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CBB Workshop: Introduction to Survival Analysis with Practical Applications in R

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Center for Bioinformatics and Biostatistics
2024-03-19

CBB offers

- **Mentoring of PhD students and postdocs**
- Seminar series and workshops
- Support in project planning and grant applications
- Joint employments including co-financing
- Networking opportunities to discuss bioinformatical and biostatistical methods and problems
 - **Drop-in on Thursdays 13:00–15:00 in Neo, room Protein**
- Please see www.ki.se/cbb for more information, or contact us at cbb@ki.se

Agenda

- 1st Session 13:00–14:00
 - The basics of survival analysis
- Coffee Break 14:00–14:30
- 2nd Session 14:30–16:00
 - Practical application in R
 - examples, graphics, and exercises

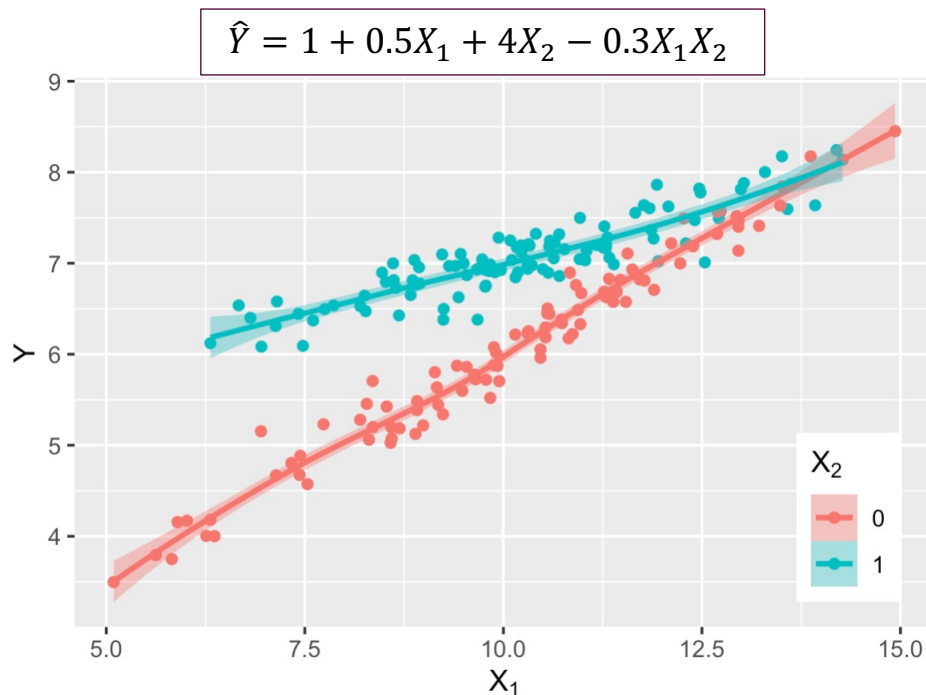
Introduction to Survival Analysis

Outline

- The most important concepts of survival analysis
 1. The type of data we use
 2. The survival function & the cumulative incidence function
 3. The hazard rate function
 4. The Cox proportional hazards regression model

Our Typical Statistical Analysis

- Typical data & research question relationship:
 - We measure some **Outcome variable (Y)**
 - We measure a set of **Covariates (X_1, X_2, \dots)**
 - Exposure/risk factor variables potentially associated with the outcome
 - We perform some statistical analysis—**Regression Model**



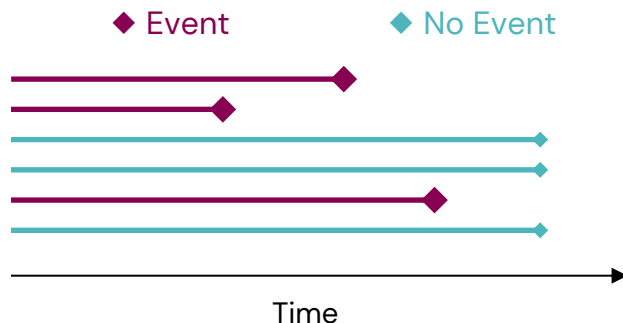
What is Survival Analysis?

- Survival Analysis is a special family of statistical analysis
- **The outcome is time**
- We measure the time until a particular event
 - Death (Mortality)
 - Diagnosis of a particular disease (Incidence)
 - Progression to a stage of disease
 - etc.



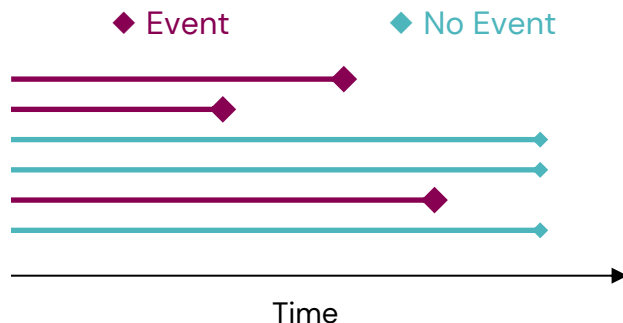
Time-to-Event Data

- Time-to-event outcome:
 1. The time duration we followed each subject
 2. Whether or not they experienced the event



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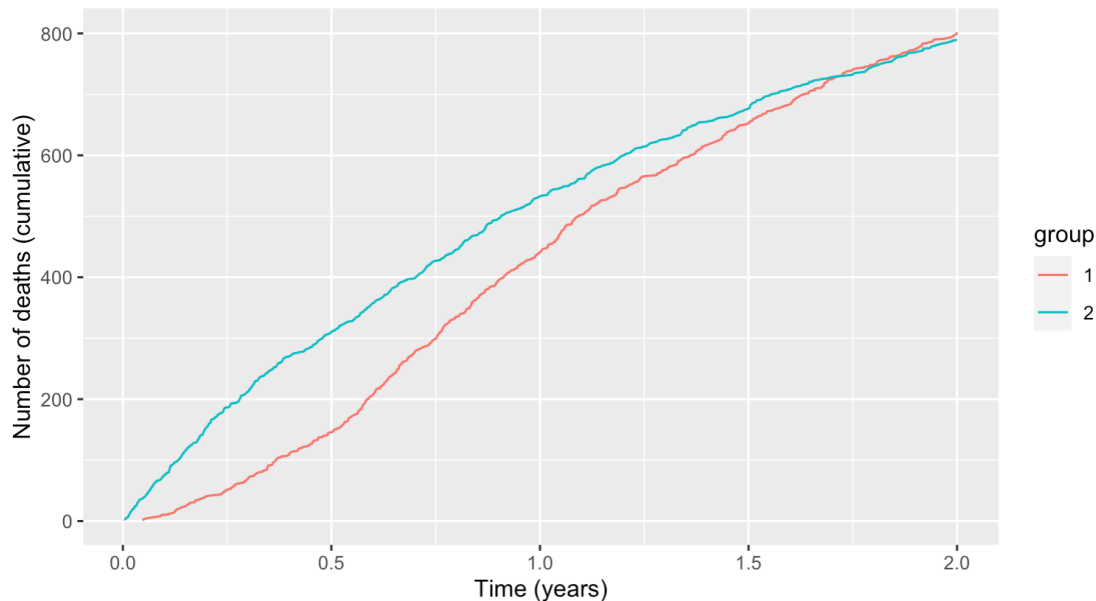


For example:

id	time	event	age	sex	bmi	masld
23	6.08	0	32	female	35.2	no
24	6.79	0	43	female	24.5	no
25	3.65	0	39	male	24.3	no
27	0.59	1	62	female	30.2	no
32	11.67	0	45	male	22.1	no
33	2.03	1	58	male	26.7	no
36	1.13	0	53	female	24.8	no
37	7.59	0	47	male	21.9	no
38	9.11	0	54	female	22.1	no
40	4.67	1	61	female	28.4	no
41	6.10	1	45	female	26.3	no

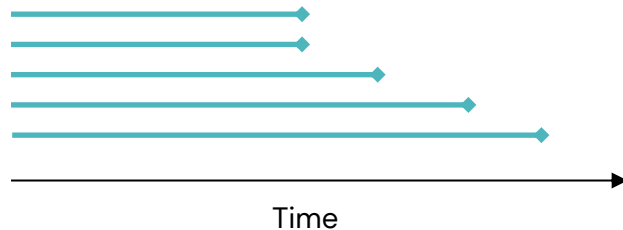
Why two-part outcome?

