

# Survival Analysis and Censored Data

# Survival analysis

**Statistical method used to analyse and model the time until an event of interest occurs**

- Provides valuable insights into the time-to-event data
- Helps to understand the factors that influence the occurrence of events
- Enables the estimation of hazard rates and comparison of survival curves between different groups
- Assists in making informed decisions
- Primary goal is to estimate the survival function
- Takes censoring into account

**Censoring refers to observations where the event of interest has not occurred for some subjects by the end of the study period**

# Survival and Censoring times

*Survival time : Time at which the event of interest occurs*

- Ex: Time at which the patient dies in a medical study
- Indicated by 'T'

*Censoring time : Time at which censoring occurs*

- Ex: Time at which the patient drops out of the study or the study ends
- Indicated by 'C'

$$\text{Observed time} = Y = \min(T, C)$$

- True survival time T is observed if  $T < C$
- Censoring time C is observed if  $T > C$

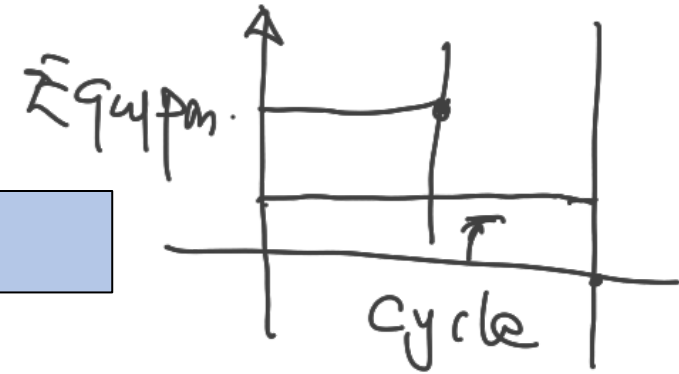
$$\text{Status indicator} = \delta = \begin{cases} 1 & \text{if } T \leq C \\ 0 & \text{if } T > C \end{cases}$$

# Types of censoring

## Right censoring

- Happens when an individual is still under observation at the end of the study
- The event of interest has not occurred for that individual
- The actual event time is unknown
- Most common type of censoring in survival analysis

*Occurs when  $T \geq Y$*



## Left censoring

- Happens when the event of interest has occurred before the start of the observation period
- Only the information that the event occurred before the study began is available
- The exact event time is unknown

*Occurs when  $T \leq Y$*

# Kaplan-Meier survival curve

Non-parametric statistic estimator

$$\text{Survival curve/Survival function} = S(t) = \Pr(T > t)$$

- Estimating survival function is complicated by the presence of censoring
- This is an approach to overcome this challenge

$$\Pr(T > d_k) = \Pr(T > d_k | T > d_{k-1}) \Pr(T > d_{k-1}) + \Pr(T > d_k | T \leq d_{k-1}) \Pr(T \leq d_{k-1})$$

Where :

- $d_1 < d_2 < \dots < d_K$  denote the  $K$  unique event times among the non-censored subjects
- $q_k$  denote the number of subjects for whom **event has occurred at time**  $d_k$
- $r_k$  denotes the number of subjects for whom the **event has not occurred** and are in the study just before  $d_k$ , **called the risk set**

$$P(A) = \overline{P(B)} \cdot P(A|B) + P(B^c) \cdot P(A|B^c)$$

# Kaplan-Meier survival curve

- It is impossible for the event to occur to the subject past time  $d_k$  if the event has not happened until an earlier time  $d_{k-1}$
- Therefore

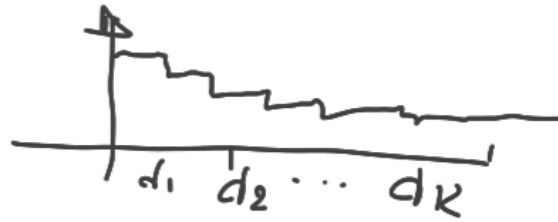
$$S(d_k) = \Pr(T > d_k) = \Pr(T > d_k | T > d_{k-1}) \Pr(T > d_{k-1})$$

- Plugging estimates of each of terms on the right side

$$\widehat{\Pr}(T > d_j | T > d_{j-1}) = (r_j - q_j) / r_j$$

$$\text{Kaplan - Meier estimator } \hat{S}(d_k) = \prod_{j=1}^k \frac{r_j - q_j}{r_j}$$

