

Lecture 4

Monday, September 25, 2023 10:01



BIOS522_Sli
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BIOS 522: Survival Analysis Methods

Lecture 4:

The hazard and cumulative hazard functions

Previously

- *Introduced the survival function*
- *Defined the Kaplan-Meier estimator*
- *Calculated the log-rank test for comparing survival curves*
- *Used R to implement these procedures*

Survival random variable

- Non-negative random variable T
- For a given population, we may want to summarize:
 - The mean of T
 - The median of T
 - The variance of T
 - The density function (pdf) for T
 - The cumulative distribution function (CDF) for T

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Survival random variable

- For time-to-event random variables, we are also interested in summarizing other key quantities
 - The survival function $S(t)$
 - Probability of surviving beyond time t
 - The **hazard function** $h(t)$
 - **Instantaneous rate of failure** among those **still at risk** at time t
 - The **cumulative hazard function** $H(t)$
 - The accumulated hazard from time 0 to time t

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Hazard function

- Among those still at risk at time t , what is the instantaneous rate of failure at time t ?

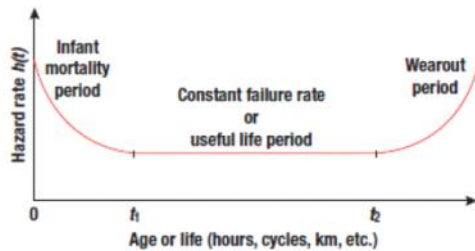
$$\begin{aligned} h(t) &= \lim_{\Delta \downarrow 0} \frac{1}{\Delta} \Pr(t \leq T < t + \Delta | T \geq t) \\ &= \frac{f(t)}{S(t)} = \frac{\text{density function}}{\text{survival function}} \end{aligned}$$

conditional on
not having failed
before time t

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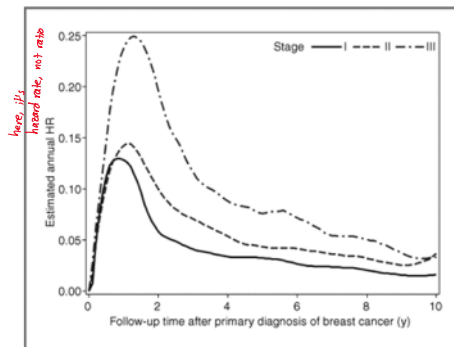
Hazard is a speedometer for risk

- Can use the hazard function to identify periods of highest risk



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Example: Breast cancer recurrence



- Smoothed hazard functions by tumor stage for first recurrence among women after primary breast cancer treatment
- Early period of elevated risk of recurrence
- At all times, highest risk is for Stage III cancers

Source: Cheng et al. (2012) Cancer, Epi, Biomarkers & Prevention [10.1158/1055-9965.EPI-11-1089](https://doi.org/10.1158/1055-9965.EPI-11-1089)

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Example: Seasonal mortality in wildlife

DOI: 10.1111/2041-210X.13305

APPLICATION

Methods in Ecology and Evolution

For everything there is a season: Analysing periodic mortality patterns with the `cyclomort` R package

Eliezer Gurarie¹ | Peter R. Thompson^{1,2} | Allicia P. Kelly³ | Nicholas C. Larter⁴ | William F. Fagan¹ | Kyle Joly⁵

- For many species, mortality risk follows a seasonal pattern
- For example, during certain times of the year, resources may be scarce, or susceptibility to predators or disease is high
- The authors create an R package to model seasonal mortality patterns

Source: Gurarie et al. (2020) Methods Ecol Evol, doi: 10.1111/2041-210X.13305

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Example: Seasonal mortality in wildlife

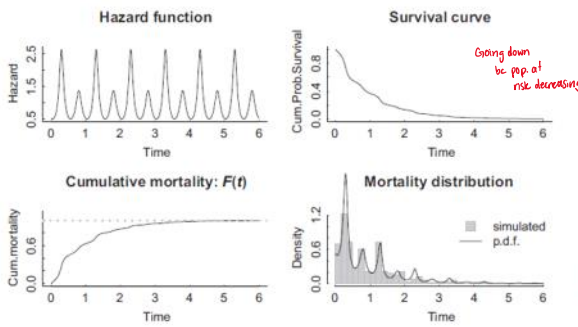


FIGURE 2 Example of a simulated multi-seasonal periodic mortality process, outputted by the `simulate_cyclosurv()` function

Check your understanding: Why do the peaks of the pdf decline over time?