- Can't we just look at the final outcome?
- Example:
Mortality over 2 years comparing 2 groups
- Not just **what** happens, but also **when**
- Who experiences more events and/or events sooner?



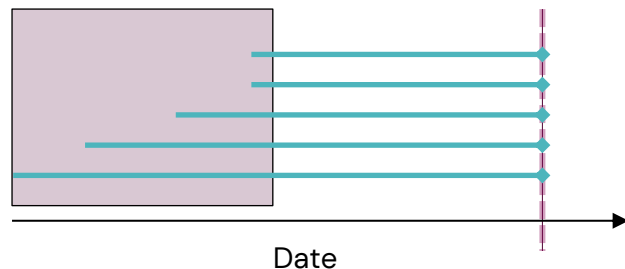
Censoring

- We don't observe events for everyone
 - We end the follow-up at some point
 - (Given enough time, would everyone have the event?)
- **"Censored at time t "** = event occurs sometime after t
 - *i.e. we didn't observe it yet*
 - *a.k.a. "lost to follow-up"*



Censoring

- We don't observe events for everyone
 - We end the follow-up at some point
 - (Given enough time, would everyone have the event?)
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 - *a.k.a. "lost to follow-up"*
- Often due to inclusion/recruitment over time and with a fixed end date



Now What?

- **What do we want to know?**
- In a typical non-survival analysis:
 1. Summary statistics (mean/median, proportions) including
 - standard errors (confidence intervals)
 - preliminary comparisons between groups (graphically or with statistical tests)
 2. Proper regression model
 - Multivariate model
 - Effect sizes (mean differences, odds ratios, etc.) and confidence intervals
 - Statistical tests for association with the outcome

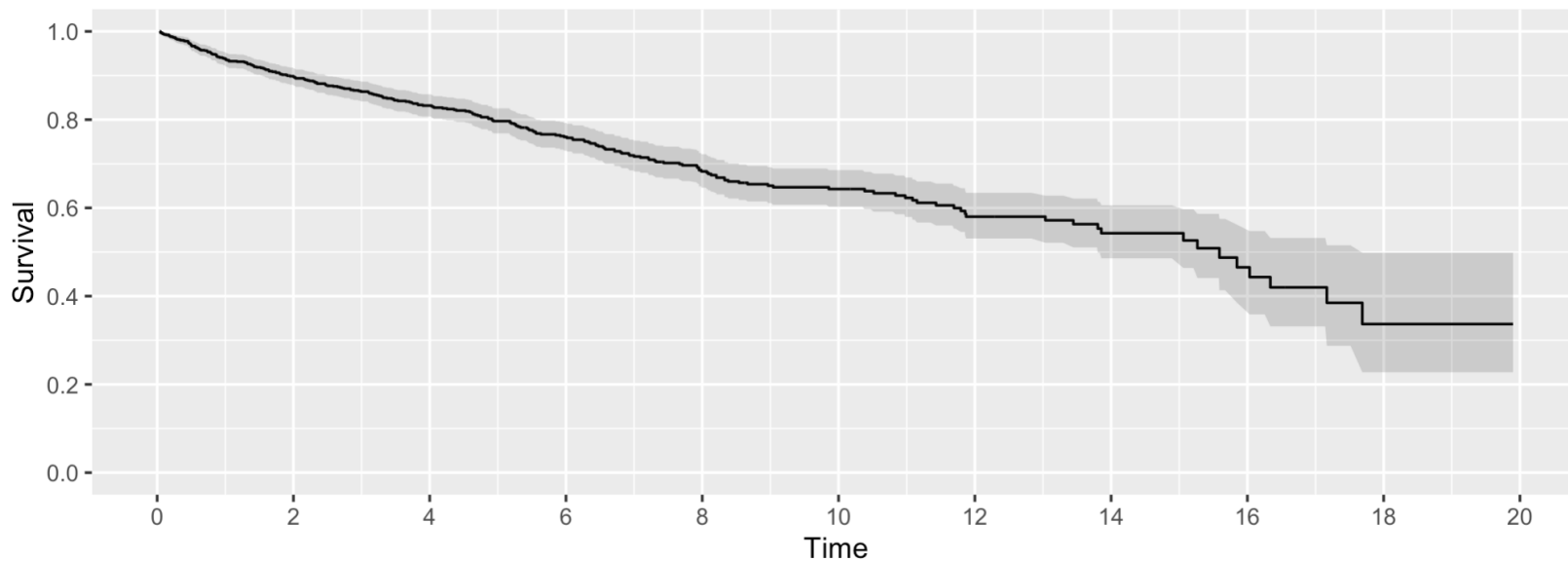
The Survival Function

The Survival Function

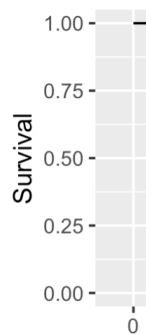
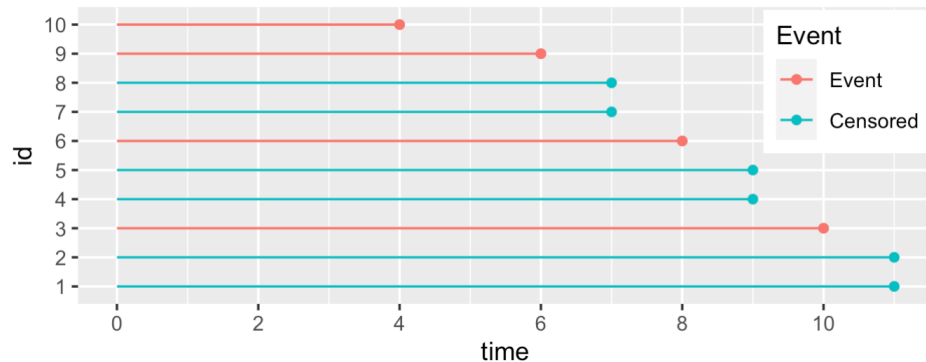
- If we measure the time to event, a reasonable question might be:
- *“What is the probability of not having the event by time t ?”*
or
“What proportion of the population will not have the event by time t ?”
- The survival function $S(t)$: The probability of being event-free by time t
- It's a probability which decreases over time
- $S(0) = 1$ if $t_1 < t_2$ then $S(t_1) \geq S(t_2)$
- *Use the time-to-event data to estimate:*

Kaplan-Meier Estimate

- Kaplan-Meier estimate of the survival function (with 95% confidence int.)

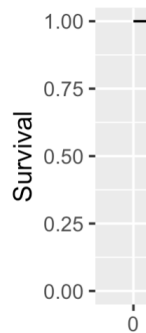
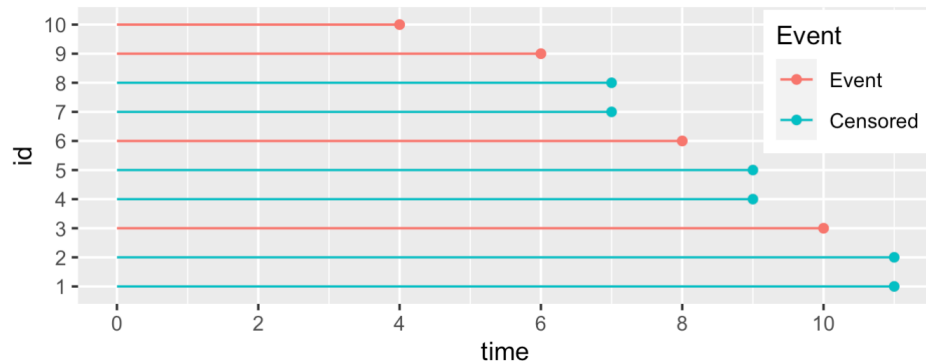


Calculating the Survival Function



Survival calculation:

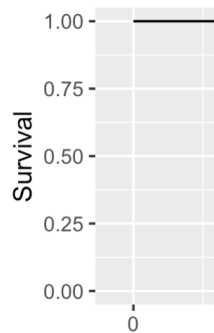
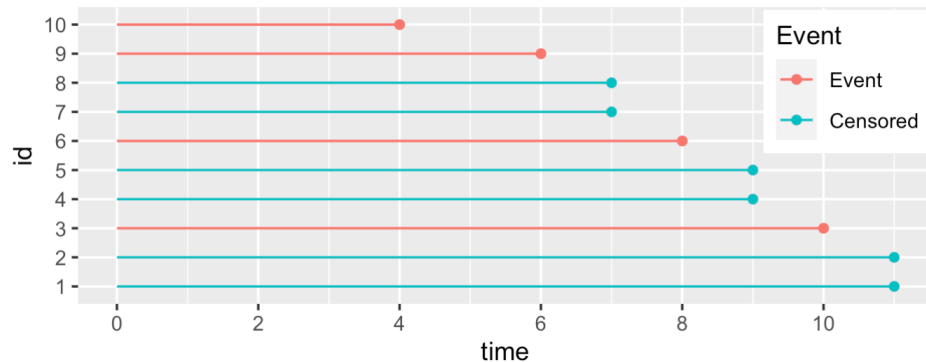
Calculating the Survival Function



