Survival Analysis Session 3: Incomplete Observations

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Recap: Survival Identities

In survival analysis we consider the random variable "X: Time until event" x = 0.1 weeks, 2.3 weeks...

We express our knowledge about the distribution of *X* in any of these functions. Knowing any single function we can derive all other via the **Survival Identities**.

f(x): Density functionThe relative likelihood of experiencing the event around time *x*.

$$F(x) = \int_0^x f(x) dx$$

F(x): Distribution function *aka Cumulative function*The probability of experiencing the event until time *x*.

$$\mathsf{F}(x) = \mathsf{P}(X {\le} x)$$

$$S(x) = \int_{x} \int f(x) dx \quad h(x) = S'(x)/S(x)$$

S(x): Survival function

The probability of *not* experiencing the event until time *x*.

$$S(x) = P(X>x)$$

h(x): Hazard function

The instantaneous rate of new events at time *x* among those who did not experience the event yet.

$$h(x) = \lim_{h \to 0} P(x \le X < x + h | X \ge x) / h$$

$$H(x) = {}_{0}\int^{x} h(x) dx$$

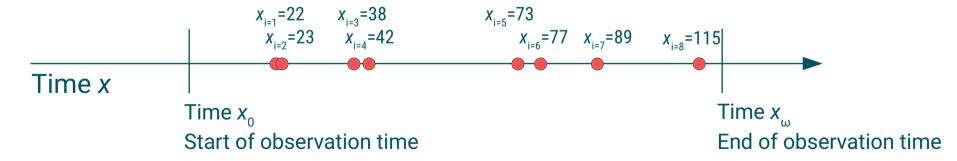
$$H(x) = -\log S(x)$$

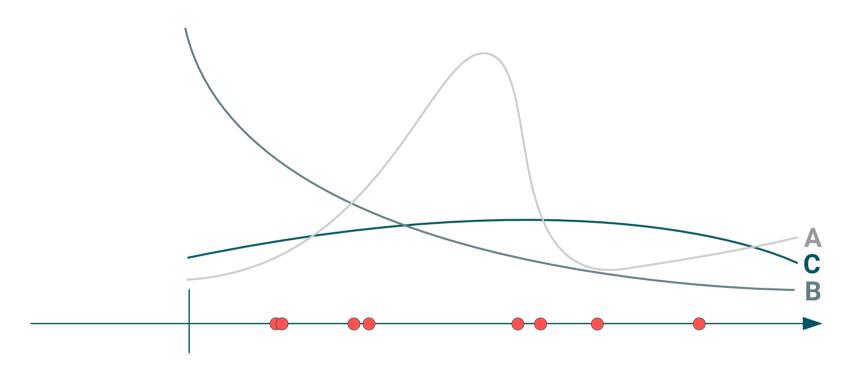


$$S(x) = \exp(-H(x))$$

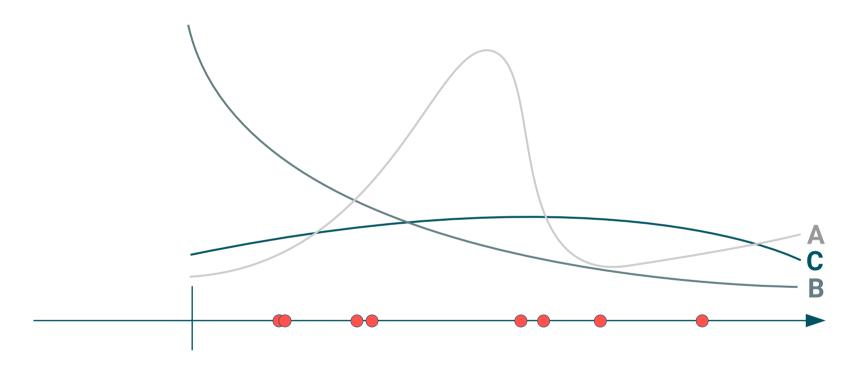
What Does Survival Data Look Like?







Which distribution most likely corresponds to the data?



Which distribution most likely corresponds to the data?

→ Maximum Likelihood Estimation

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Maximum Likelihood Estimation

We fit a model f_{θ} to the data \mathbf{x} by choosing model parameters $\mathbf{\theta}$ which maximize the **likelihood function** L, i.e. which make the observed data most probable.

