

Fitting the Cox Proportional Hazards model

SURVIVAL ANALYSIS IN PYTHON



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Hazard function and hazard rate

Hazard function $h(t)$: describes the probability that event happens at some time, given survival up to that time.

Hazard rate: the instantaneous rate of event occurring

$$h(t) = -\frac{d}{dt}\log S(t)$$

The **hazard function** $h(t)$ and the **survival function** $S(t)$ can be derived from each other.

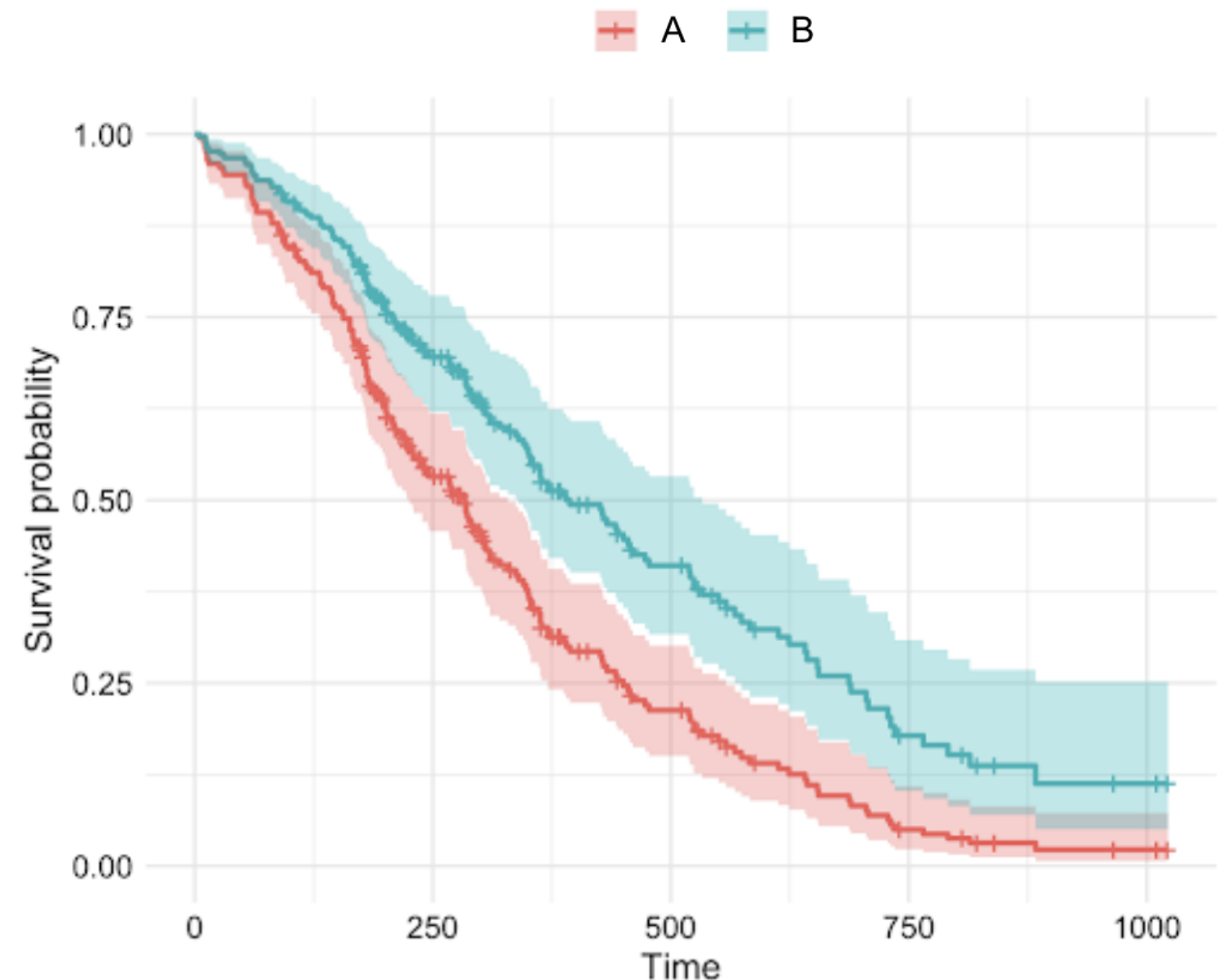
The proportional hazards assumption

The proportional hazards assumption: all individuals' hazards are proportional to one another.

In the case of individual A and individual B :

$$h_A(t) = ch_B(t)$$

1. There is a **baseline hazard function** and other hazards are specified with **scaling factors**.
2. The **relative survival impact** associated with **a variable** does not change with time (time-invariant).



The Cox Proportional Hazards model

Based on the proportional hazards assumption:

$$h(t|x) = b_0(t) \exp \left(\sum_{i=1}^n b_i (x_i - \bar{x}_i) \right)$$

$b_0(t)$: population-level baseline hazard function that changes with time.

$\exp \left(\sum_{i=1}^n b_i (x_i - \bar{x}_i) \right)$: the linear relationship between covariates and the log of hazard, does NOT change with time.

- The Cox Proportional Hazards (Cox PH) model is a **regression model** that regresses **covariates** on time-to-event/duration.

Data requirement for Cox PH model

- **Durations:** the lifetime/duration of the individuals.
- **Events:** whether the event has been observed (1=Yes, 0=No, censored).
 - If not supplied, the model assumes no subjects are censored.
- **Covariates:** continuous or one-hot encoded categorical variables for the regression.

Fitting the Cox PH model

1. Import and instantiate the `CoxPHFitter` class

```
from lifelines import CoxPHFitter  
coxph = CoxPHFitter()
```

2. Call `.fit()` to fit the estimator to the data

```
coxph.fit(df, duration_col, event_col)
```

3. Access other properties to check model summary, covariate, coefficients, predict, plot, etc.

```
coxph.summary()  
coxph.predict()
```

Example Cox PH model

- DataFrame: `mortgage_df`
- Covariates:
 - `house`
 - `principal`
 - `interest`
 - `property_tax`
 - `credit_score`
- Other columns: `duration` , `paid_off`

```
from lifelines import CoxPHFitter
coxph = CoxPHFitter()
coxph.fit(df=mortgage_df,
          duration_col="duration",
          event_col="paid_off")
```

Custom model

Filter the `DataFrame` :

```
new_df = mortgage_df.loc[:,  
    mortgage_df.columns!="house"]  
coxph.fit(df=new_df,  
    duration_col="duration",  
    event_col="paid_off")
```

Use the `formula` parameter:

```
coxph.fit(df=mortgage_df,  
    duration_col="duration",  
    event_col="paid_off",  
    formula="principal + interest  
    + property_tax + credit_score")
```

- More convenient and clearer, but doesn't scale to large number of covariates.