Survival calculation:

- $S(0) = \frac{10}{10} = 1$

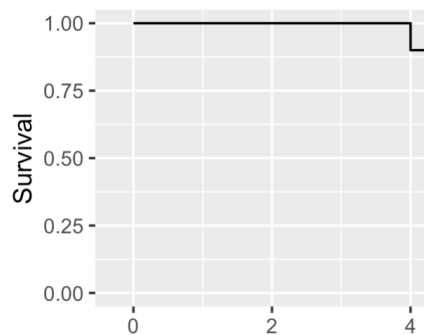
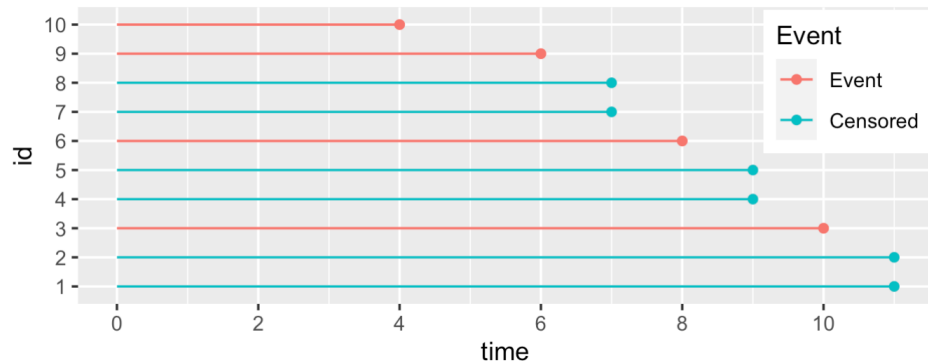
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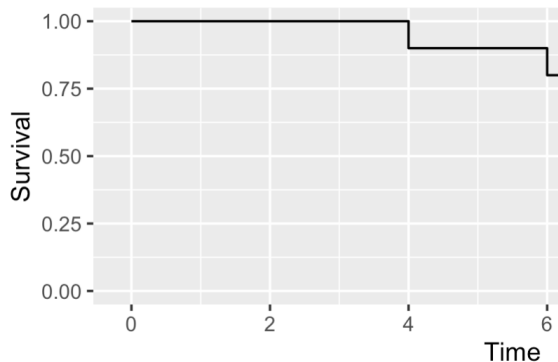
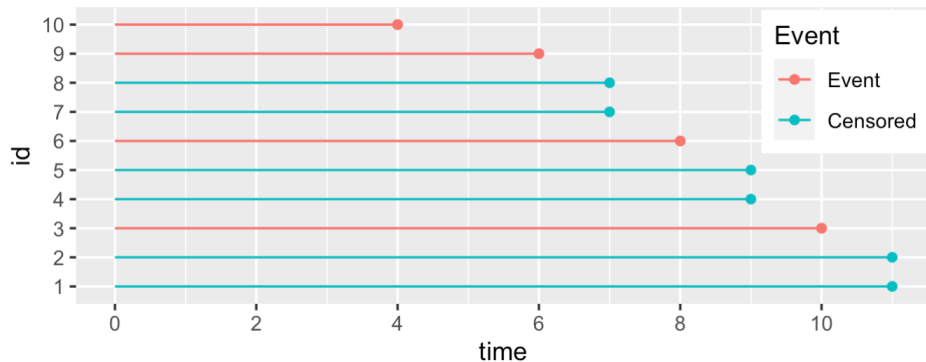
Calculating the Survival Function



Survival calculation:

- $S(0) = \frac{10}{10} = 1$
- $S(1) = \frac{10}{10} = 1$
- $S(4) = \frac{9}{10} = 0.9$

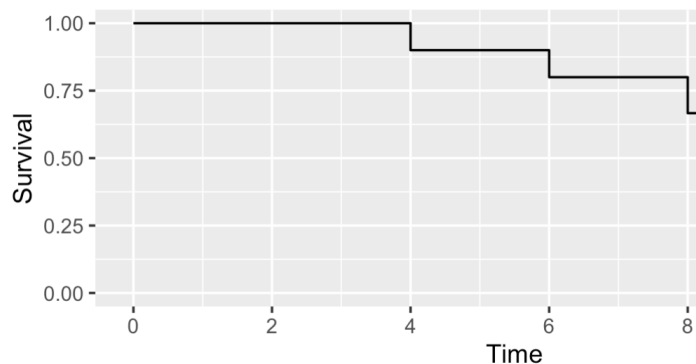
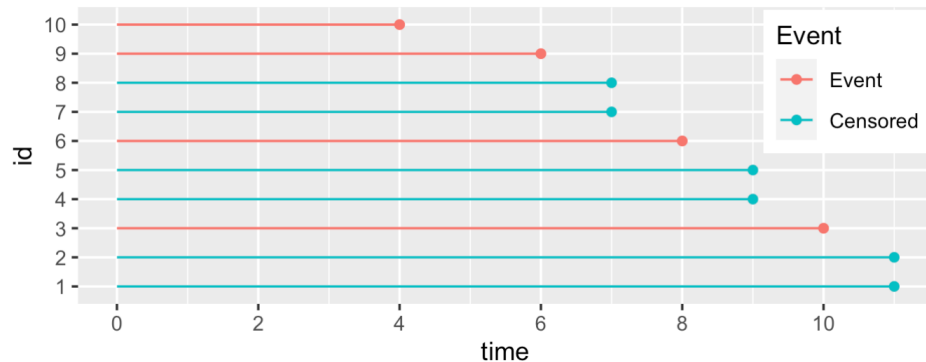
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- $S(6) = S(4) \cdot \frac{8}{9} = 0.9 \cdot 0.89 = 0.8$

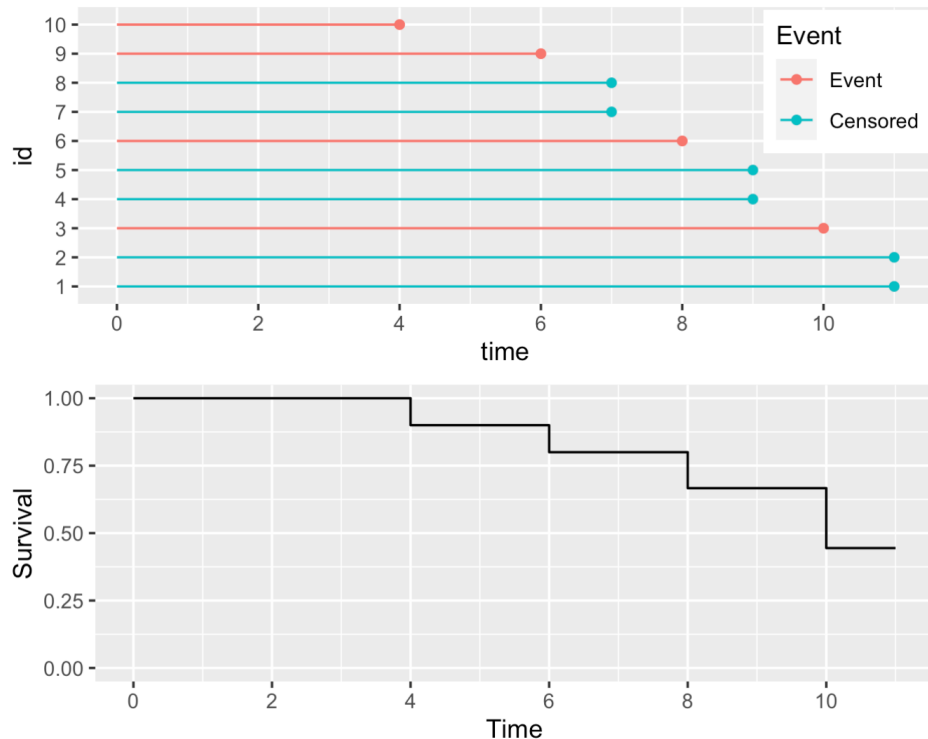
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- $S(8) = S(6) \cdot \frac{5}{6} = 0.8 \cdot 0.83 = 0.67$