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Cumulative hazard function

- How much hazard has accumulated between time 0 and time t ?

$$H(t) = \int_{u=0}^{u=t} h(u) du$$

- Has a convenient relationship to $S(t)$

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



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Any one function fully describes the distribution...

Survival function

$$\begin{aligned} S(t) &= \Pr(T > t) \\ S(t) &= 1 - F(t) \\ S(t) &= e^{-H(t)} \end{aligned}$$

Cumulative distribution function

$$F(t) = \Pr(T \leq t)$$

Probability density function

$$\begin{aligned} f(t) &= \frac{d}{dt} F(t) \\ f(t) &= h(t)S(t) \end{aligned}$$

Hazard function

$$\begin{aligned} h(t) &= \frac{f(t)}{S(t)} \\ h(t) &= -\frac{d}{dt} \ln[1 - F(t)] \\ h(t) &= -\frac{d}{dt} \ln S(t) \end{aligned}$$

Cumulative hazard function

$$\begin{aligned} H(t) &= \int_{u=0}^{u=t} h(u) du \\ H(t) &= -\log S(t) \end{aligned}$$

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KM is non-parametric, but:

Common parametric survival distributions

- Failure time random variable T ($T \geq 0$)
- $T \sim \text{Exponential}(\lambda)$ or $T \sim \text{Exp}(\lambda)$
- $T \sim \text{Weibull}(\lambda, \gamma)$
- $T \sim \text{LogLogistic}(\lambda, \gamma)$
- There are other parametric survival distributions out there (e.g. gamma, Gompertz-Makeham, log-normal, generalized F, Pareto), but we won't discuss these in this course



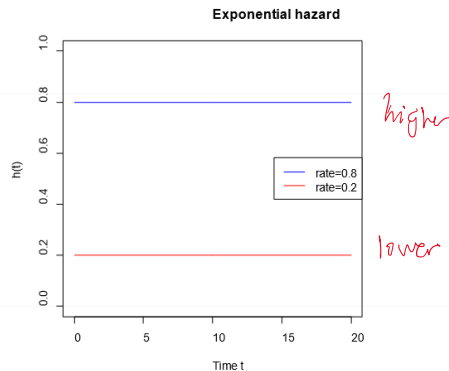
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Exponential distribution

- Constant hazard function

$$h(t) = \lambda$$

- λ is called the **rate parameter**
- For the exponential distribution, λ is also the hazard rate



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Exponential distribution

- “Memoryless” *bc risk of failure doesn't depend on the past*
- It can be hard to justify the constant hazard assumption in practice
- **Examples of exponential distributions in the real world:**
 - Time until an earthquake occurs
 - Length (in minutes) of long-distance business telephone calls
 - The amount of time (in months) a car battery lasts
 - The amount of time (in minutes) a postal clerk spends with a customer



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Weibull distribution

- By adding a second parameter, we allow for greater flexibility in our hazard function

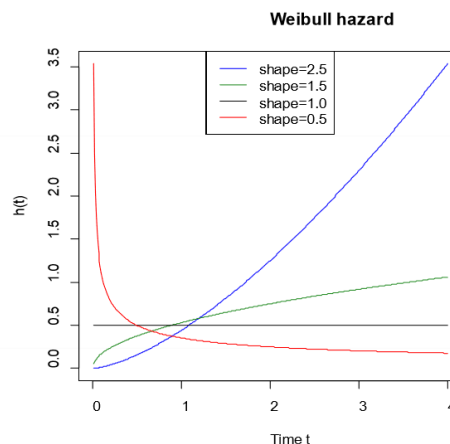
$$h(t) = \lambda\gamma(\lambda t)^{\gamma-1}$$

- λ is still called the **rate parameter**
- γ is called the **shape parameter** *→ second parameter alters the shape*
- Note that the hazard rate for the Weibull distribution is not λ , but rather the hazard rate is calculated from λ, γ, t *Hazard rate is not λ !*

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Weibull distribution

- More flexible than the exponential
- The Weibull distribution accommodates three distinct possibilities*:
 1. If something is going to fail it will most likely fail at the start
 2. The rate of failure is fairly constant
 3. Failure becomes more likely as time goes on.