Interpret coefficients

```
print(coxph.summary)
```

```
<lifelines.CoxPHFitter: fitted with 1808 observations, 340 censored>
```

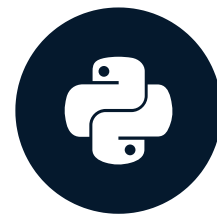
| | | coef | exp(coef) | se(coef) | z | p |
|-----------|--------------|-------|-----------|----------|--------|------|
| covariate | house | -0.38 | 0.68 | 0.19. | -1.98 | 0.05 |
| | principal | -0.06 | 0.94 | 0.02 | -2.61 | 0.01 |
| | interest | 0.31 | 1.37 | 0.31 | 1.02 | 0.31 |
| | property_tax | -0.15 | 0.86 | 0.21 | -0.71 | 0.48 |
| | credit_score | -0.43 | 0.65 | 0.38 | -1.14. | 0.26 |

- **Hazard ratio:** e^{coef}
 - A one unit increase in `interest` from its median value -> the hazards change by the a factor of $e^{0.31} = 1.37$, which is a 37% increase compared to the baseline hazards.

Let's practice!
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Interpreting the Cox PH model

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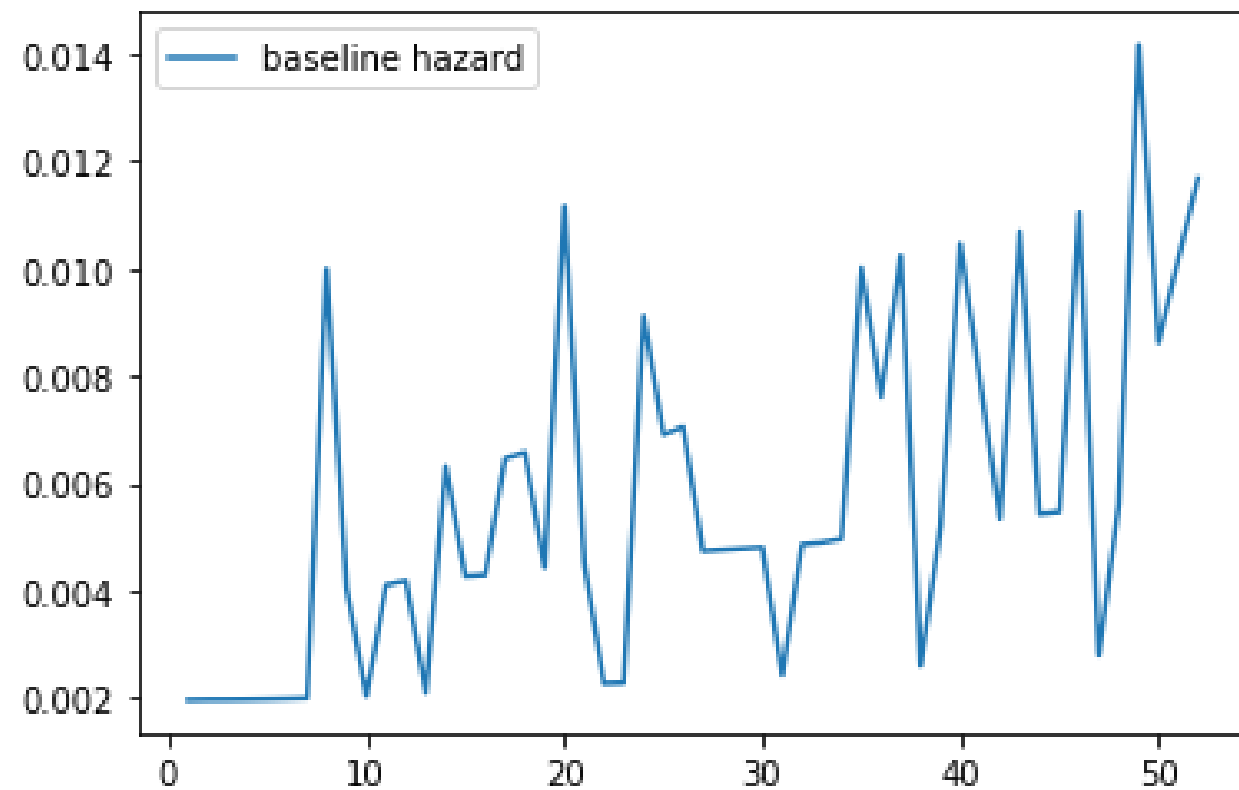
The baseline hazards

- **Hazard ratio:** how much hazard increases or decreases relative to *baseline* hazards.
- **Baseline hazards:** the risk for individuals at the baseline levels of covariates.
 - *Baseline* \neq setting covariates to 0
 - *Baseline* means setting covariates to their averages (median for `lifelines`)