Product over all observations

Probability density given parameters $oldsymbol{ heta}$

$$L(\theta|\mathbf{x}) = \prod_{i} f_{\theta}(x_{i})$$
 Single observed survival time

In practice we often maximize the **log-likelihood** for convenience:

$$\log L(\boldsymbol{\theta}|\boldsymbol{x}) = \log \prod_{i} f_{\boldsymbol{\theta}}(x_{i}) = \sum_{i} \log f_{\boldsymbol{\theta}}(x_{i})$$

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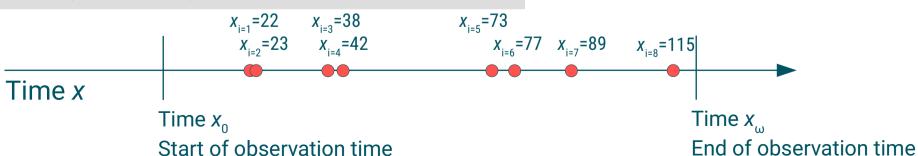
Probability density given parameters $\boldsymbol{\theta}$ $L(\boldsymbol{\theta}|\boldsymbol{x}) = \prod_{i} f_{\boldsymbol{\theta}}(\boldsymbol{x}_{i})$ Single observed survival time

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→ data **x**



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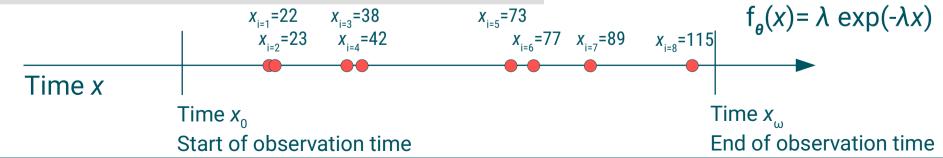
Product over all observations

Probability density given parameters $\boldsymbol{\theta}$ $L(\boldsymbol{\theta}|\boldsymbol{x}) = \prod_{i} f_{\boldsymbol{\theta}}(\boldsymbol{x}_{i})$ Single observed survival time

 $\log L(\boldsymbol{\theta}|\boldsymbol{x}) = \log \prod_{i} f_{\theta}(x_{i}) = \sum_{i} \log f_{\theta}(x_{i})$

You need:

- → data x
- \rightarrow a probability density f_{θ} ("the model") parameterized by...



Which distribution most likely corresponds to the data?

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Maximum Likelihood Estimation

We fit a model f_{θ} to the data x by choosing model parameters θ which maximize the **likelihood function** L, i.e. which make the observed data most probable.

Product over all observations

Probability density given parameters
$$\boldsymbol{\theta}$$

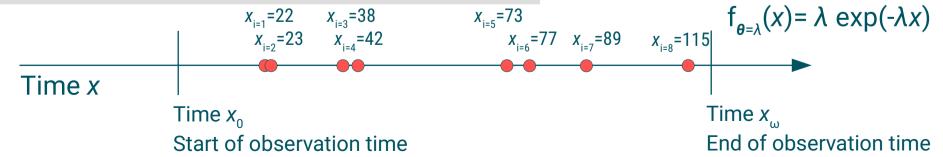
$$L(\boldsymbol{\theta}|\boldsymbol{x}) = \prod_{i} f_{\boldsymbol{\theta}}(\boldsymbol{x}_{i})$$
 Single observed survival time

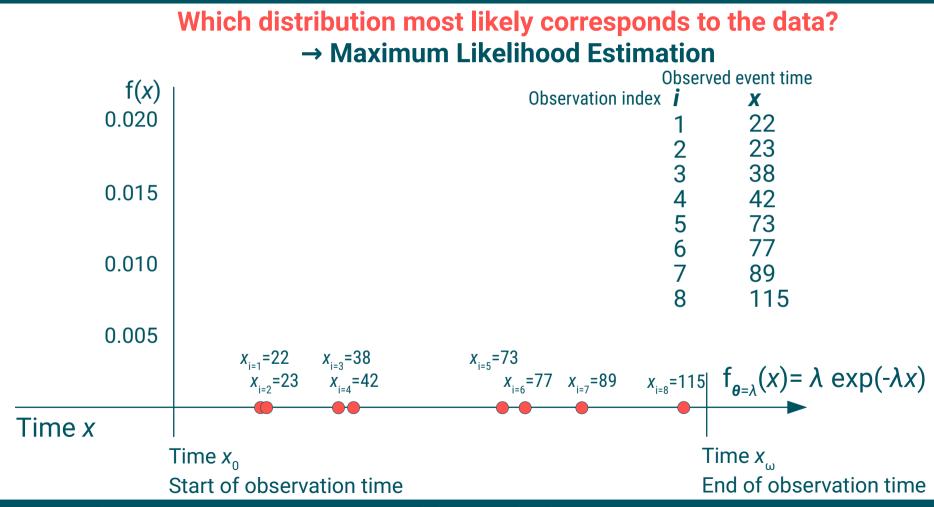
In practice we often maximize the **log-likelihood** for convenience: $| \mathbf{a} \cdot \mathbf{c} \cdot$