*<https://doi.org/10.1111/j.1740-9713.2018.01123.x>

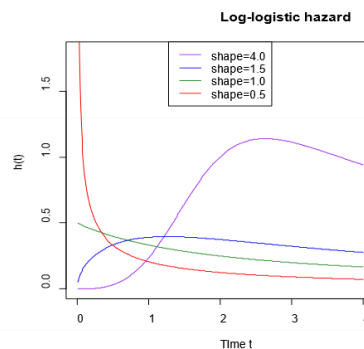
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Log-logistic distribution

Even more complex

- Another (different) two-parameter distribution
not a generalization of exponential,
no flat line $h(t) = \frac{\lambda\gamma(\lambda t)^{\gamma-1}}{1 + (\lambda t)^\gamma}$

$\left\{ \begin{array}{ll} \text{decreasing from } \infty, & \text{if } \gamma < 1 \\ \text{decreasing from } \lambda, & \text{if } \gamma = 1 \\ \text{increasing then decreasing,} & \text{if } \gamma > 1 \end{array} \right.$



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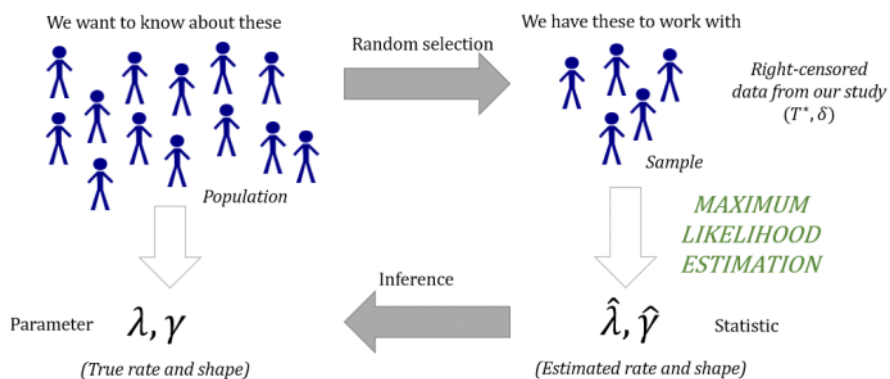
Distribution	Hazard Function	Cumulative Hazard Function	Survival Function
Any	$h(t)$	$H(t) = \int_0^t h(u) du$	$S(t) = e^{-H(t)}$

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Distribution	Hazard Function	Cumulative Hazard Function	Survival Function
Exponential	$h(t) = \lambda$	$H(t) = \lambda t$	$S(t) = e^{-\lambda t}$
Weibull	$h(t) = \lambda \gamma (\lambda t)^{\gamma-1}$	$H(t) = (\lambda t)^\gamma$	$S(t) = e^{-(\lambda t)^\gamma}$
Log-logistic	$h(t) = \frac{\lambda \gamma (\lambda t)^{\gamma-1}}{1 + (\lambda t)^\gamma}$	$H(t) = \log(1 + (\lambda t)^\gamma)$	$S(t) = \frac{1}{1 + (\lambda t)^\gamma}$

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Statistical inference

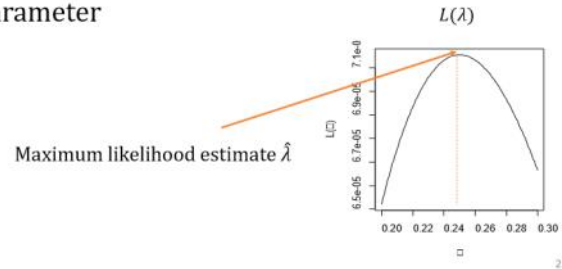


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<https://www.cliffsnotes.com/study-guides/statistics/sampling/populations-samples-parameters-and-statistics>

