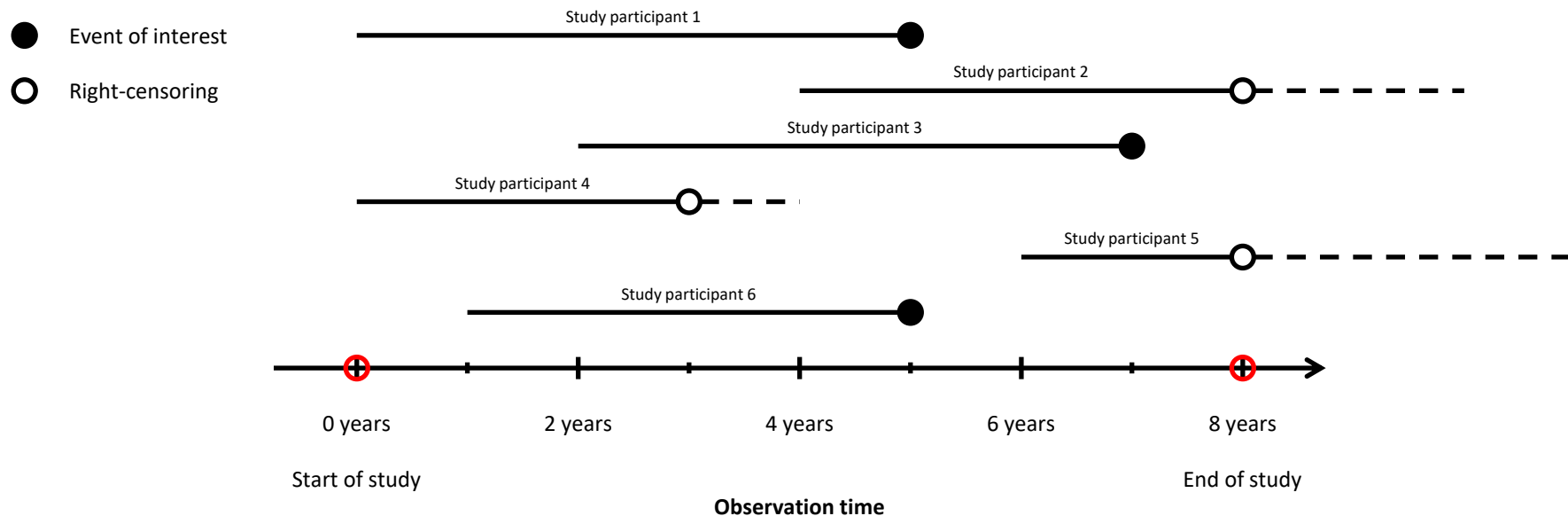
X \implies Event occurs

Right Censoring

Right-censored: true survival time is equal to or greater than observed survival time



Right Censoring, cont.



Right Censoring, cont.

- Right-censoring due to **study termination** or **loss to follow-up**
- Several possible reasons for being lost to follow-up
 - Not responding to questionnaires or attending scheduled hospital visits
 - Study withdrawal
 - Moving or emigration
 - Death (by a cause other than that being studied)
- A common phenomenon
 - Routinely handled in survival analysis

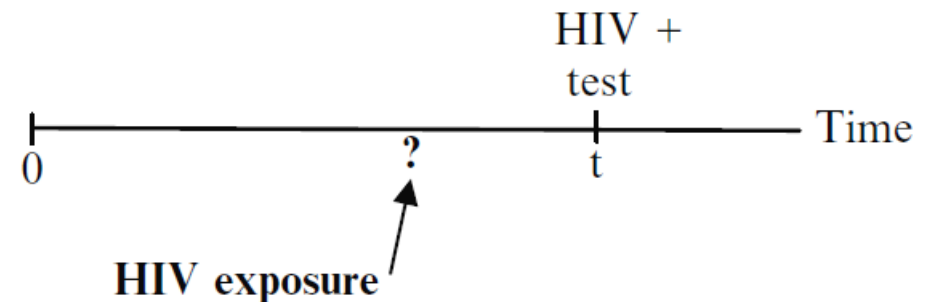
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Left Censoring

Left-censored: true survival time is less than or equal to the observed survival time

- If a person is left-censored at time t , we know they had an event between time 0 and t , but we do not know the exact time of event.