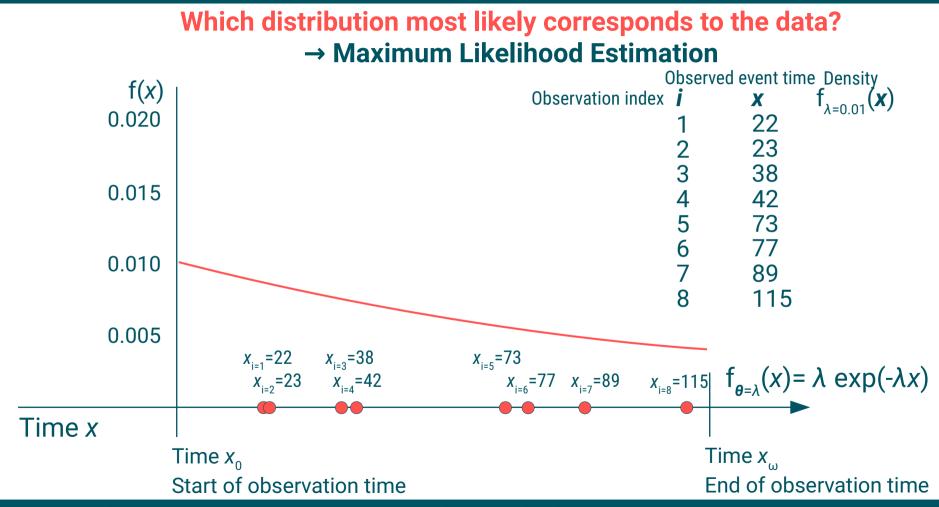
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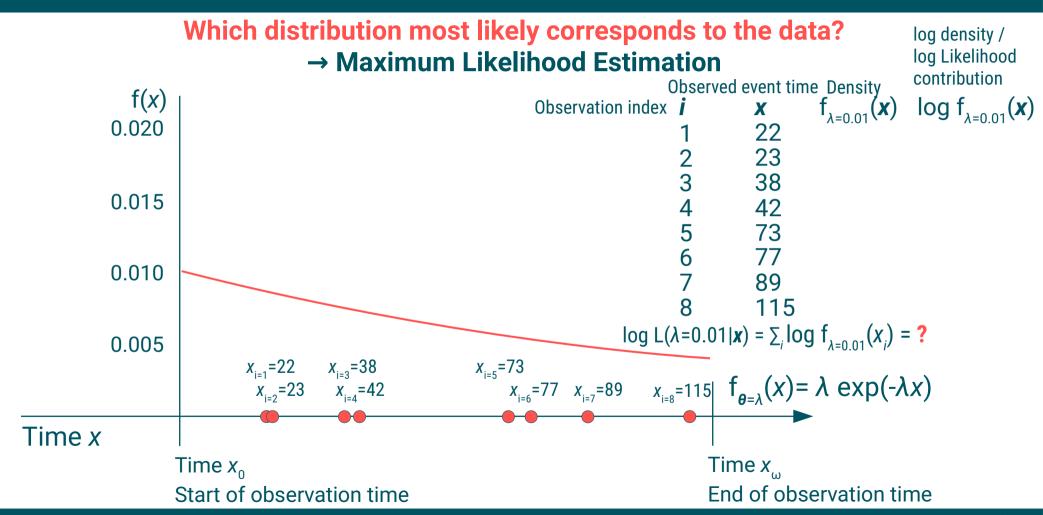
You need:

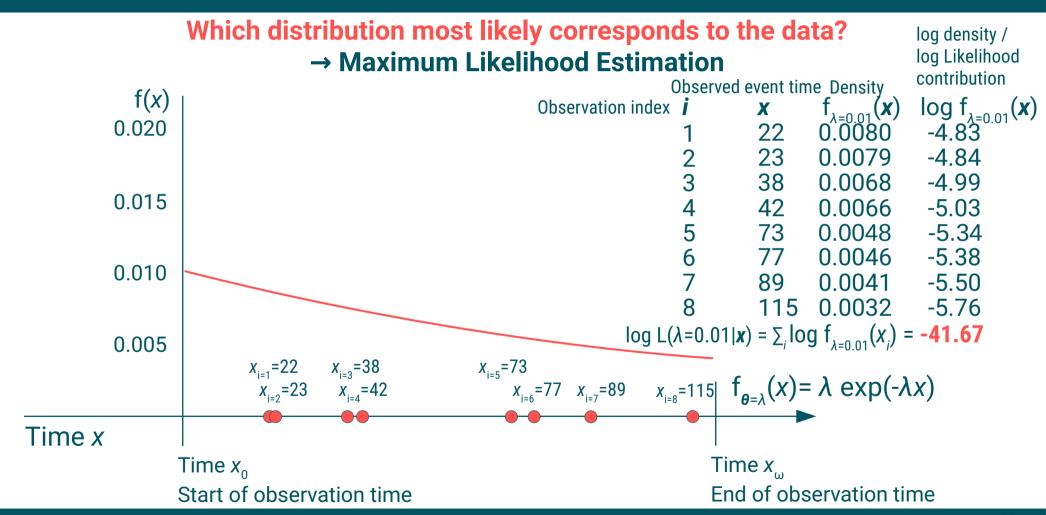
- → data **x**
- \rightarrow a probability density f_{θ} ("the model") parameterized by...
- → ...a set of parameters **0**