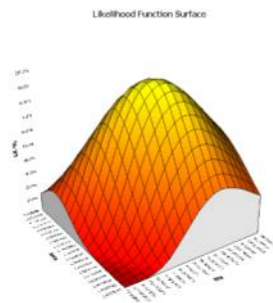
Maximum likelihood estimation

- Likelihood function $L(\lambda)$ is the probability of observing data if the true parameter is λ
- Function of the *data* and the *parameter(s)*
- Example with one parameter



Maximum likelihood estimation

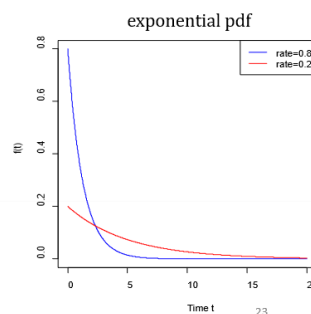
- If there is more than one parameter, then the maximum likelihood estimate is where all parameters are simultaneously maximized
- Where does $L(\lambda, \gamma)$ maximize?



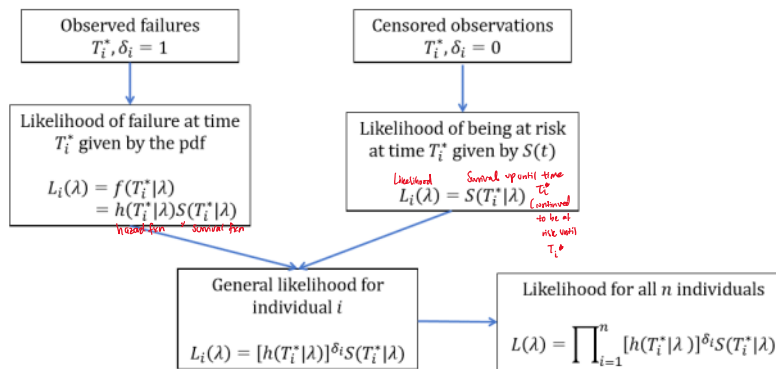
Building the likelihood function

- General likelihood function for time-to-event data
- If there were NO censoring in the data:

$$L(\lambda) = \prod_{i=1}^n f(T_i|\lambda)$$



Likelihood for right-censored data



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General Likelihood: $L(\lambda) = \prod_{i=1}^n [h(T_i^* | \lambda)]^{\delta_i} S(T_i^* | \lambda)$

Distribution	Hazard Function	Survival Function	Likelihood Function
Exponential	$h(t) = \lambda$	$S(t) = e^{-\lambda t}$	$L(\lambda) = \prod_{i=1}^n \lambda^{\delta_i} e^{-\lambda T_i^*}$ <i>hazard fun raised to delta_i</i> <i>survival fun</i>
Weibull	$h(t) = \lambda \gamma (\lambda t)^{\gamma-1}$	$S(t) = e^{-(\lambda t)^\gamma}$	$L(\lambda, \gamma) = \prod_{i=1}^n [\lambda \gamma (\lambda T_i^*)^{\gamma-1}]^{\delta_i} e^{-(\lambda T_i^*)^\gamma}$
Log-logistic	$h(t) = \frac{\lambda \gamma (\lambda t)^{\gamma-1}}{1 + (\lambda t)^\gamma}$	$S(t) = \frac{1}{1 + (\lambda t)^\gamma}$	$L(\lambda, \gamma) = ?$ <i>will not write out for in-class activity</i>

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In practice...