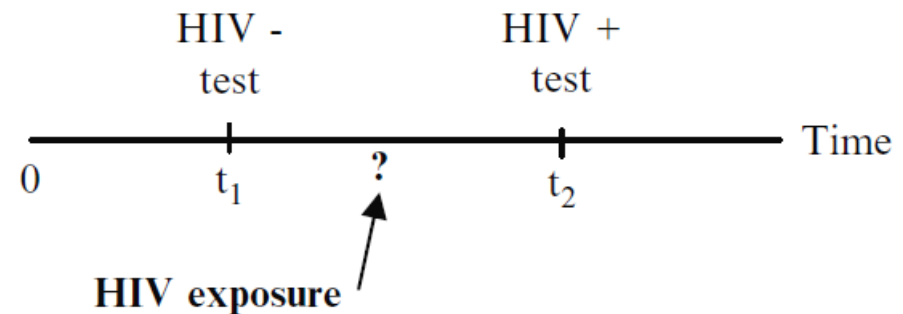


Event occurs between 0 and t
but
do not know the exact time.

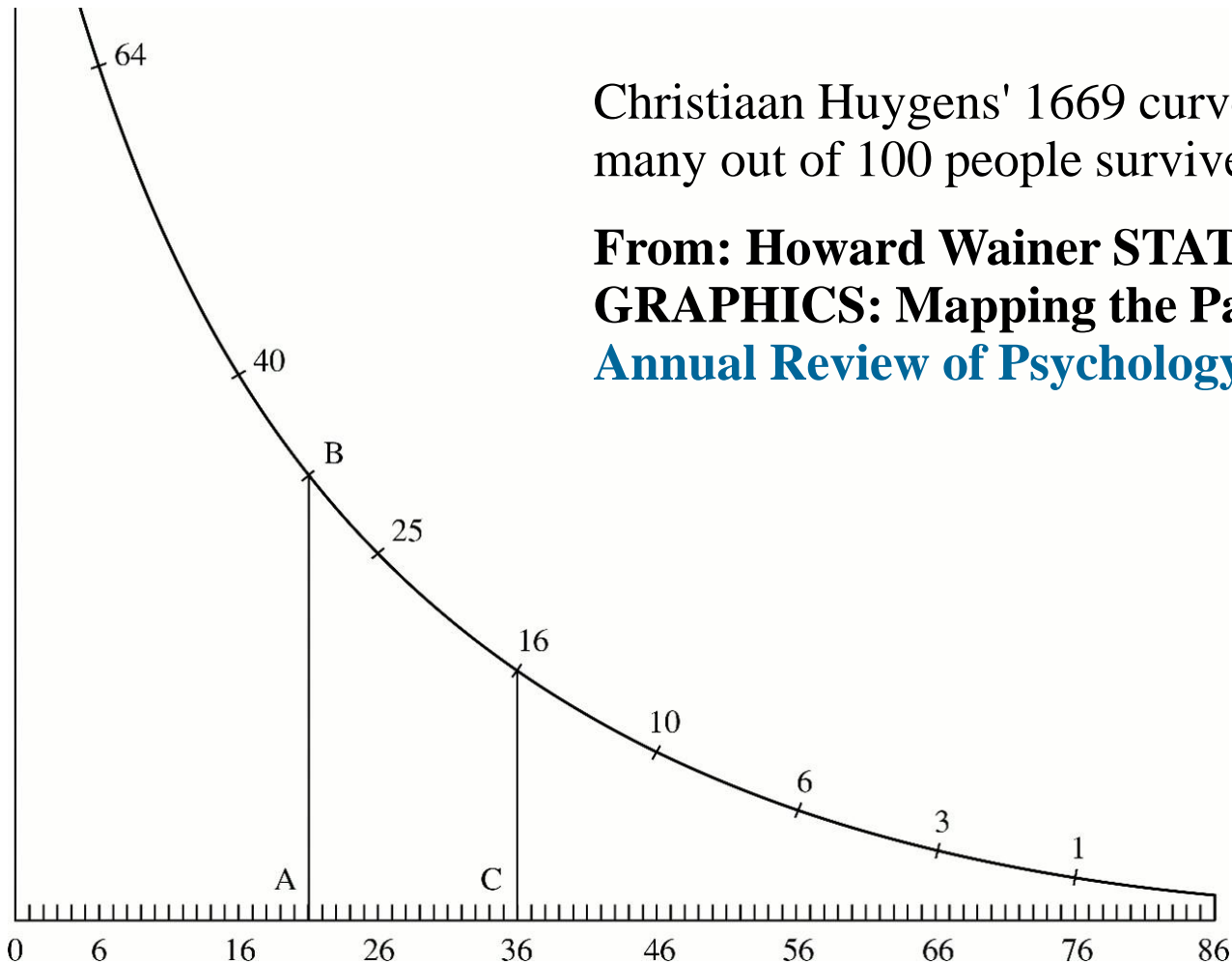
Interval Censoring

Interval-censored: true survival time is within a known time interval

A subject may have had two HIV tests, where he/she was HIV negative at the time (say, t_1) of the first test and HIV positive at the time (t_2) of the second test. In such a case, the subject's true survival time occurred after time t_1 and before time t_2 , i.e., the subject is interval censored in the time interval (t_1, t_2) .



Early example of survival analysis, 1669



Christiaan Huygens' 1669 curve showing how many out of 100 people survive until 86 years.

From: Howard Wainer STATISTICAL GRAPHICS: Mapping the Pathways of Science.
Annual Review of Psychology. Vol. 52: 305-335

What is survival analysis?

- Statistical methods for analyzing longitudinal data on the occurrence of events.
- Events may include death, injury, beginning of illness, recovery from illness (binary variables) or transition above or below the clinical threshold of a meaningful continuous variable.
- Accommodates data from randomized clinical trial or cohort study design.

Objectives of survival analysis

- **Estimate time-to-event for a group of individuals**, such as time until second heart-attack for a group of MI patients.