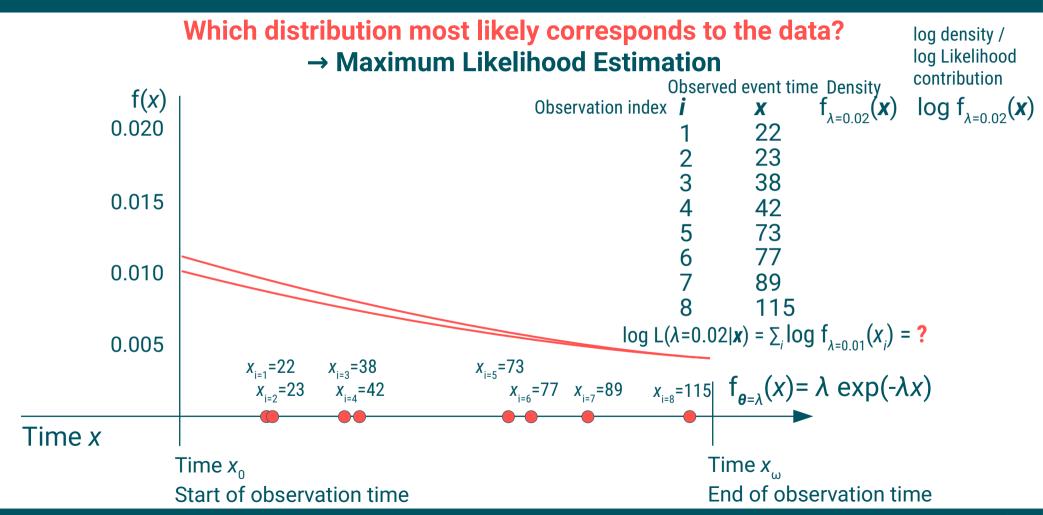


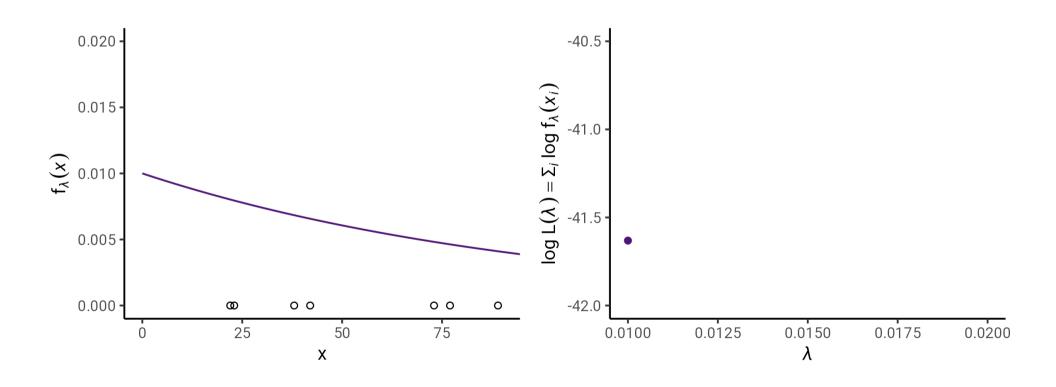


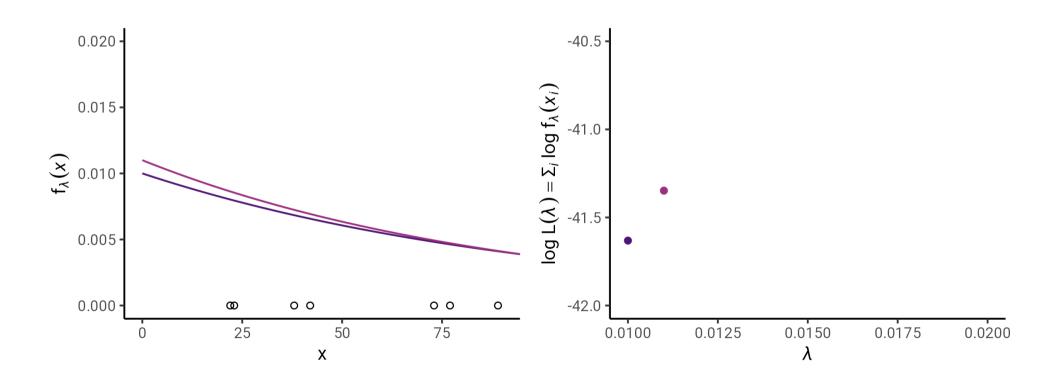


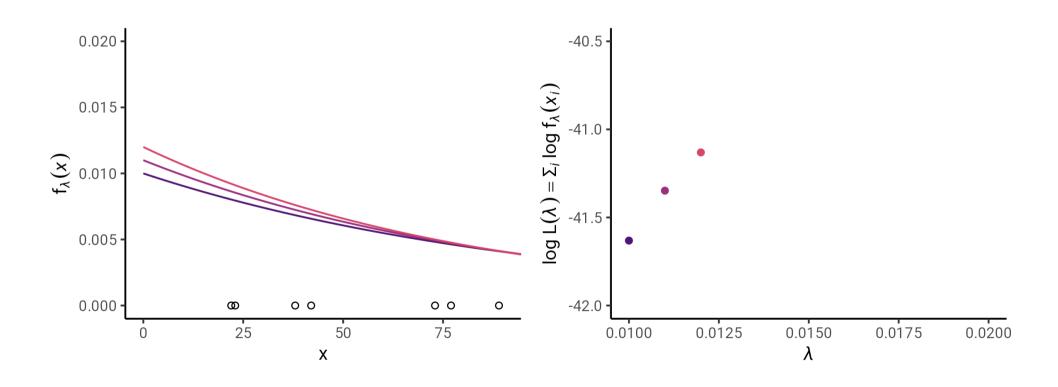


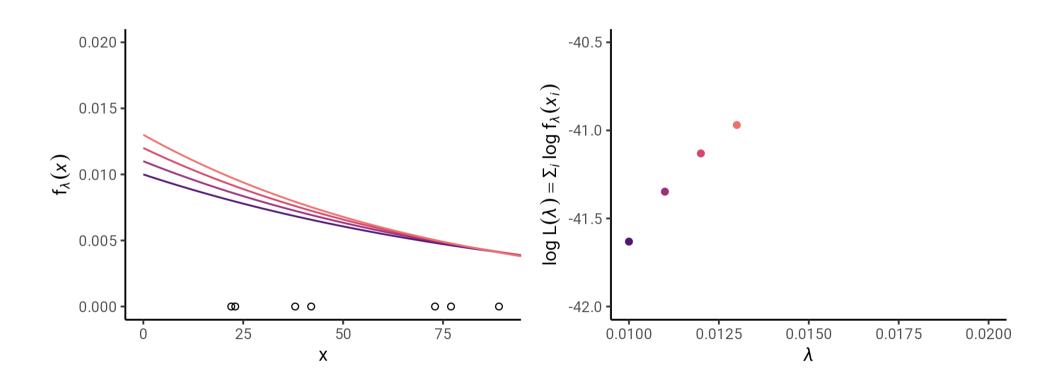


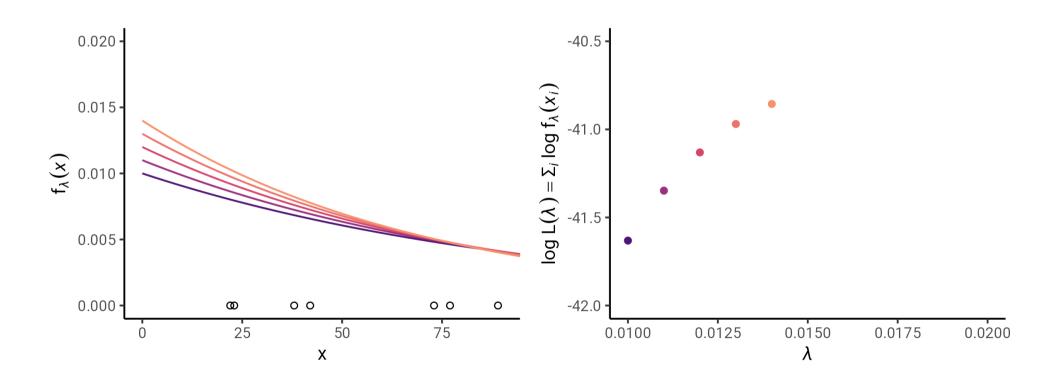