The baseline functions

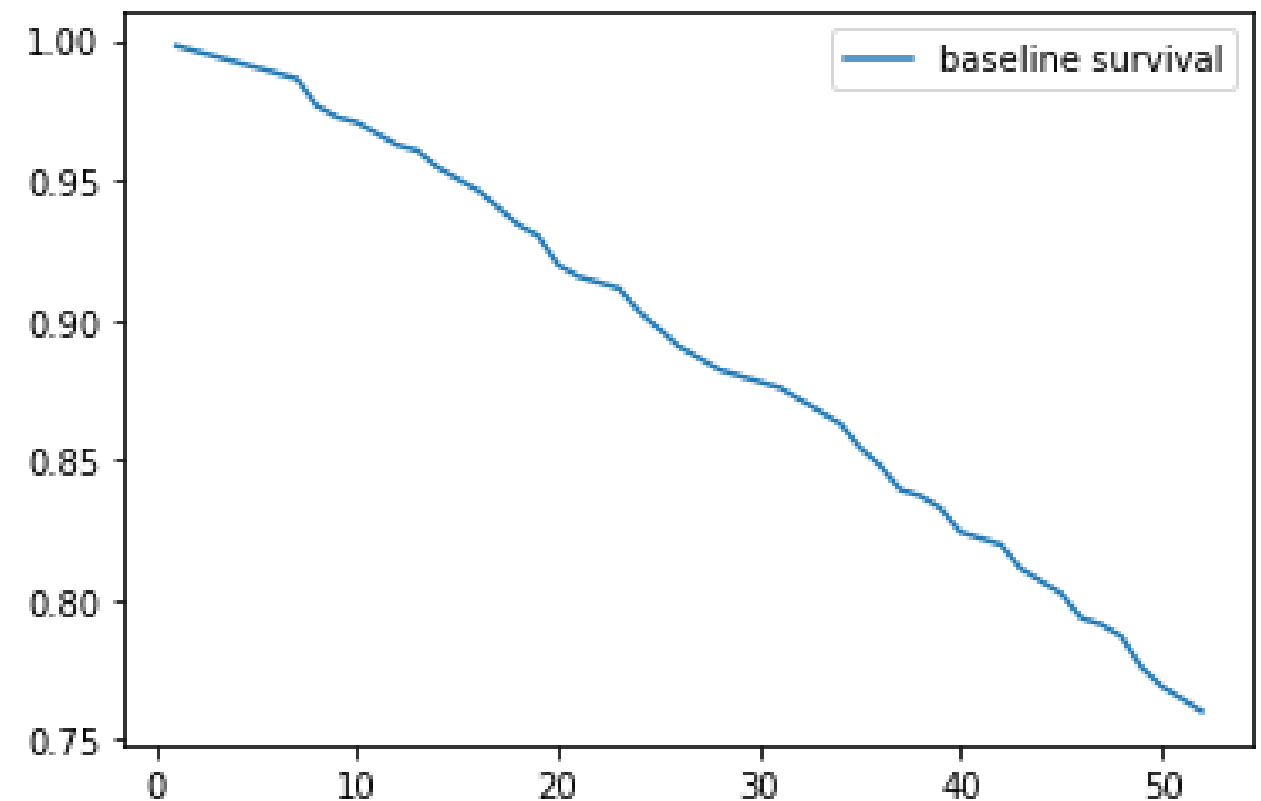
Baseline hazard function

```
model.baseline_hazard_.plot()  
plt.show()
```



Baseline survival function

```
model.baseline_survival_.plot()  
plt.show()
```



Interpret the hazard ratio

- **Hazard ratio:** e^{coef} , how much hazard changes relative to the *average* individual when covariates change.

| | Calculation | Example |
|------------------------------|---------------------|---|
| Coefficient | x | 0.405 |
| Hazard ratio | e^x | $e^{0.405} = 1.5$ |
| Hazards interpretation | $e^x - 1$ | $1.5 - 1 = 0.5$ -> 50% increase in hazards |
| Survival time interpretation | $\frac{1}{e^x} - 1$ | $\frac{1}{1.5} - 1 = 0.67 - 1 = -0.23$ -> 23% decrease in survival time |

Visualize the hazard ratio

```
.plot_partial_effects_on_outcome()
```

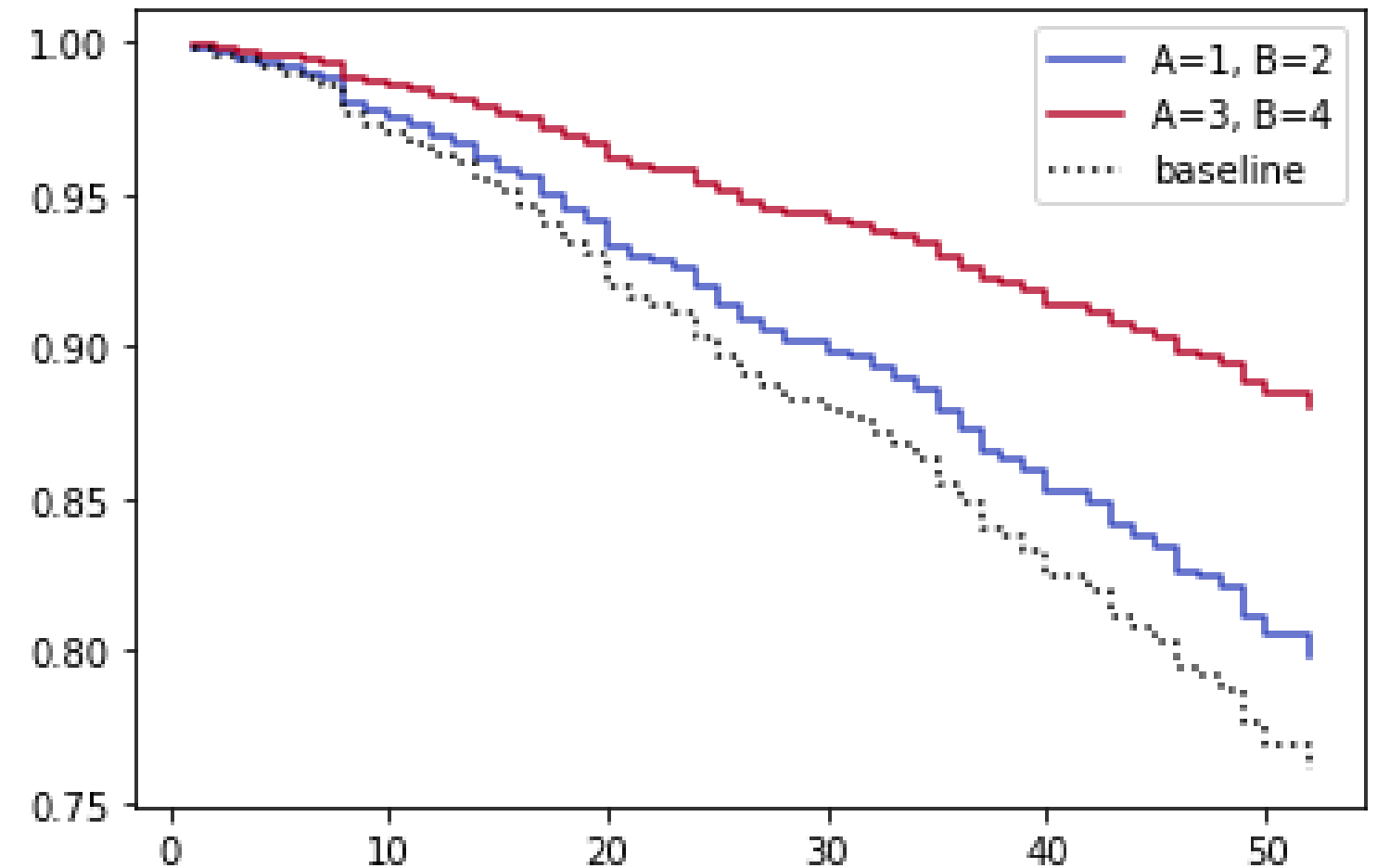
- `covariates` (string or list): name(s) of the covariate(s) in the original dataset that we wish to vary.
 - If there are multiple covariates, pass them in as **a list**.
- `values` (1d or 2d iterable): values we wish the covariate to take on.
 - If there are multiple covariates, pass the values as **pairs/tuples of values**.