- Most of the time we will rely on statistical software to compute the maximum likelihood estimate
- For the exponential, we can derive the maximum likelihood estimate from the likelihood function

- The maximum likelihood estimate is the incidence rate:

$$\hat{\lambda} = \frac{\sum_{i=1}^n \delta_i}{\sum_{i=1}^n T_i^*}$$

total # failures
total amount of person-time

will prove this in HW

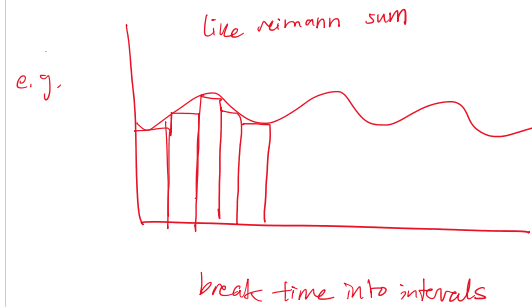
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Another nonparametric estimator of $S(t)$

- With a nonparametric estimator of $H(t)$, we can estimate $S(t)$

$$H(t) = \int_{u=0}^{u=t} h(u) du$$

- What if we broke time into small intervals, and assumed that the hazard is constant within each interval?



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Interval

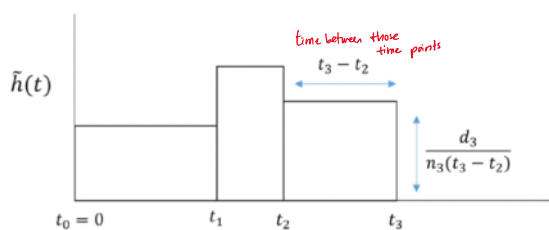
- Use the same time intervals as the Kaplan-Meier estimator
- Within each interval, we assume the hazard rate is constant and estimate the incidence rate

$$\frac{\text{\# of events in the interval}}{\text{person-time follow-up in the interval}} = \frac{d_j}{n_j(t_j - t_{j-1})}$$

failures (pointing to d_j)
pp't at risk (pointing to n_j)
one interval (pointing to $t_j - t_{j-1}$)

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Nelson-Aalen estimator



Each rectangle has area:

$$\frac{d_j}{n_j(t_j - t_{j-1})} \times (t_j - t_{j-1}) = \frac{d_j}{n_j}$$

The estimated cumulative hazard is:

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

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Example: AIDS hemophiliac cohort

Ordered follow-up times: 2, 3+, 6, 6, 8, 10+, 15, 15, 16, 27, 30, 32 months

Unique failure/ censoring time	Number at risk n_j during $(t_{j-1}, t_j]$	Number of deaths d_j at t_j	Number censored c_j at t_j	Cumulative hazard contribution $\frac{d_j}{n_j}$	Nelson-Aalen estimate
$t_0 = 0$					$t = [0, 2)$ $\hat{H}(t) = 0$
$t_1 = 2$	$n_1 = 12$	$d_1 = 1$	$c_1 = 0$	$\frac{d_1}{n_1} = \frac{1}{12}$	$t = [2, 3)$ $\hat{H}(t) = 0.083$
$t_2 = 3$	$n_2 = 11$	$d_2 = 0$	$c_2 = 1$	$\frac{d_2}{n_2} = \frac{0}{11}$	$t = [3, 6)$ $\hat{H}(t) = 0.083$
$t_3 = 6$	$n_3 = 10$	$d_3 = 2$	$c_3 = 0$	$\frac{d_3}{n_3} = \frac{2}{10}$	$t = [6, 8)$ $\hat{H}(t) = 0.283$

↳ At $t=3$, before
anyone fails,
cannot look to the future

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Breslow estimator

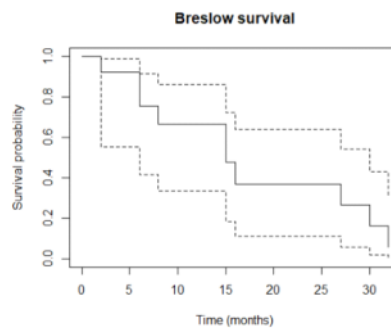
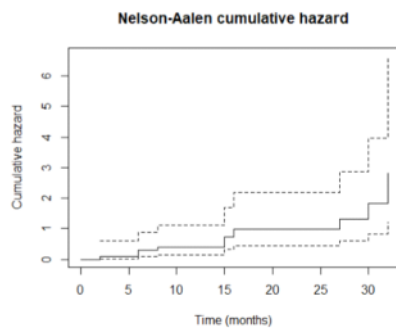
- Use the Nelson-Aalen estimator to estimate survival

$$\tilde{S}(t) = \exp(-\tilde{H}(t))$$

- Produces an estimate of survival that is similar to, though not identical to, the Kaplan-Meier estimator