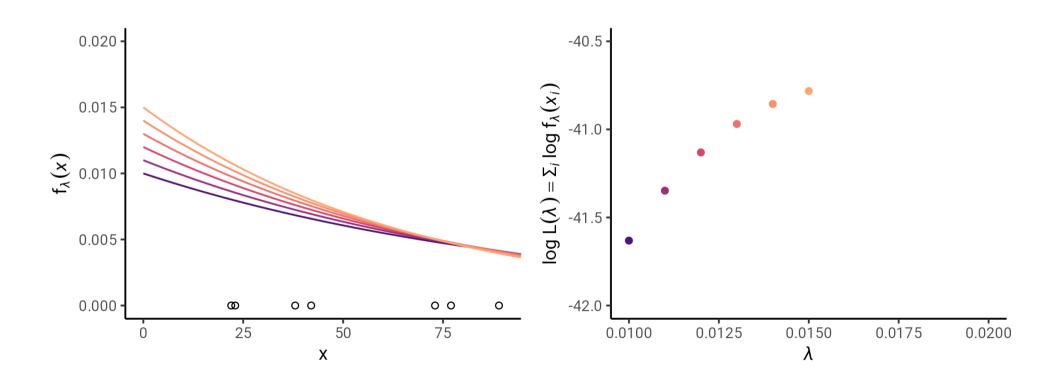


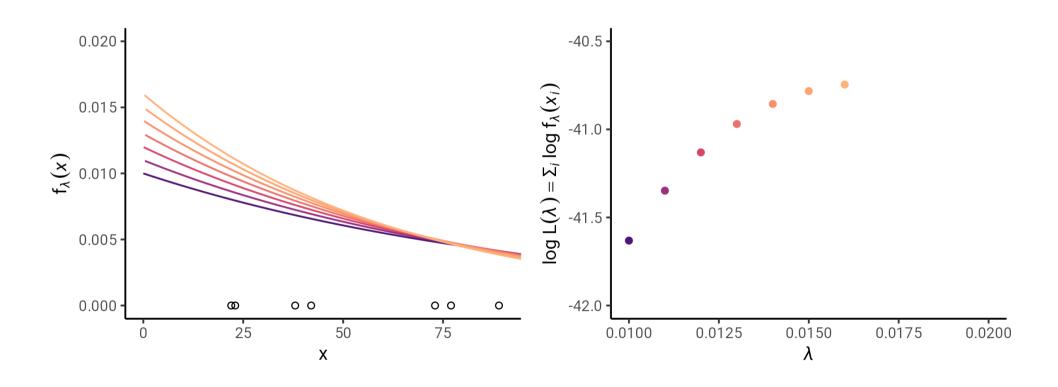


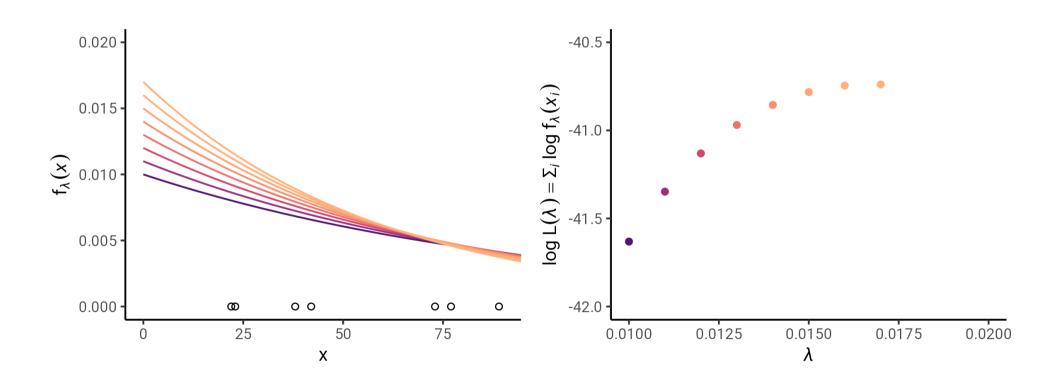


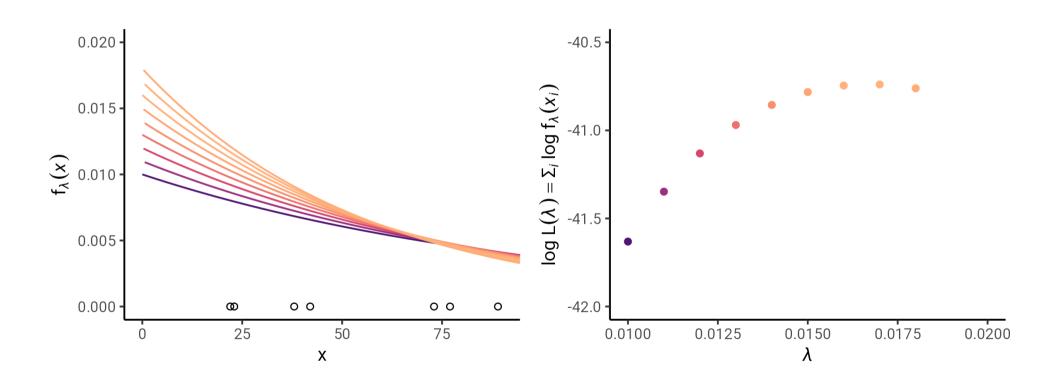


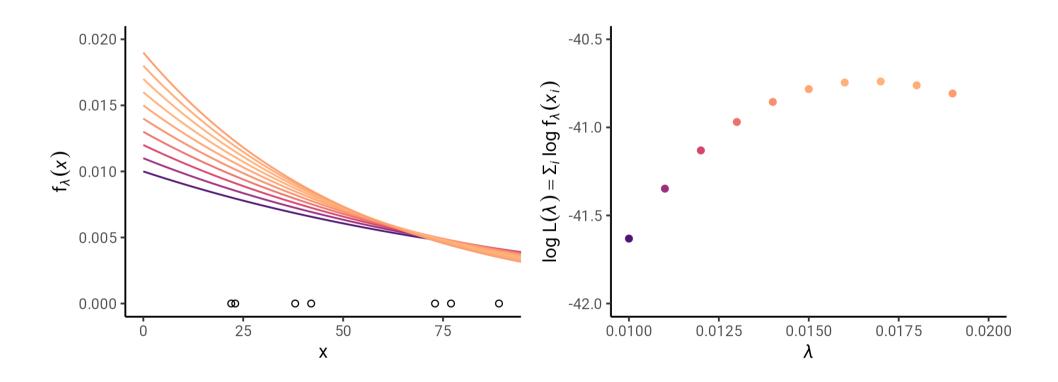