Visualize the hazard ratio

The model has covariates **A** , **B** , **C** , and we wish to vary

- **A** over 1, 2
- **B** over 3, 4

```
model.plot_partial_effects_on_outcome(  
    covariates=["A", "B"],  
    values=[[1, 2],  
            [3, 4]]  
)  
plt.show()
```



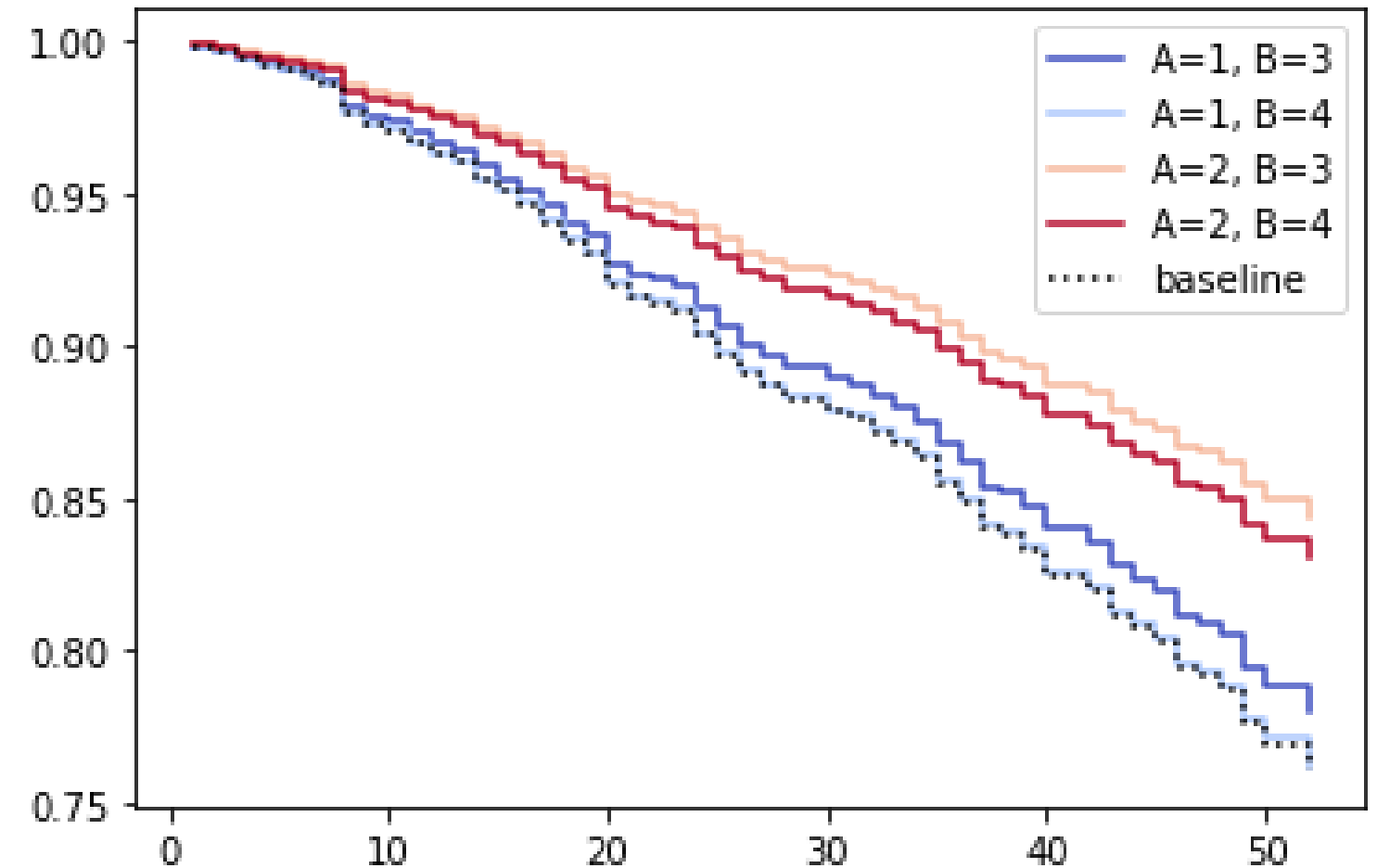
Wrong...

Visualize the hazard ratio

The model has covariates **A** , **B** , **C** , and we wish to vary

- **A** over 1, 2
- **B** over 3, 4

```
model.plot_partial_effects_on_outcome(  
    covariates=["A", "B"],  
    values=[[1, 3],  
            [1, 4],  
            [2, 3],  
            [2, 4]]  
)  
plt.show()
```



Correct!