Calculating the Survival Function

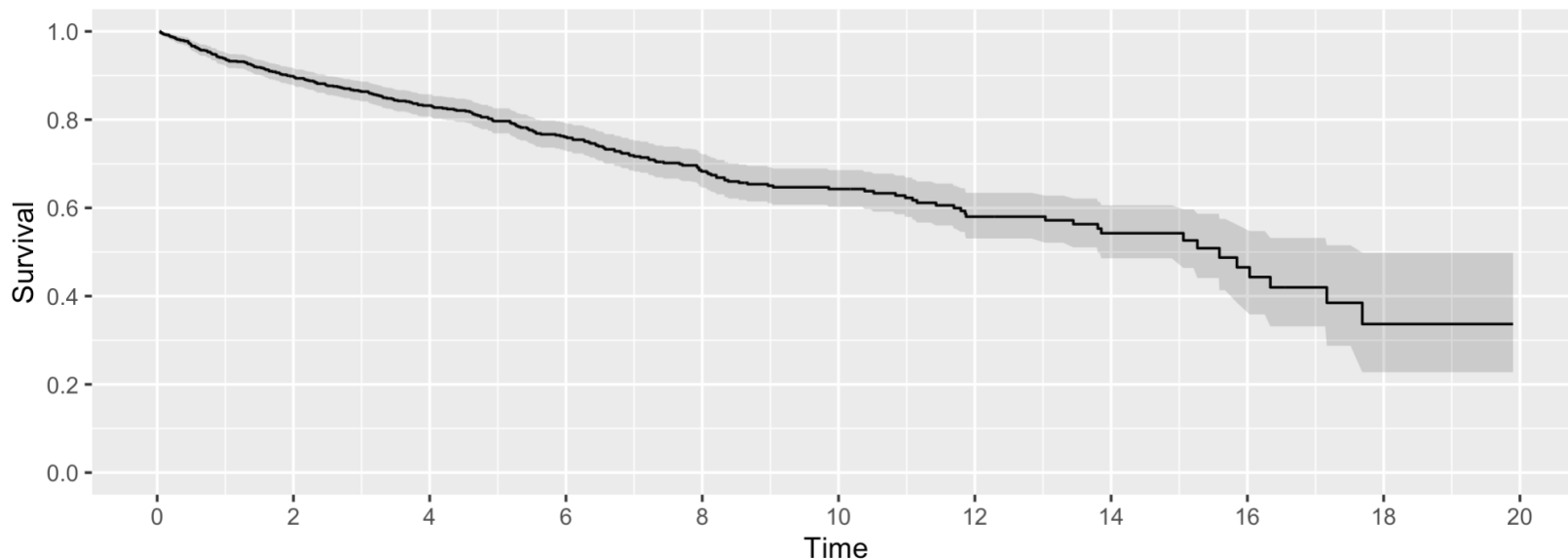


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- $S(8) = S(6) \cdot \frac{5}{6} = 0.8 \cdot 0.83 = 0.67$
- $S(10) = S(8) \cdot \frac{2}{3} = 0.67 \cdot 0.67 = 0.44$

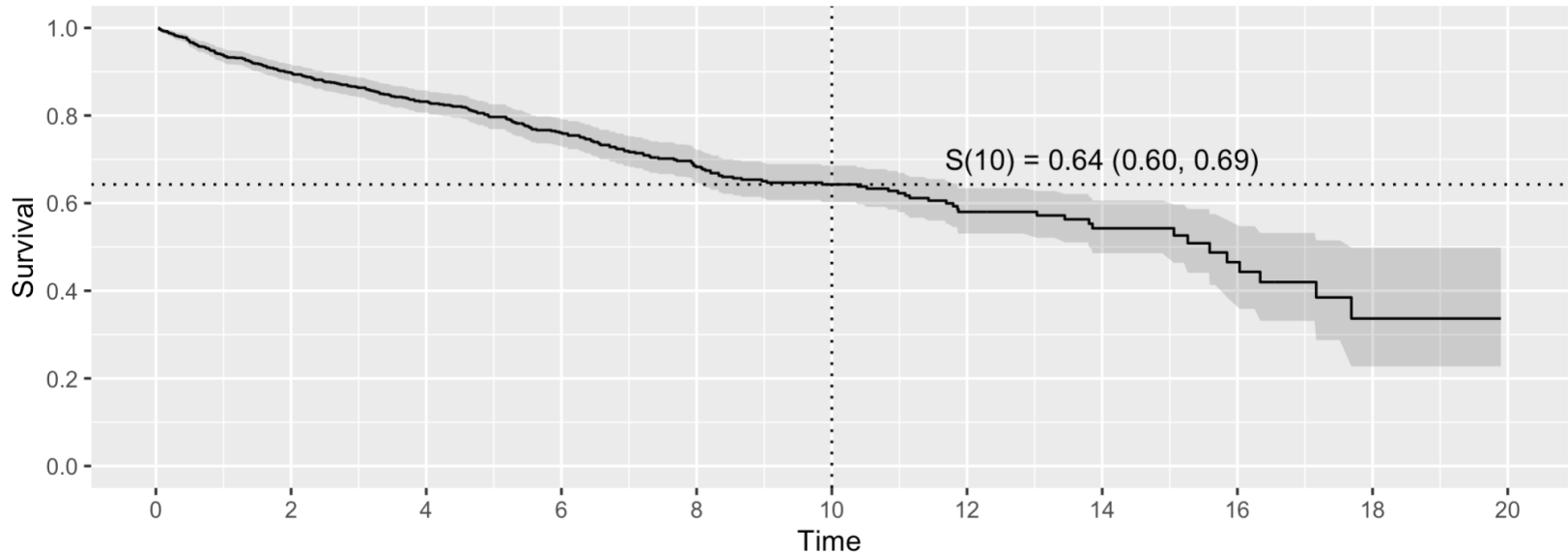
Kaplan-Meier Estimate

- Kaplan-Meier estimate of the survival function (with 95% confidence int.)



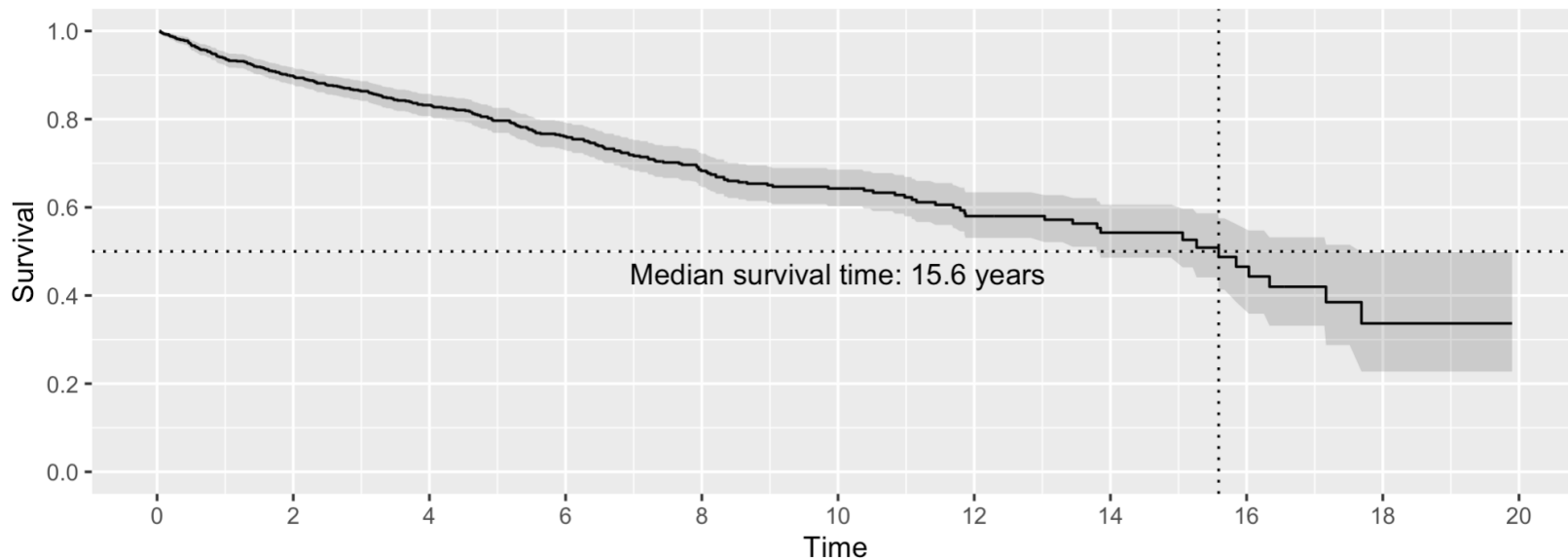
Kaplan-Meier Estimate (2)

- Kaplan-Meier estimate of the survival function (with 95% confidence int.)