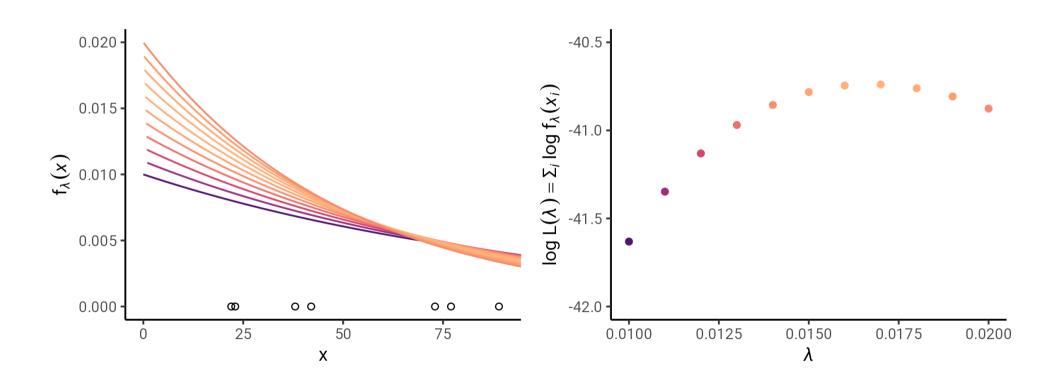






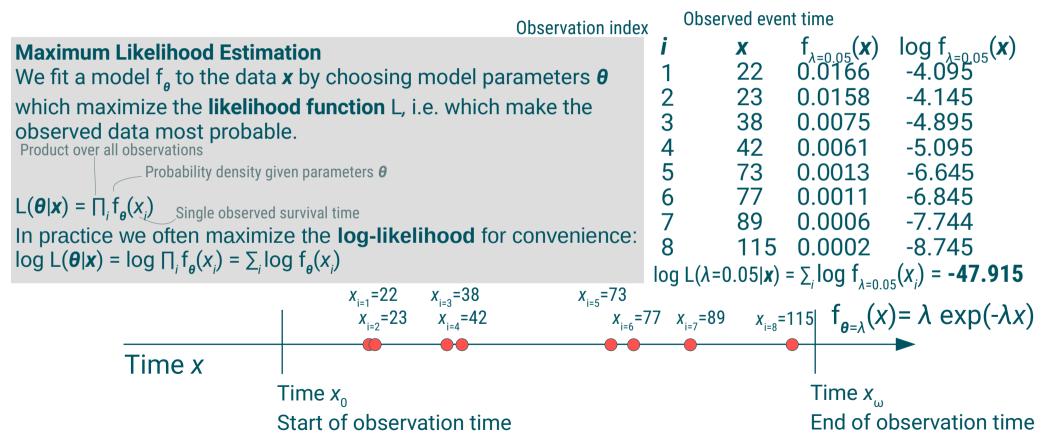
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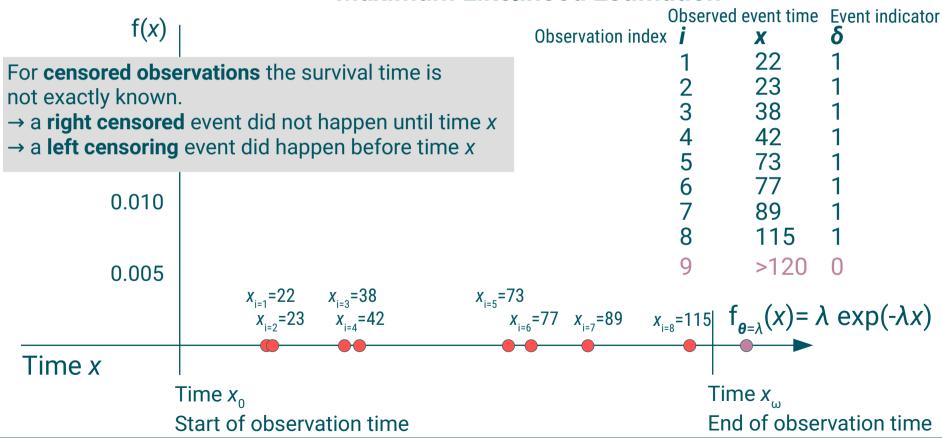
Censoring

But what if some people did not experience the event during the observation time?