Let's practice!
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The proportional hazards assumption

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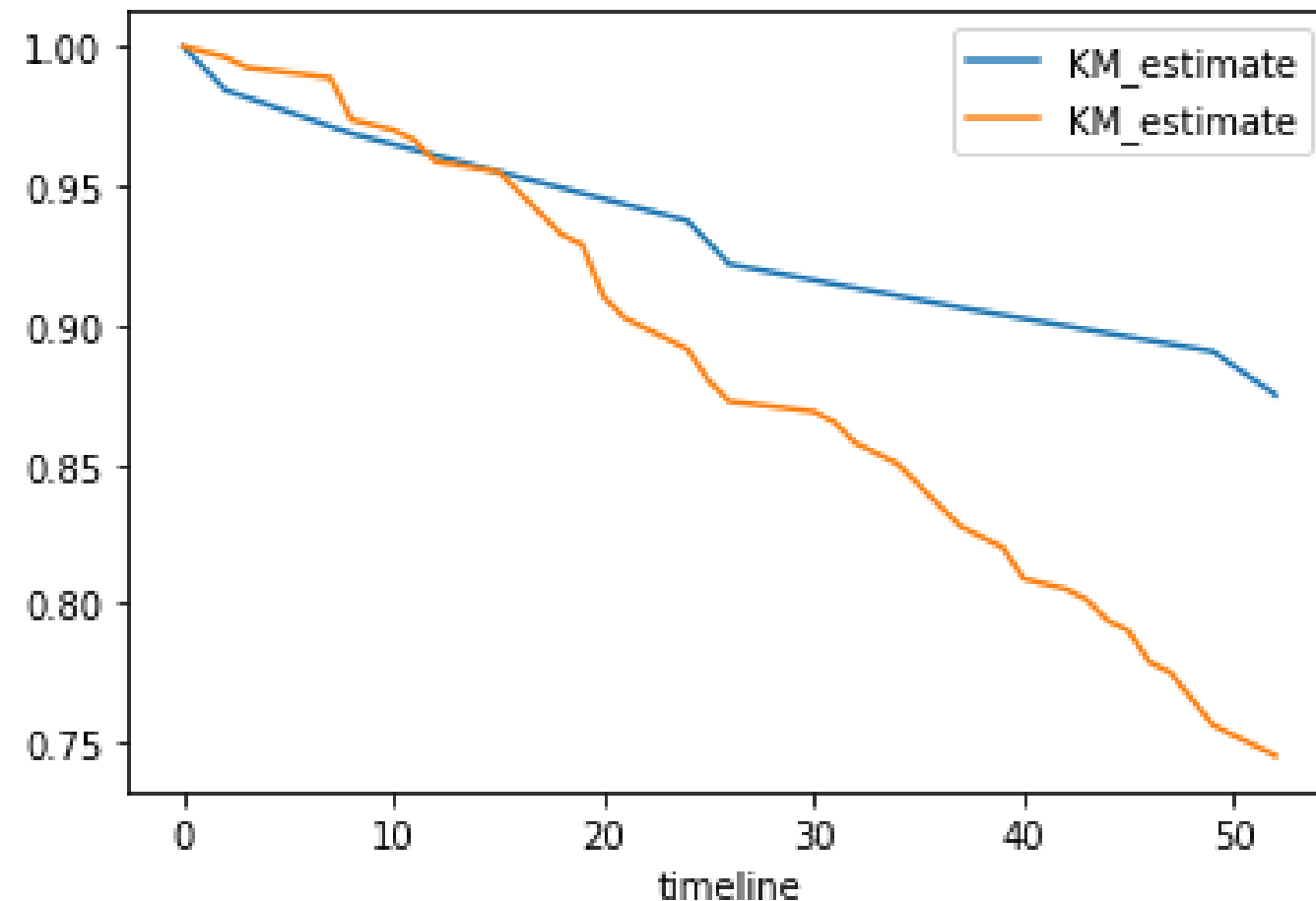
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Use the Kaplan-Meier curves

If the covariate only has a few values, inspect each group's Kaplan-Meier curve.

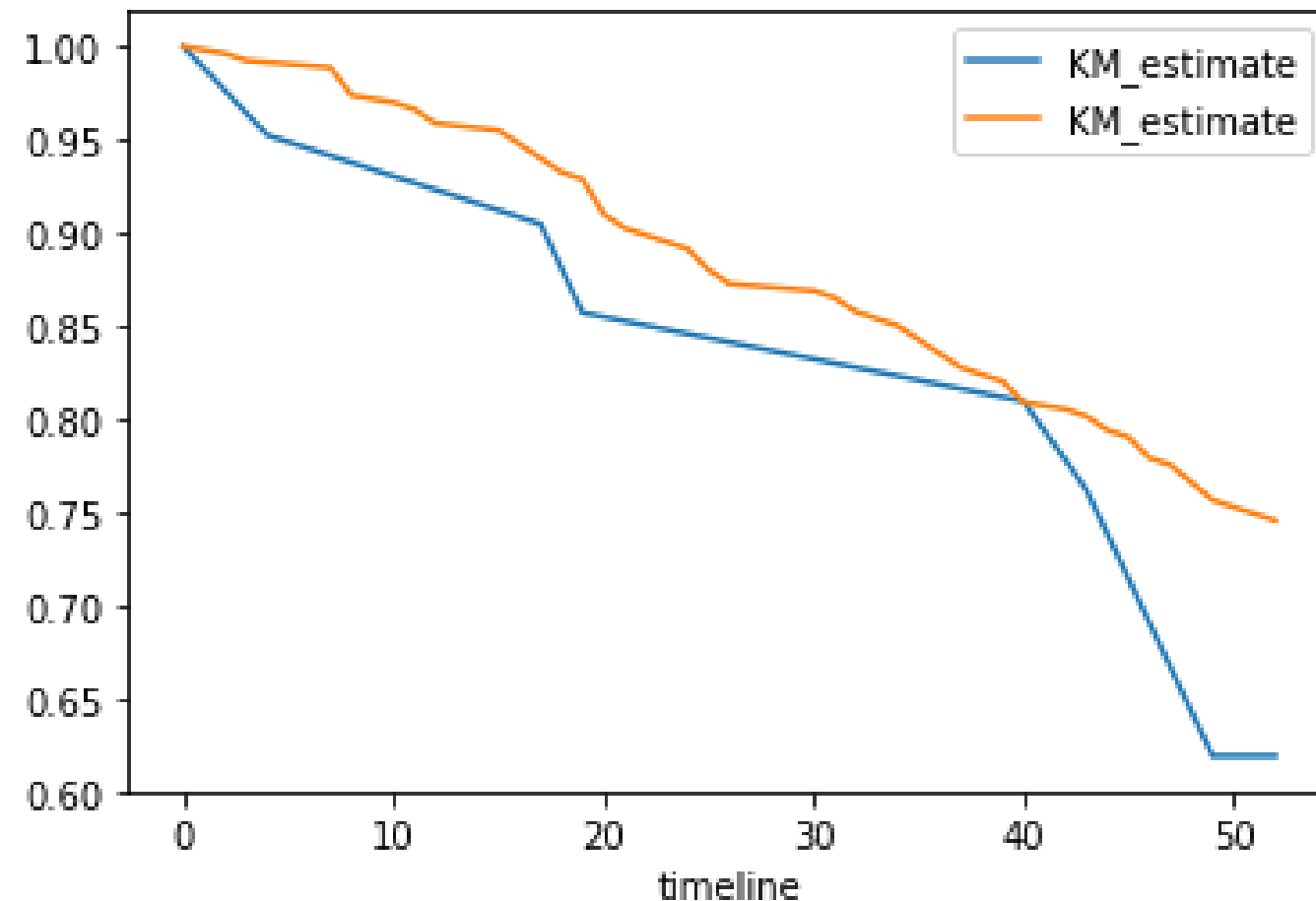
- Curves intersect: **fails** the proportional hazards assumption.



