



GILLINGS SCHOOL OF
GLOBAL PUBLIC HEALTH

An Introduction to Censored Covariates

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STA779: Applied Survival Analysis (Spring 2023)

February 28, 2023

REVIEW: Censoring

Patient #

1. _____

2. _____

3. _____

4. _____

5. _____

Years in Study

REVIEW: Censoring

Patient #

1.



= event

2.



3.



4.

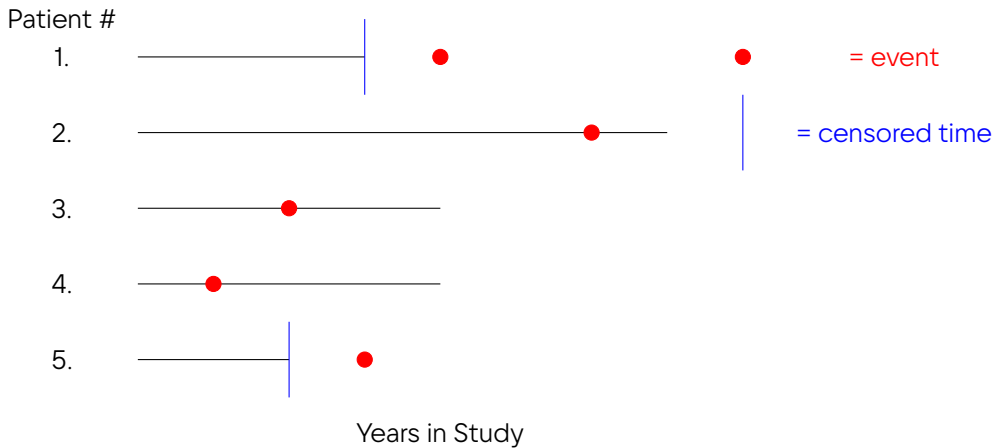


5.

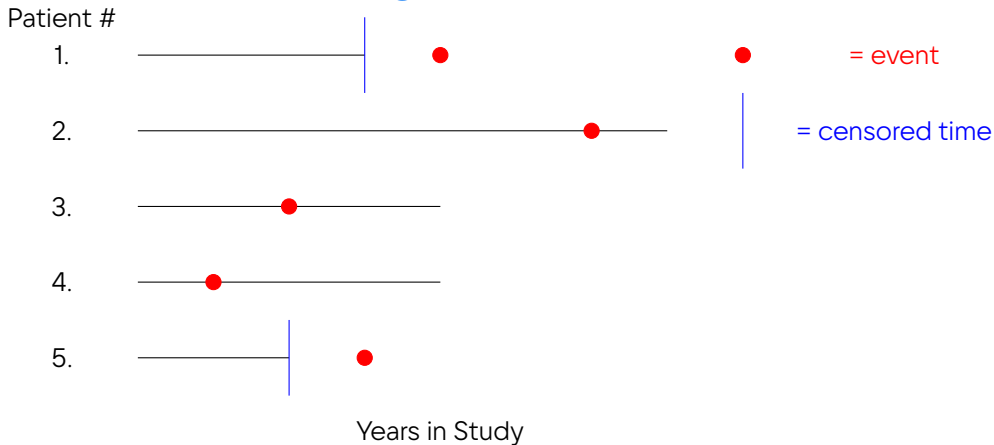


Years in Study

REVIEW: Censoring



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What kind of censoring is this?: *Right?, Left?, Interval?*

REVIEW: Censoring

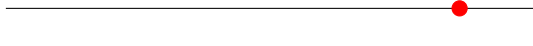
Patient #

1.



= event

2.



= censored time

3.



4.



5.



Years in Study

What kind of Type I censoring could this be?: *Fixed? Progressive? Generalized?*

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- What you will *primarily* learn about in this class (except today)
- The whole idea around survival analysis, where the *outcome* you are interested in is some 'time until event' (or some function of that time)
- It is often the case that you will adjust for some other factors, or covariates, that could affect this outcome
- or maybe you want to see which factors determine survival of this event

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- What if I want to predict height based on someone's weight?

How do you determine where to put variables in a model?

- What if I want to measure the linear association between height and weight?
- This depends heavily on your **research question and analysis goal**

Censored Covariates

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- The patient's diagnosis time here, is a covariate, based on our research question
- AND it is right-censored, since not all patients will be diagnosed during a Huntington disease observational study
- Another example: What if I want to measure the impact of income on college matriculation but income is top-coded, meaning anyone with an income over a certain number, let's say \$250K, is simply labeled as \geq \$250K.