- **To compare time-to-event between two or more groups**, such as treated vs. placebo MI patients in a randomized controlled trial.
- **To assess the relationship of co-variables to time-to-event**, such as: does weight, insulin resistance, or cholesterol influence survival time of MI patients?

Note: expected time-to-event = $1/\text{incidence rate}$

Why use survival analysis?

- **Logistic regression** can predict the presence or absence of events but not time until events and it can not handle time dependent covariates.
- **Linear regression** can not handle censoring well or time dependent covariates or the fact that time can only be positive.

Survival Analysis Steps

- Get some data and make sure it is valid.
- Estimate the survival/hazard functions.
- Compare the functions between groups.
- Assess the impact of predictors on survival rates.

Survival Function or Curve

- Let T denote the survival time
- $S(t) = P(\text{surviving longer than time } t) = P(T > t)$
- *The function $S(t)$ is also known as the cumulative survival function. $0 \leq S(t) \leq 1$*

$$\hat{S}(t) = \frac{\text{number of patients surviving longer than } t}{\text{total number of patients}}$$

$$\therefore S(t) = \int_t^{\infty} f(x) dx ; \quad \text{where } f(x) = \text{pdf}$$

Survival function

- The goal of survival analysis is to estimate and compare survival experiences of different groups.
- Survival experience is described by the cumulative survival function:

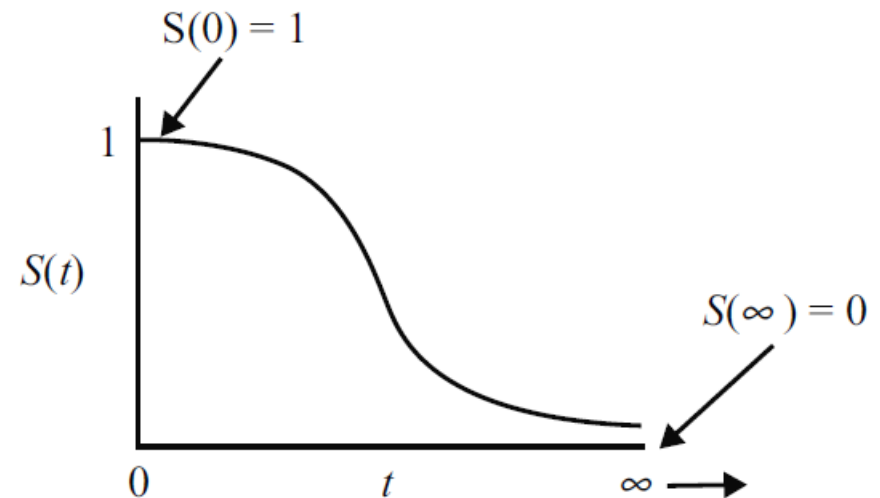
$$S(t) = 1 - P(T \leq t) = 1 - F(t)$$

$F(t)$ is the CDF of $f(t)$, and is “more interesting” than $f(t)$.

- **Example:** If $t=100$ years, $S(t=100)$ = probability of surviving beyond 100 years.

Characteristics survivor functions


- They are **nonincreasing**; that is, they head downward as t increases;
- At time $t=0$, $S(0)=1$; that is, at the start of the study, since no one has gotten the event yet, the probability of surviving past time 0 is one;
- At time $t=\infty$, $S(\infty)=0$; that is, theoretically, if the study period increased without limit, eventually nobody would survive, so the survivor curve must eventually fall to zero.



Hazard Function

- The hazard function $h(t)$ of survival time T gives the *conditional failure rate*
- The hazard function is also known as the *instantaneous failure rate, force of mortality, and age-specific failure rate*
- *The hazard function gives the risk of failure per unit of time during the aging process*

Hazard Function

Given 

$$h(t) = \lim_{\Delta t \rightarrow 0} \frac{P(t \leq T < t + \Delta t \mid T \geq t)}{\Delta t}$$

Conditional probabilities: $P(A|B)$

$$\begin{aligned} &P(t \leq T < t + \Delta t \mid T \geq t) \\ &= P(\text{individual fails in the interval} \\ &\quad [t, t + \Delta t] \mid \text{survival up to time } t) \end{aligned}$$

$$h(t) = \frac{f(t)}{1 - F(t)} = \frac{f(t)}{S(t)}$$

Cumulative Hazard Function

The cumulative hazard function is defined as

$$H(t) = \int_0^t h(x)dx$$

Relationship of $S(t)$ and $h(t)$

$$f(t) = S(t) \times h(t)$$

Relations

$$\text{Hazard from density and survival: } h(t) = \frac{f(t)}{S(t)}$$

$$\text{Survival from density: } S(t) = \int_t^{\infty} f(u) du$$

$$\text{Density from survival: } f(t) = -\frac{dS(t)}{dt} e^{(-\int_0^t h(u) du)}$$

$$\text{Density from hazard: } f(t) = h(t) e^{(-\int_0^t h(u) du)}$$

$$\text{Hazard from survival: } h(t) = -\frac{d}{dt} \ln S(t)$$

Example

Suppose, $f(t) = e^{-t}$; $t \geq 0$

Find $S(t)$, and $h(t)$.

We know that, $S(t) = 1 - F(t)$

$$F(t) = \int_0^t f(x) dx = \int_0^t e^{-x} dx = 1 - e^{-t}$$

$$S(t) = 1 - (1 - e^{-t}) = e^{-t}$$

$$f(t) = h(t) \times S(t)$$

$$h(t) = \frac{f(t)}{S(t)} = \frac{e^{-t}}{e^{-t}} = 1$$

Practice

Suppose, $f(t) = \lambda e^{-\lambda t}$; $t \geq 0$

Find $S(t)$, and $h(t)$.

Thanks for Your Attention