Use the Kaplan-Meier curves

If the covariate only has a few values, inspect each group's Kaplan-Meier curve.

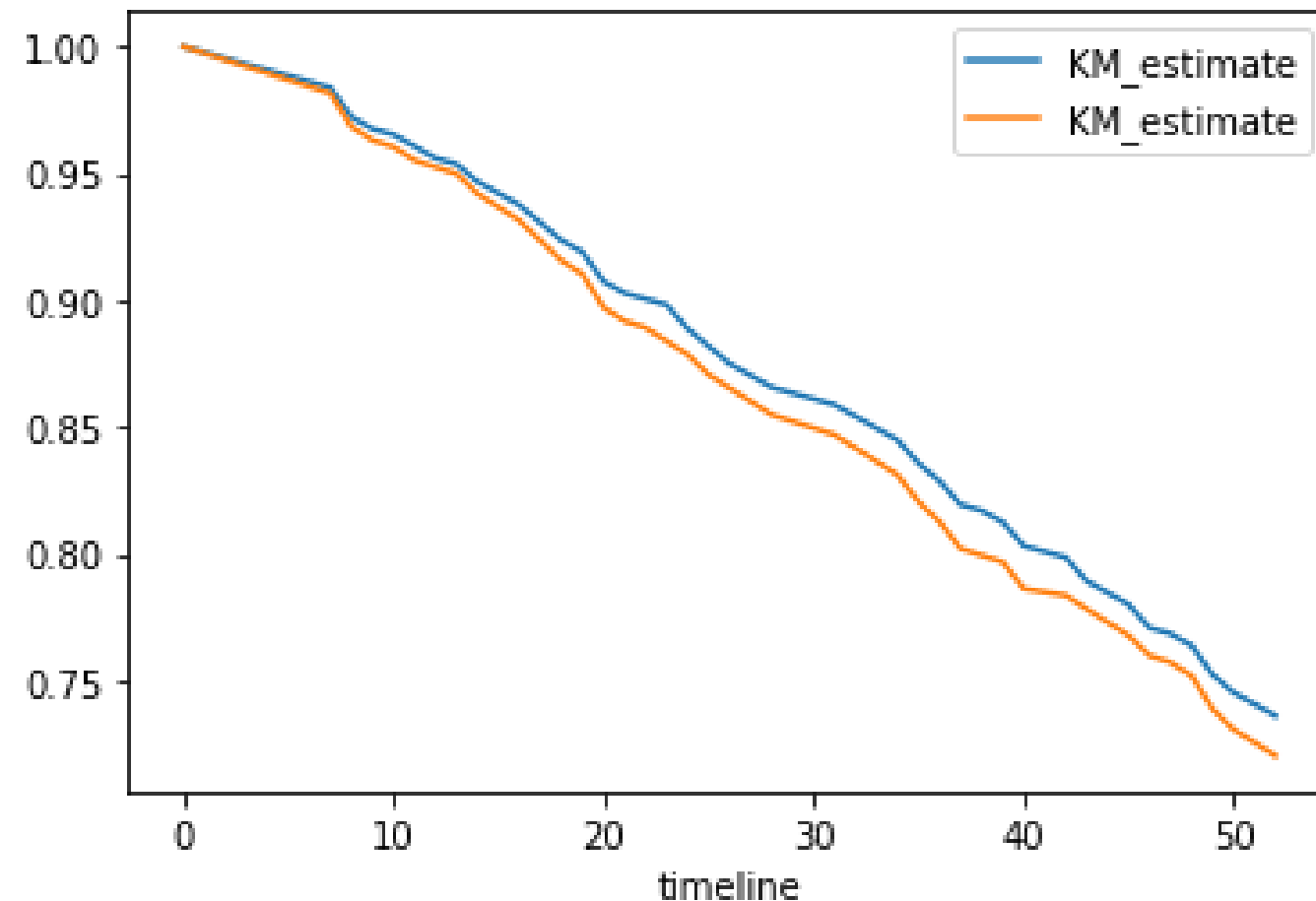
- Curves have different shapes: **fails** the proportional hazards assumption.



Use the Kaplan-Meier curves

If the covariate only has a few values, inspect each group's Kaplan-Meier curve.

- Curves have similar shapes and are parallel: **satisfies** the proportional hazards assumption.



.check_assumptions()

If the covariates are continuous, use the `.check_assumptions()` method.

- Parameters
 - `training_df` : the original DataFrame used in the call to fit the model.
 - `p_value_threshold` : the threshold to use to alert the user of violations (default: 0.01, recommended: 0.05).

.check_assumptions()

```
model.check_assumptions(training_df, p_value_threshold=0.05)
```

```
1. Variable 'A' failed the non-proportional test: p-value is 0.0007.
```

```
Advice 1: ...
```

```
Advice 2: ...
```

```
2. Variable 'B' failed the non-proportional test: p-value is 0.0063.
```

```
Advice 1: ...
```

```
Advice 2: ...
```


When the proportional hazards assumption fails

- Usually, it's a reasonable assumption and violations do not impact model performance significantly.
- If it fails, try other modeling frameworks, such as the Weibull AFT model, and compare their AIC scores.