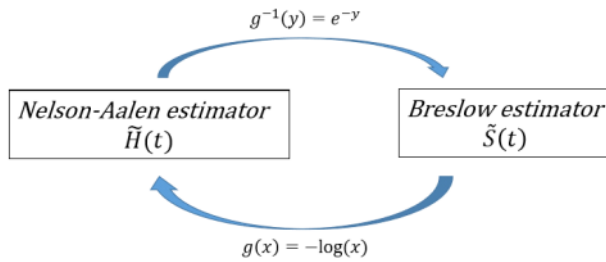
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Nelson-Aalen curve example



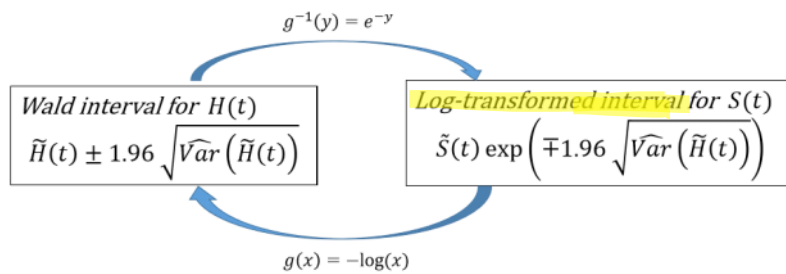
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Point estimation



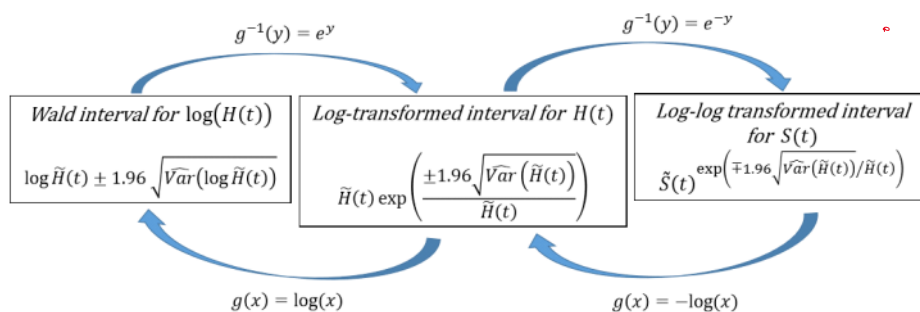
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Confidence intervals



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Confidence intervals, continued



Complementary log-log: $\log(-\log(x))$

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→ how can we connect this to 36

The NEW ENGLAND JOURNAL OF MEDICINE

ADVERTISEMENT IN THIS ISSUE MAY 15, 2002 VOL. 346, NO. 20

Prostate Cancer Mortality at 11 Years of Follow-up

Ernst H. Schemper, M.D., James Huggins, M.D., Margaret J. Rhodes, M.D., Yasuo I. Tamaoka, M.D., Stefano Cefis, M.D., Vito Lenzi, M.D., Marco Rivaroli, M.D., Marco Logani, M.D., Hans Lilja, M.D., Marco Zappa, M.D., Victor D'Amico, M.D., Franco Bazzocchi, M.D., Alvaro Paez, M.D., Luis Montano, M.D., Chien H. Cheng, M.D., Giammarco Ricci, M.D., Nigel Cunniff, M.D., Arnold Hirsch, M.D., John R. Stellard, M.D., Thomas von der Kietz, M.D., Fred W. Ruzza, M.D., Bert G. Monstorf, M.D., Ulf Håkan Larsson, M.D., Anders Therasme, M.D., Kenneth Taylor, M.D., Marcia Hakola, Ph.D., Sue M. Klein, Ph.D., Henry J. Iscovich, M.D., and Paolo Manter, M.D. (For the EORTC Investigators)