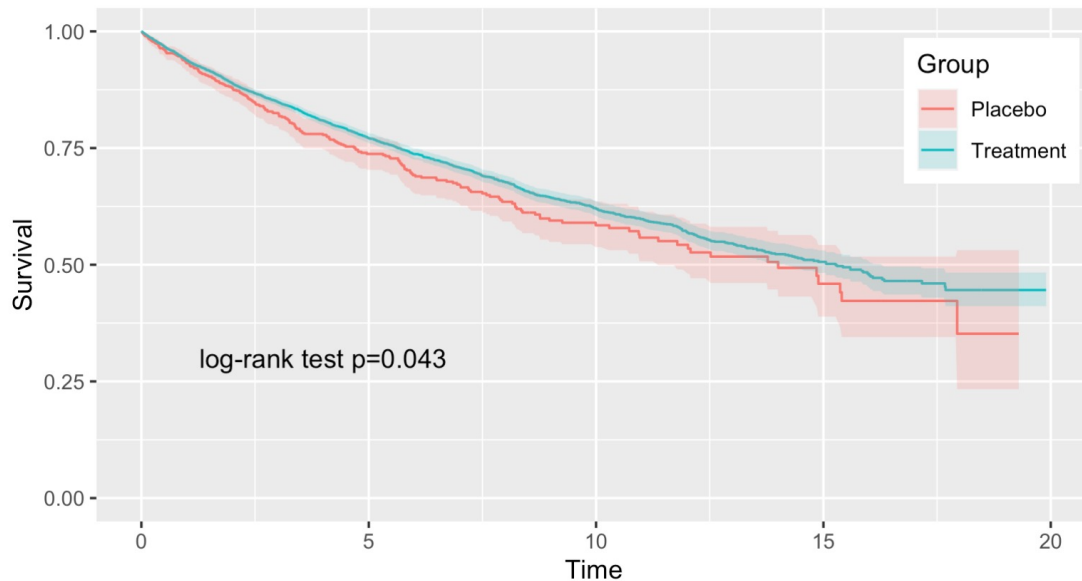
Median survival time

- Another interesting statistic could be the **Median Survival Time**:
 - The time until 50% have experienced the event.
 - For which t is $S(t)=0.5$?



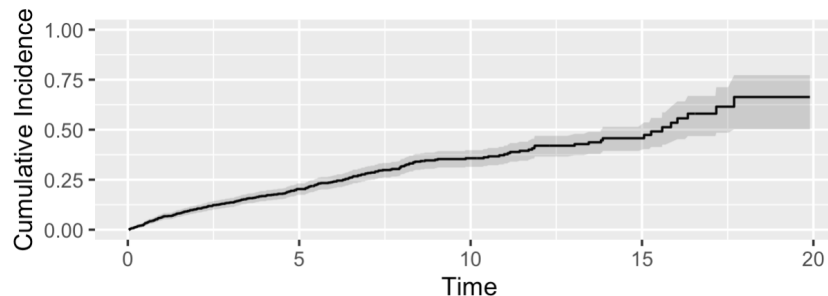
Comparing Survival Curves

- We can compare survival curves between groups
- We can do a statistical test—the **log-rank test**—for difference between the curves



The Cumulative Incidence Function

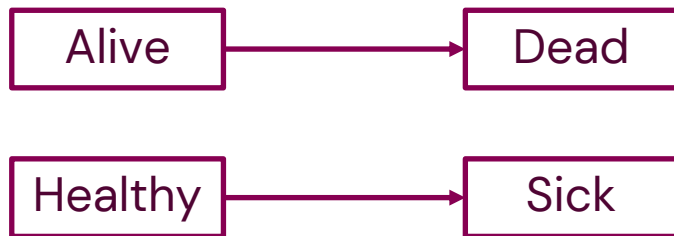
- “The risk”
- $1 - S(t)$
- The proportion who **have had the event** by t
- or equivalently the probability of having the event by time t
- Study Survival or Cumulative Incidence depending on context
 - Which more interesting: the event-havers or the event-free?
 - Cancer researchers favor survival (often survival after cancer dx)
 - Hepatology favors cumulative incidence (who is at greater risk of dx)



The Hazard Rate

Different View: Transition Between States

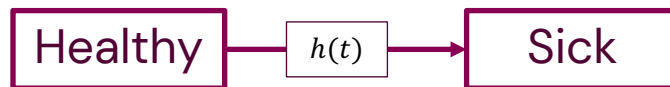
- Subject starts in one state and at some timepoint moves to a different state



- We study the transition (the arrow)
- $S(t)$ is the proportion still in the 1st state at time t

The Hazard Rate

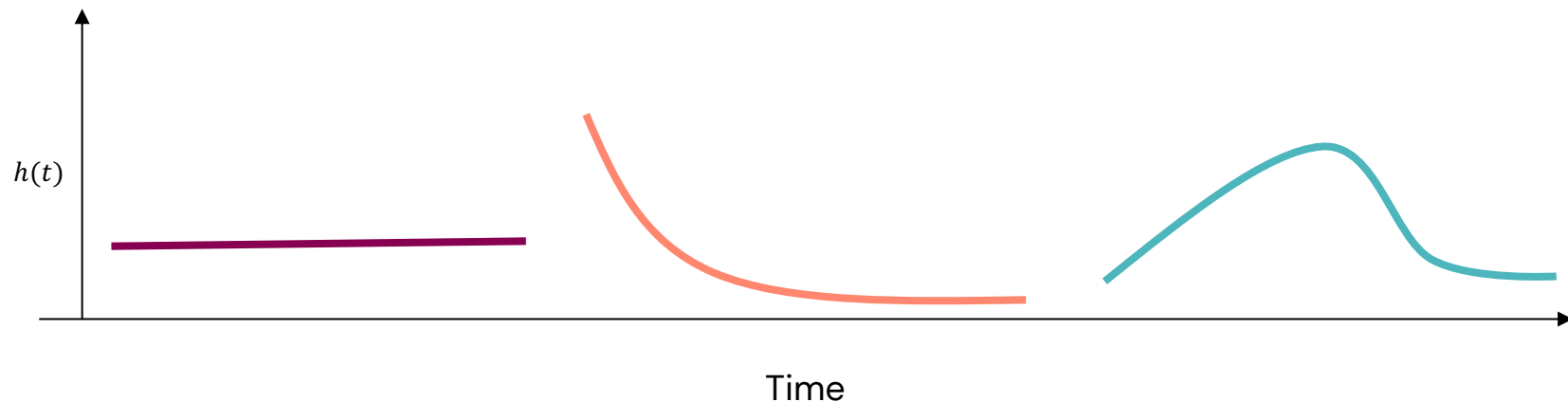
- The rate of transition is determined by the hazard rate
- Hazard rate function = $h(t)$



- The hazard is a velocity
- $h(t) \geq 0$
- Higher hazard rate means both more events over time and events sooner

Hazard Function Examples

- The Hazard function can look very different depending on the mechanics of the transition



The Cumulative Hazard

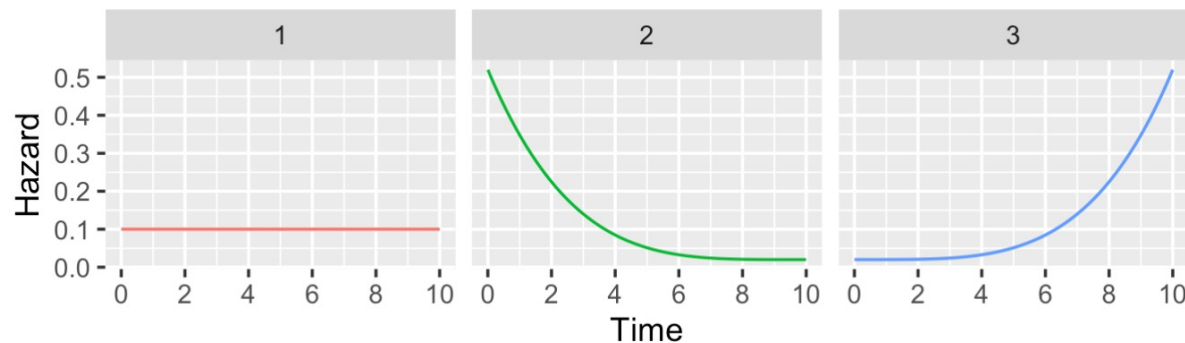
- The total accumulated hazard from 0 to t :

$$H(t) = \int_0^t h(u) du$$

- If the hazard is the velocity, the cumulative hazard is the total distance travelled
- How much hazard were you exposed to?
 - A lot over a short time?
 - A little over a long time?