Types of censored covariates

- **Limit of Detection Censoring**

- Also called top-coded or bottom-coded, like the income example
- A lot of biological markers often have a lower limit of detection, meaning the device to measure the factor cannot pick up levels below a certain threshold
- Often, observations have the same limit of detection, but they can also have different limits
- These limits are known a priori, and are considered constants
- Similar to Type I censoring that you discussed last week

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- **Random Censoring**

- Like the Huntington disease example
- Censoring values differ for each observation, and we don't consider these values known a priori
- Instead, we consider the set of these values as a random variable, with it's own distribution

Some Notation

- Let X denote the censored covariate of interest
- Let C denote the censoring value (considering random censoring)
- We observe $W = \min(X, C)$ and $\Delta = I(X \leq C)$
- Let Y be the outcome of interest and Z be the set of fully-observed covariates

Missing Data vs. Censored Data

- Censored Data \subset Missing Data

Missingness and Censoring Mechanisms

Why don't we know X ? *This is the most important question to answer when considering missing data techniques*

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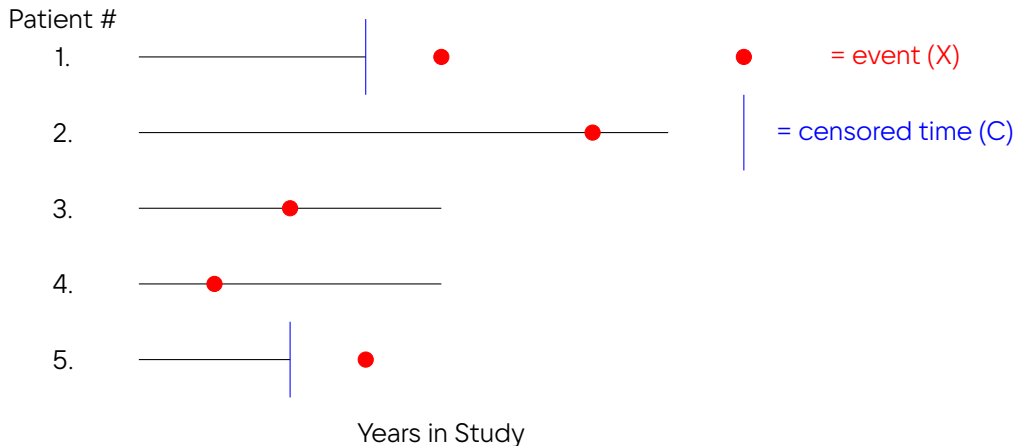
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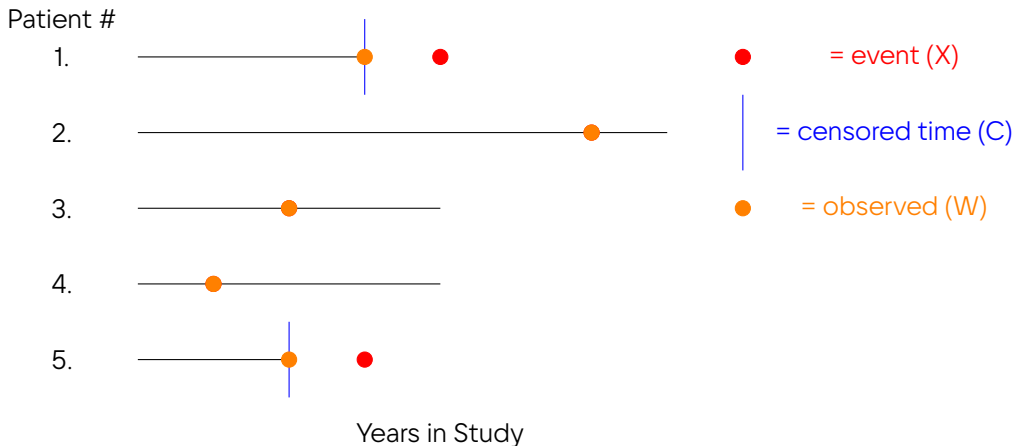
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- Always **MNAR**
- $\Delta \not\perp X$

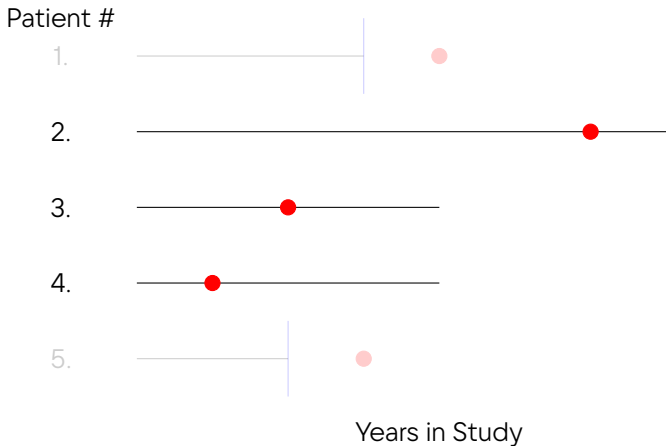
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- i.e. $\Delta \perp\!\!\!\perp Y | X, Z$
- What happens if our COVARIATE is censored?
- What happens if our OUTCOME is censored (i.e., switch the roles of X and Y ?)

Let's Try it!

Click here for the code on my GitHub!

Thank you! Any questions?

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