- **Goal:** Investigators conducted a large **randomized** study to examine the value of monitoring **prostate-specific antigen (PSA)** as part of routine screening for reducing prostate-cancer mortality.
Therneau C, M.D., Vera-Rubio, M.D., Morton-Russell, M.D., Morton-Lucas, M.D., Hains-Burns, M.D., Lacey, G.M., Evans-Richter, M.D., Moore-Paul, M.D., Lissa-Mattin, M.D., Bingham, M.D., Gorman-Ross, M.D., Sigurd-Carlsson, M.D., Arnold-Vitres, M.D., Kavanagh, M.D., Bostrom, M.D., Pineda-Munoz, M.D., Serrin, G., Björnstrand, M.D., Gill-Fabian, M.D., Andrews-Stiles, M.D., Kinnison-Tyler, M.D., Manti-Nakanga, Ph.D., Saito M. Moun, Ph.D., Harty J. and Anne Arnesen, M.D. for the EPOPC investigators*
- **Population:** The study included **162,388 men** between the ages of 55 and 69 years at entry randomized to receive either **PSA-based screening or standard screening** (control group). The trial was conducted in eight European countries.

Source: Schroder et al. (2012) NEJM [10.1056/nejmoa1113135](https://doi.org/10.1056/nejmoa1113135)

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was **date of prostate-**
al registries to identify
with prostate cancer
ndomization.
lysis, the only predictor
PSA-based screening or
Nelson-Aalen method
ath from prostate cancer.

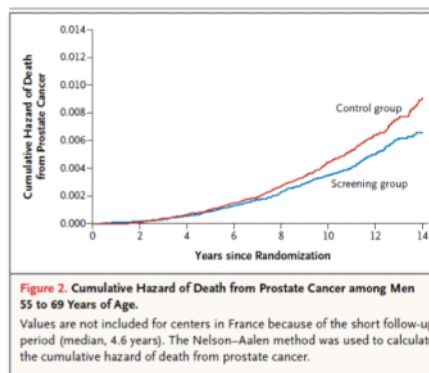
- **Outcome variable:** The primary outcome was **date of prostate-cancer mortality**, assessed using national registries to identify the official cause of death in participants with prostate cancer diagnosis. The time origin was time of randomization.
- **Predictor variables:** For the primary analysis, the only predictor variable considered was screening arm (PSA-based screening or control).
- **Statistical analysis:** Researchers used the **Nelson-Aalen method** to calculate the cumulative hazard of death from prostate cancer.

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The graph displays the cumulative hazard of death from prostate cancer for two groups over a 14-year period. The x-axis represents 'Years since Randomization' from 0 to 14. The y-axis represents the cumulative hazard, ranging from 0.000 to 0.014. The 'Control group' is represented by a red line, and the 'Screening group' is represented by a blue line. Both curves start at (0,0) and increase over time. The Control group curve is consistently higher than the Screening group curve, indicating a higher cumulative hazard. The curves are relatively flat until around year 6, after which they rise more steeply. The Control group curve reaches approximately 0.009 at year 14, while the Screening group curve reaches approximately 0.0065.

Years since Randomization	Control group (Cumulative Hazard)	Screening group (Cumulative Hazard)
0	0.000	0.000
2	0.0005	0.0005
4	0.0010	0.0010
6	0.0015	0.0015
8	0.0025	0.0020
10	0.0040	0.0030
12	0.0060	0.0045
14	0.0090	0.0065

- **Results:** Figure 2 summarizes the Nelson-Aalen cumulative hazard curve. These two curves begin to gradually separate starting approximately 7 years after randomization. Authors note that there is evidence that PSA-based screening significantly reduced mortality from prostate cancer but did not affect all-cause mortality (*not shown in figure*).



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Looking ahead

- The hazard function is the basis of the **Cox proportional hazards model** – the most popular regression model for time-to-event data
- Regression models allow us to model the effects of multiple covariates simultaneously, including continuous covariates
- Next week will be the first of several weeks on the Cox proportional hazards regression model

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Today's activity

- Small groups
- Wordle word problem!
- More transformed confidence intervals
- Sketching hazard functions

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