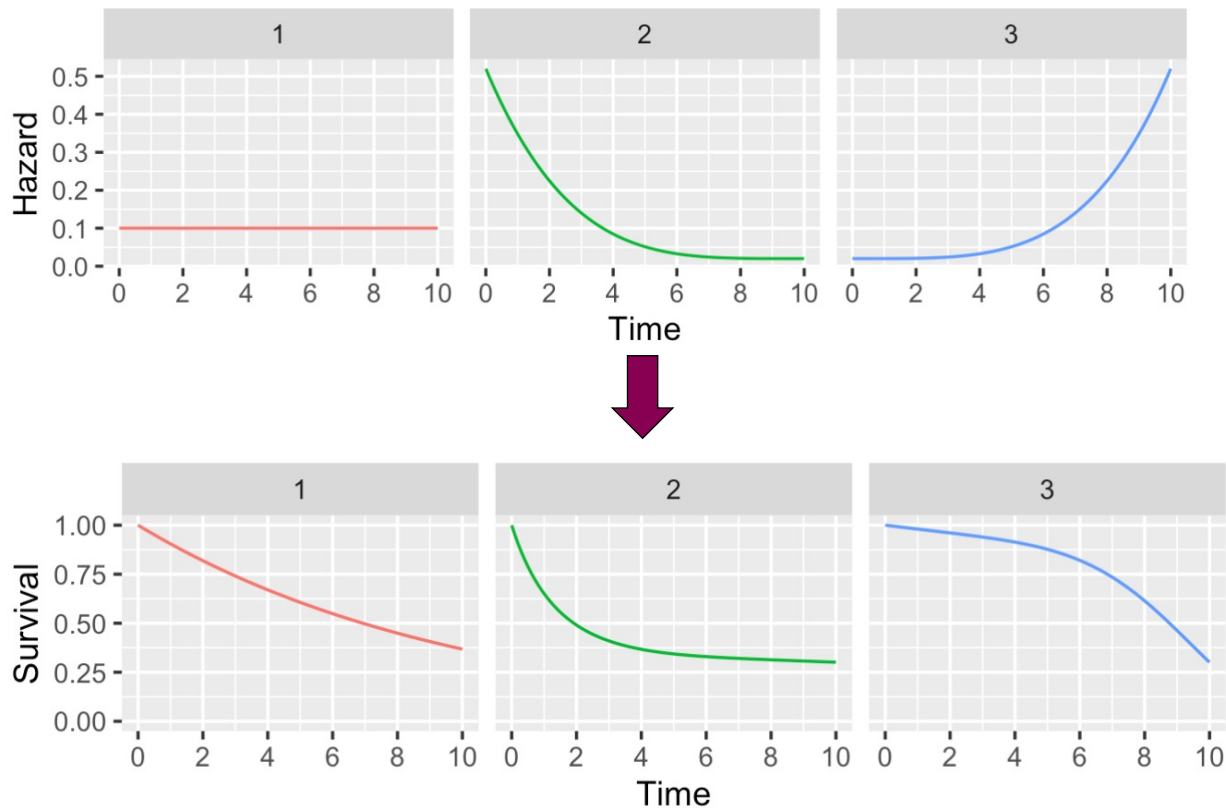
Hazard & Survival

- How is the hazard function related to the survival function?



Hazard & Survival

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

Two Measures of Interest

- When doing survival analysis, we want to study two quantities:
 1. The Survival (or Cumulative Incidence) function over time for summary and interesting comparisons
 - More practical scale
 - e.g. survival after 5 years
 - treatment vs. no treatment
 - by smoking and age group (4 comparisons)
 2. The Hazard function
 - Useful scale for exposure/risk factors
 - Example: carcinogens for developing cancer

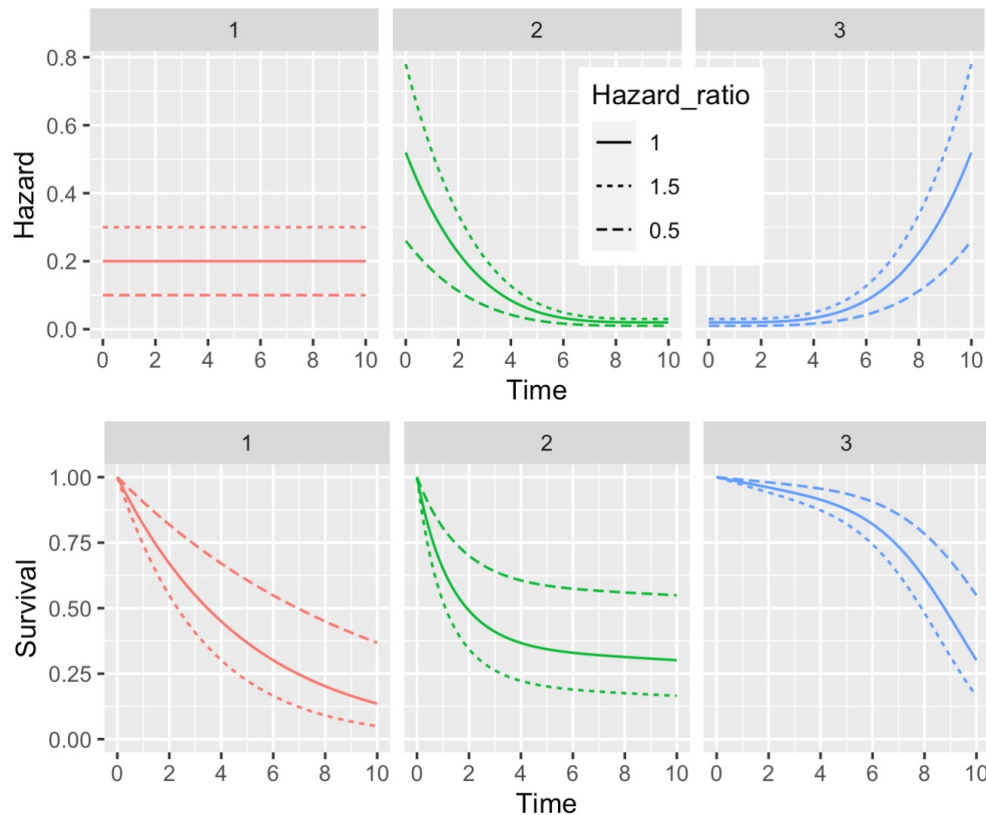
Cox Proportional Hazards Regression

Proportional Hazards

- How do we compare hazards $h_1(t)$ and $h_2(t)$?
- Hazard ratio: $HR(t) = \frac{h_1(t)}{h_2(t)}$
- **Proportional hazards:**
 - If the hazard ratio is constant over time
 - $h_1(t) = HR \cdot h_2(t)$

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Cox Proportional Hazards Regression Models

- The workhorse for inference in survival analysis
- All subjects have an unknown hazard function specific to them
- We impose a rule:
 - There is a common underlying shape to all the hazards, $h_0(t)$
 - All subjects' hazards are then **proportional to $h_0(t)$** according to their covariates/risk factors
- Subject i 's hazard = $h_0(t) \cdot HR_i$

Cox Proportional Hazards Regression Models (2)

- Subject i 's hazard = $h_0(t) \cdot HR_i$
- Cox regression is regression on $\ln(HR_i)$:

$$\ln(HR_i) = \beta_1 X_{1i} + \beta_2 X_{2i} + \cdots \beta_m X_{mi}$$

- The β :s are estimated using the time-to-event outcome and the covariates
- $h_0(t)$ is not estimated! ("semi-parametric", "partial log-likelihood")

Interpreting Cox Models

$$\ln(HR_i) = \beta_1 X_{1i} + \beta_2 X_{2i} + \cdots \beta_m X_{mi}$$

- The β :s are “log-hazard ratios”
- **e^{β_j} is the hazard ratio from a 1 unit increase in covariate X_j**
 - Binary (0/1): Change from treatment group 1 to treatment group 2
 - Continuous: Change in BMI from 25 to 26
 - Multiplicative, e.g. 2 unit change $\Rightarrow e^{\beta_j \cdot 2} = e^{\beta_j} \cdot e^{\beta_j}$, etc.
- Like all other regressions we can get confidence intervals and p-values for each β