→ Censored observations

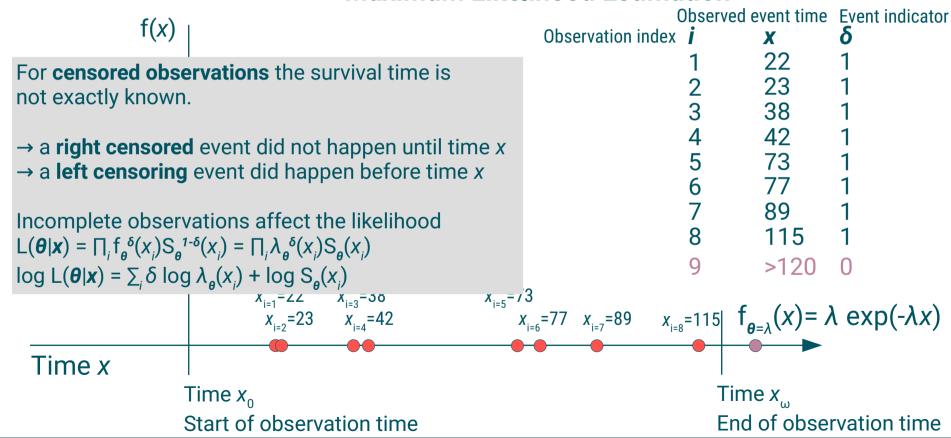
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Materials for this lecture

github.com/jschoeley/survival_analysis-ur-ss22

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