Let's practice!
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Predicting with the Cox PH model

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Predict median survival times

After calling `.fit()` to fit model to the data:

- `.predict_median()` : predicts the median lifetimes for subjects
 - If the survival curve does not cross 0.5, the median survival time is `inf`.
- Parameters:
 - `X` : the DataFrame to predict with.
 - `conditional_after` : an array or list of values that represent how long subjects have already lived for.

Predict median survival times

```
model.predict_median(X, conditional_after)
```

```
0      inf
1    44.0
2    46.0
3      inf
4    48.0
...
500    inf
```

Predict the survival function

- `.predict_survival_function()` : predicts the survival function for subjects, given their covariates.
- Parameters:
 - `X` : the DataFrame to predict with.
 - `conditional_after` : an array or list of values that represent how long subjects have already lived for.

Predict the survival function

```
model.predict_survival_function(X, conditional_after)
```

| | 0 | 1 | 2 | 3 | 4 | ... | 500 |
|-----|----------|----------|----------|----------|----------|-----|----------|
| 1.0 | 0.997616 | 0.993695 | 0.994083 | 0.999045 | 0.997626 | ... | 0.998865 |
| 2.0 | 0.995230 | 0.987411 | 0.988183 | 0.998089 | 0.995250 | ... | 0.997728 |
| 3.0 | 0.992848 | 0.981162 | 0.982314 | 0.997133 | 0.992878 | ... | 0.996592 |
| 4.0 | 0.990468 | 0.974941 | 0.976468 | 0.996176 | 0.990507 | ... | 0.995455 |
| 5.0 | 0.988085 | 0.968739 | 0.970639 | 0.995216 | 0.986392 | ... | 0.993476 |

Why are survival predictions useful?

- Proactive failure prevention, forecasting models, etc.

Key steps

1. Preprocess the data and one-hot encode any categorical variables.
2. Split data into train and test (common split is 80% train and 20% test).
 - The proportions of censored data should be similar in both sets.
3. Fit the Cox PH model to train.

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Congratulations!

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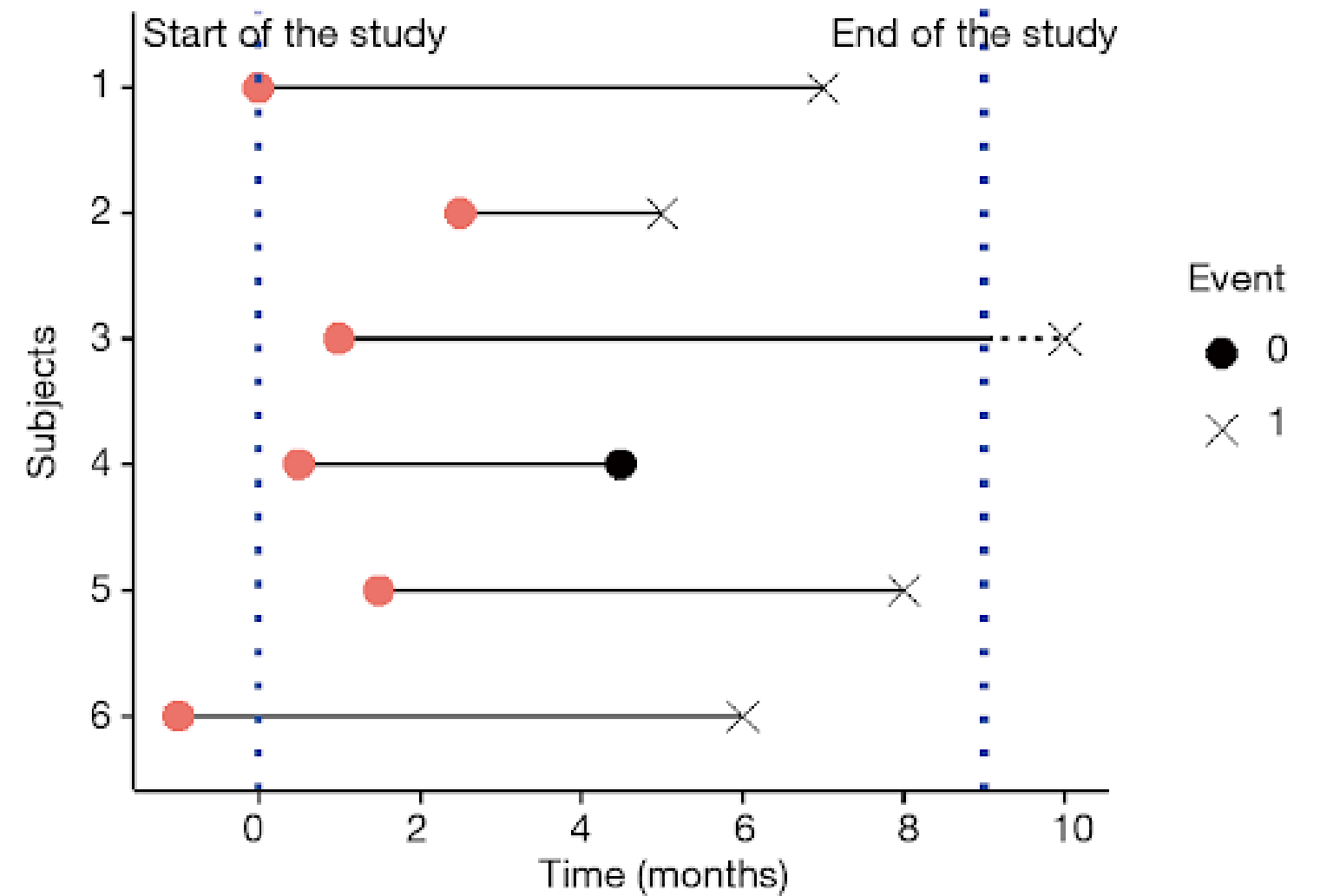
Senior Data Scientist at Ripple

Why survival analysis?