Pros and Cons of the Cox Model

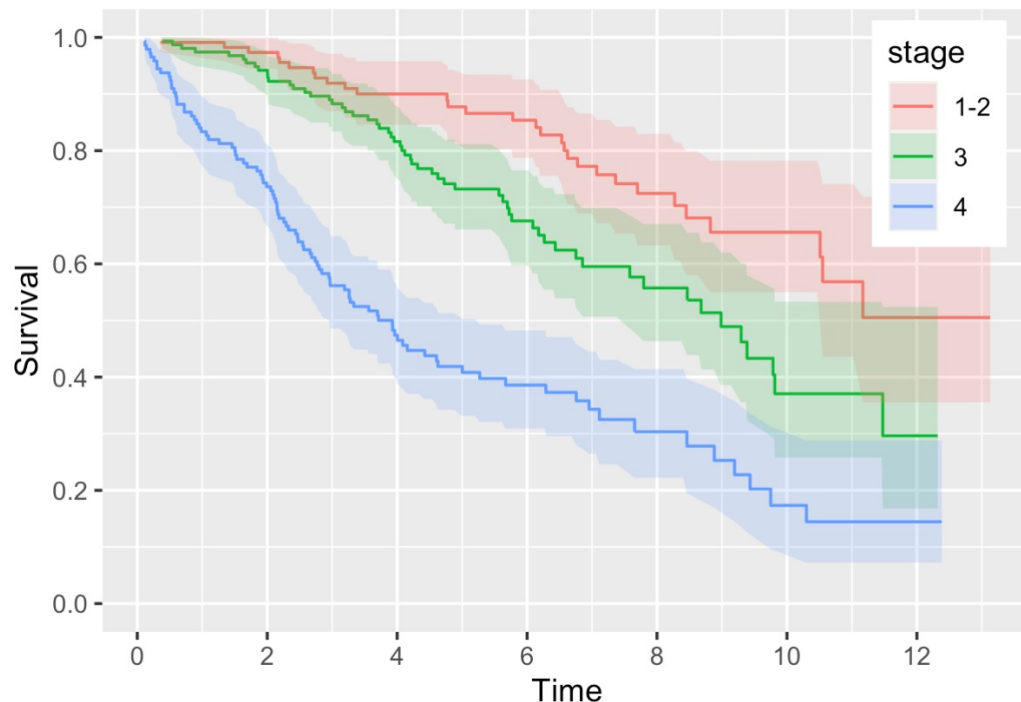
- Pros:
 - We don't need to model $h_0(t)$
 - Very efficient compared to fully parametric alternatives
- Cons:
 - Hinges on the proportional hazards assumption
 - No time-varying effects

Example: Primary Biliary Cholangitis & Death

- PBC: Autoimmune disease destroying the bile ducts in the liver
- Causes fibrosis (scarring) \Rightarrow Cirrhosis \Rightarrow Liver failure \Rightarrow Death
- *What is your prognosis based on your current fibrosis stage (1-4)?*
- 412 PBC patients from the Mayo clinic
- Stage 1+2: 113 Stage 3: 155 Stage 4 (cirrhosis): 144
- 182 died or underwent liver transplant over up to 12 years (median 5 years)
- `data("pbc", package="survival")` in R

Example: Primary Biliary Cholangitis & Death (2)

- Kaplan-Meier curves:
- Can do log-rank test, but...
 - Age is potential confounder!
 - risk of death
 - fibrosis progression (time)
 - Overall quantification of the difference
- Cox regression let's us do all this **and** test for differences between stages



Example: Primary Biliary Cholangitis & Death (3)

- Cox model: $\ln(HR) = \beta_1[age] + \beta_2[sex(f)] + \beta_3[stage\ 3] + \beta_4[stage\ 4]$

Estimates:

Hazard Ratios:

Significance tests:

Example: Primary Biliary Cholangitis & Death (3)

- Cox model: $\ln(HR) = \beta_1[age] + \beta_2[sex(f)] + \beta_3 \cdot 0 + \beta_4 \cdot 0$

Stage 1-2:

Estimates:

Hazard Ratios:

Significance tests:

Example: Primary Biliary Cholangitis & Death (3)

- Cox model: $\ln(HR) = \beta_1[age] + \beta_2[sex(f)] + \beta_3 \cdot 1 + \beta_4 \cdot 0$

Stage 3:

Estimates:

Hazard Ratios:

Significance tests:

Example: Primary Biliary Cholangitis & Death (3)

- Cox model: $\ln(HR) = \beta_1[age] + \beta_2[sex(f)] + \beta_3 \cdot 0 + \beta_4 \cdot 1$

Stage 4:

Estimates:

Hazard Ratios:

Significance tests:

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Estimates:

	coef	exp(coef)
age	0.011160	1.011222
sexf	-0.276887	0.758140
stage3	0.588064	1.800500
stage4	1.463718	4.321997

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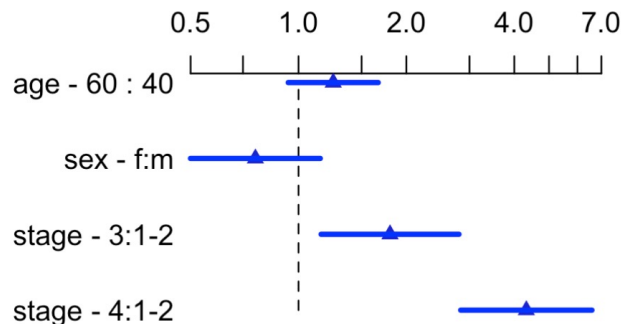
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Significance tests:

$$HR\ age\ 60\ vs\ 40: e^{0.011 \cdot (60-40)} = 1.25$$

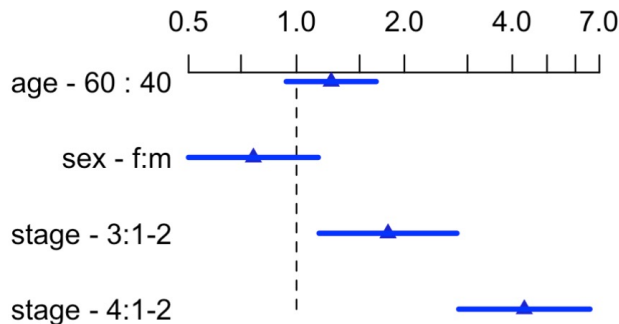
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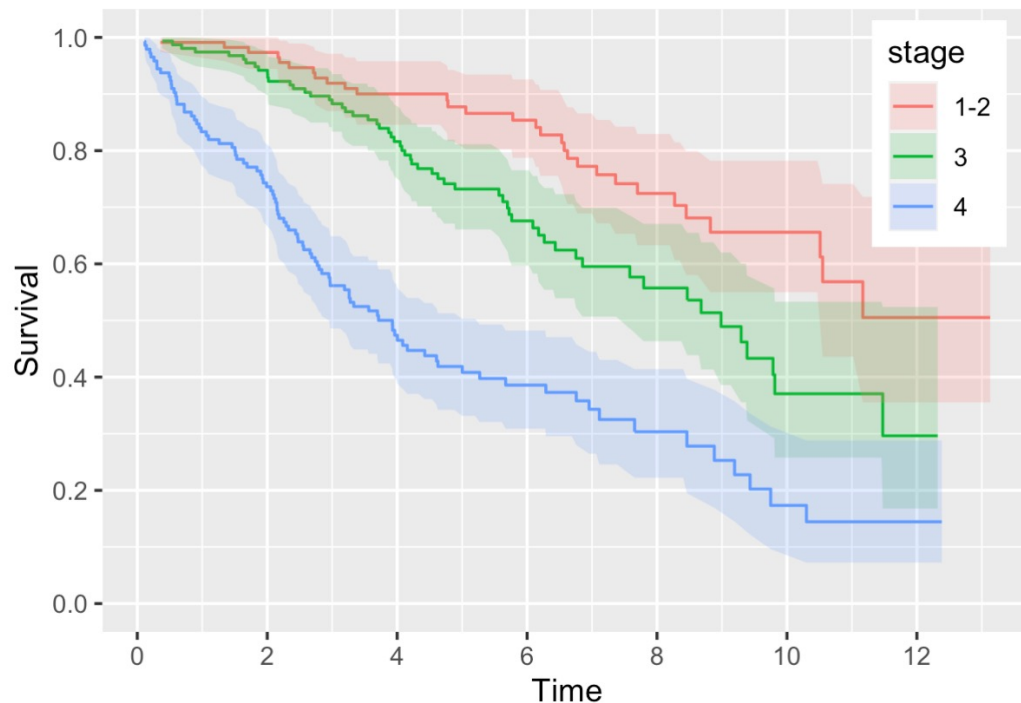
Significance tests:

Factor	Chi-Square	d.f.	P
age	2.29	1	0.1301
sex	1.70	1	0.1929
stage	55.57	2	<.0001
TOTAL	69.00	4	<.0001

$$HR\ age\ 60\ vs\ 40: e^{0.011 \cdot (60-40)} = 1.25$$

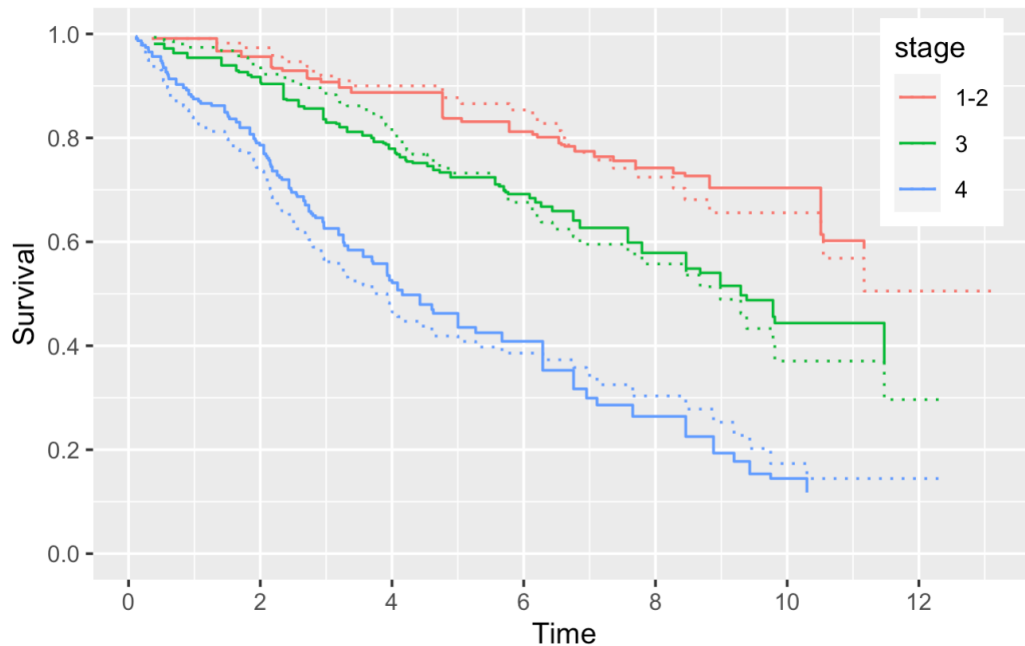
Checking the Proportional Hazards Assumption

- Do the curves look like they have proportional hazards?



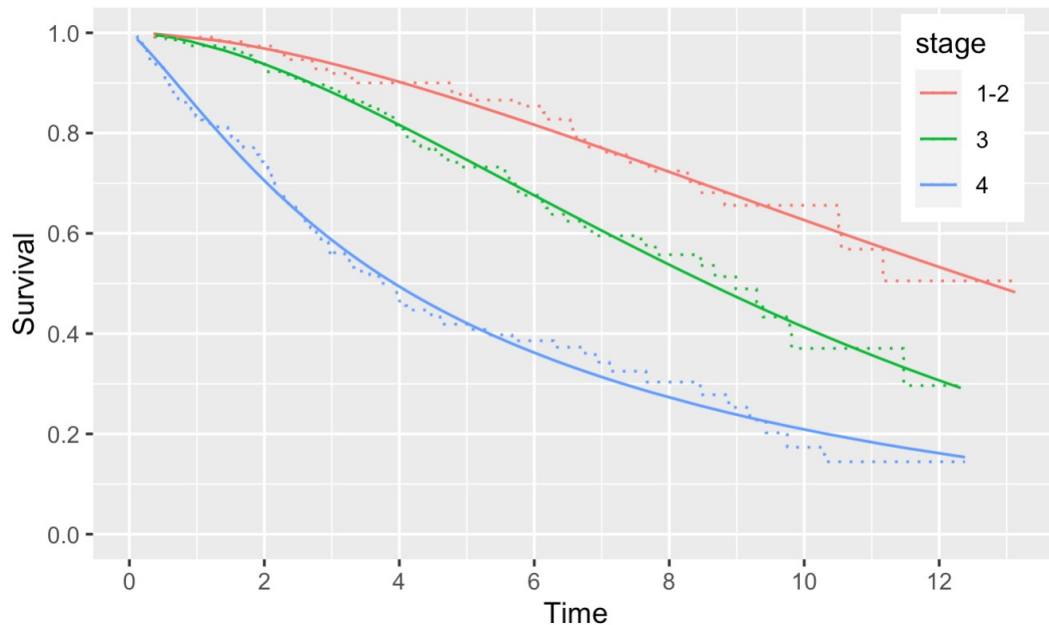
Checking the Proportional Hazards Assumption

- **Do the curves look like they have proportional hazards?**
- If they did, according to the Cox model:
- Important to check!
- What to do?
 - Flexible parametric survival model (advanced)



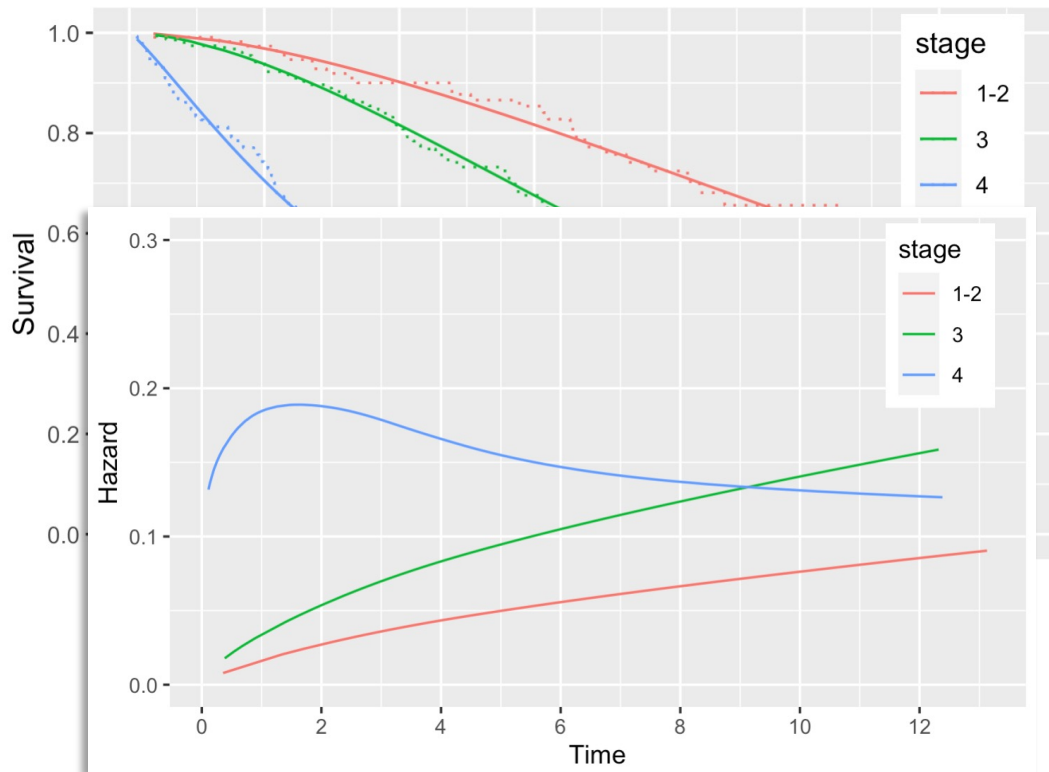
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Summary

- The most important concepts of survival analysis
 1. The type of data we use:
 - Time-to-event
 2. The survival function & the cumulative incidence function:
 - Summary statistics
 3. The hazard rate function
 - The mechanism behind the survival
 4. The Cox proportional hazards regression model
 - All the regression goodness!



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Institutet**