Use cases

- Estimate time to event
- Measure how factors affect time to event

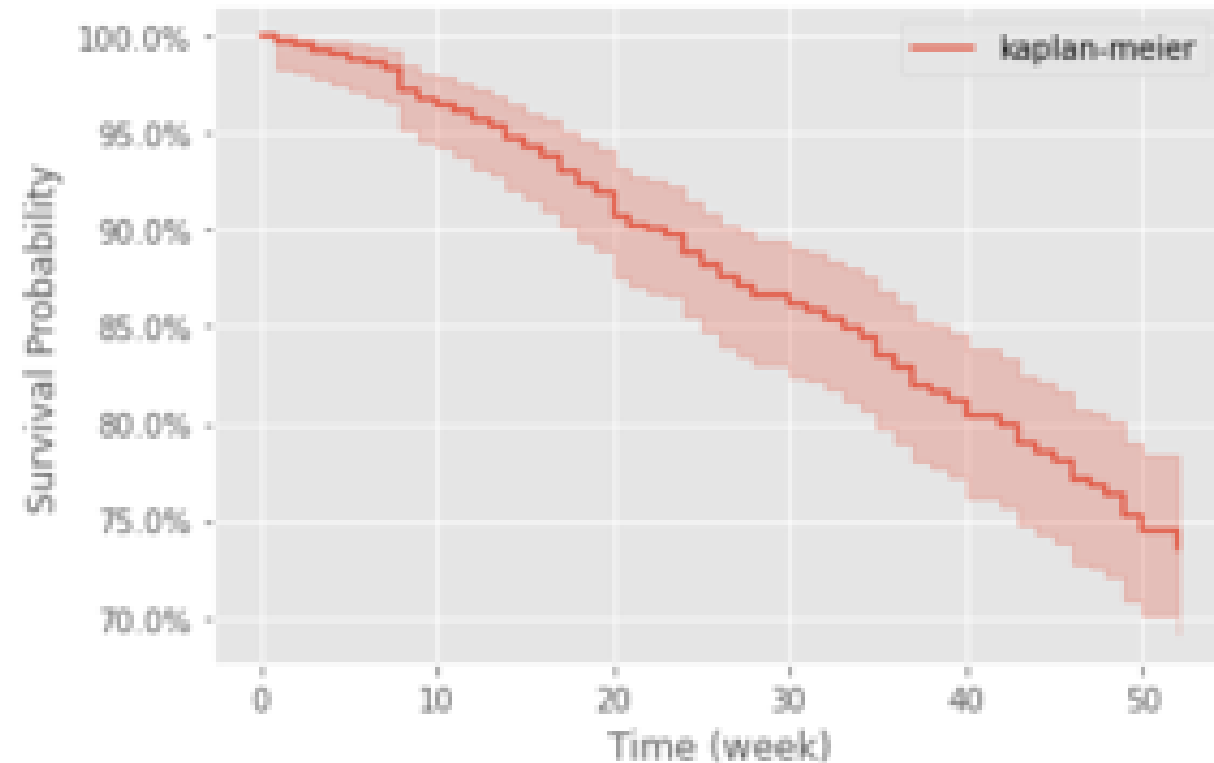
Censoring



Estimate survival curves

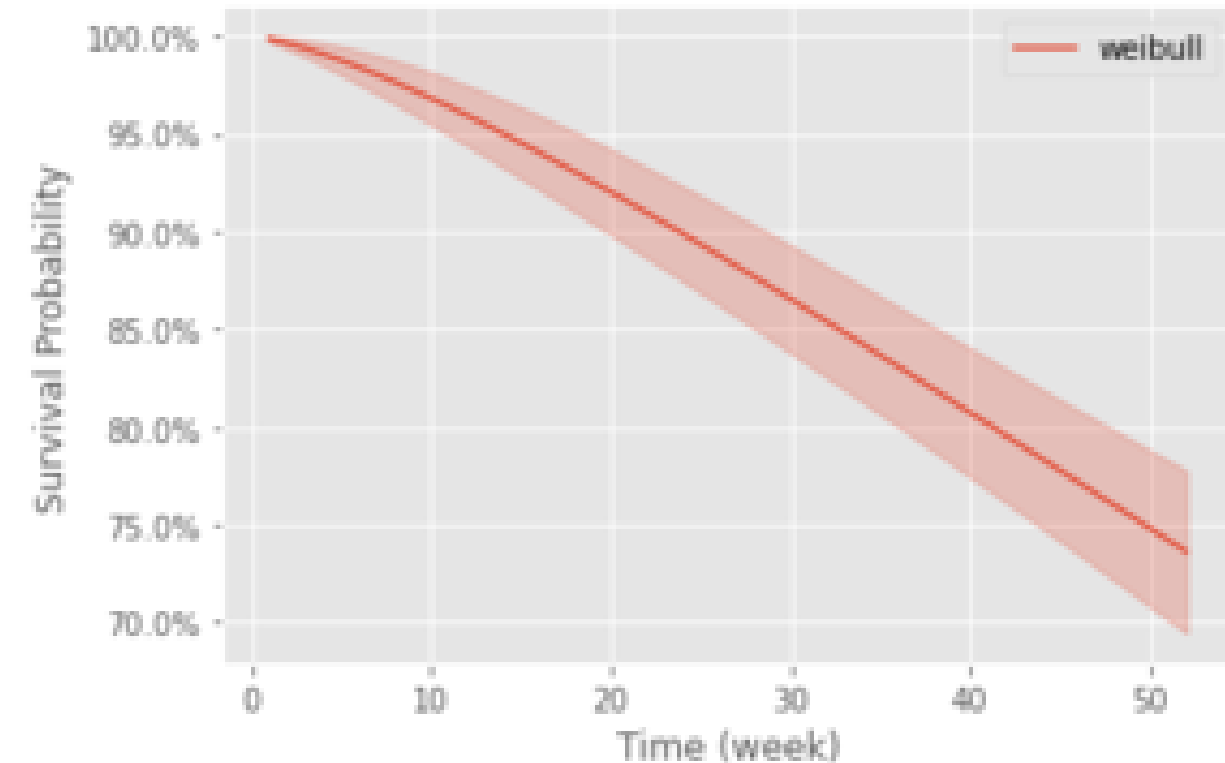
Kaplan-Meier estimator

- Non-parametric model



Weibull model

- Parametric model



The lifelines package

Kaplan-Meier estimator

```
from lifelines import KaplanMeierFitter
```

```
kmf = KaplanMeierFitter()  
kmf.fit()
```

Weibull model

```
from lifelines import WeibullFitter
```

```
wb = WeibullFitter()  
wb.fit()
```

Survival curve with covariates

Methods

- Log-rank test
- Weibull model
- Cox Proportional-Hazards (Cox PH) model

Thank you!

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