- Kaplan-Meier survival curve has a step-like shape since  $\hat{S}(t) = \hat{S}(d_k)$  for times between  $d_k$  and  $d_{k+1}$



# Log-Rank test

- Used to compare the survival curves of two or more groups or treatment arms
- EX: In case of cancer study, it is used to compare the survival of males to that of females to a treatment
- The idea of log-rank test statistic is
  - $H_0: E(X) = \mu$  for some random variable  $X$
  - Test statistic is of the form ('1' denotes the group)
  - When the sample size is large,  $W$  has approximately a standard normal distribution

$$W = \frac{X - \mu}{\sqrt{\text{Var}(X)}}, \quad X = \sum_{k=1}^K q_{1k}$$

$$\text{Expected value of } X = \mu = \sum_{k=1}^K \frac{r_1 k}{r_k} q_k$$

# Log-Rank test

- Variance of  $q_{1k}$  is

$$Var(q_{1k}) = \frac{q_k(r_{1k}/r_k)(1 - r_{1k}/r_k)(r_k - q_k)}{r_k - 1}$$

$$Var\left(\sum_{k=1}^K q_{1k}\right) \approx \sum_{k=1}^K Var(q_{1k}) = \sum_{k=1}^K \frac{q_k(r_{1k}/r_k)(1 - r_{1k}/r_k)(r_k - q_k)}{r_k - 1}$$

- Log-rank test statistic is

$$W = \frac{\sum_{k=1}^K q_{1k} - \frac{r_{1k}}{r_k} q_k}{\sqrt{\sum_{k=1}^K \frac{q_k(r_{1k}/r_k)(1 - r_{1k}/r_k)(r_k - q_k)}{r_k - 1}}}$$



# Regression models with a survival response

## Hazard function

- Also called as hazard rate/ force of mortality
- Defined as

$$h(t) = \lim_{\Delta t \rightarrow 0} \frac{\Pr(t < T \leq t + \Delta t | T > t)}{\Delta t}$$

- $T$  = Unobserved survival time
- $\Delta t$  is a small number

$$h(t) \approx \frac{\Pr(t < T \leq t + \Delta t | T > t)}{\Delta t}$$

- From

- $\Pr(A|B) = \frac{\Pr(A \cap B)}{\Pr(B)}$
- $S(t) = \Pr(T > t)$

$$h(t) = \lim_{\Delta t \rightarrow 0} \frac{\Pr((t < T \leq t + \Delta t) \cap (T > t)) / \Delta t}{\Pr(T > t)}$$

# Regression models with a survival response

## Hazard function

$$h(t) = \lim_{\Delta t \rightarrow 0} \frac{Pr(t < T \leq t + \Delta t) / \Delta t}{Pr(T > t)} = \frac{f(t)}{S(t)}$$

- $f(t)$  = Probability density function *i.e.*, Instantaneous rate of death at time  $t$
- The likelihood associated with the  $i$ th observation is

$$L_i = \begin{cases} f(y_i) & \text{if the } i\text{th observation is not censored} \\ S(y_i) & \text{if the } i\text{th observation is censored} \end{cases}$$

$$L_i = f(y_i)^{\delta_i} S(y_i)^{1-\delta_i}$$

- Assuming that the  $n$  observations are independent

$$L = \prod_{i=1}^n f(y_i)^{\delta_i} S(y_i)^{1-\delta_i} = \prod_{i=1}^n h(y_i)^{\delta_i} S(y_i)$$

# Regression models with a survival response

## Proportional Hazards

- Assumption (exponential survival)

$$h(t|x_i) = h_0(t) \exp \left( \sum_{j=1}^p x_{ij} \beta_j \right)$$

$T \leftarrow x$

Risk function

$$= h_0(t) \cdot \underline{e^{\alpha(x_i, \beta)}}$$

- $h_0(t) \geq 0$  is called the baseline hazard (unspecified function)
- $\underline{x_i}$  is the feature vector
- $\exp \left( \sum_{j=1}^p x_{ij} \beta_j \right)$  is the relative risk for the feature vector  $\mathbf{x_i} = (x_{i1}, \dots, x_{ip})^T$
- The hazard function is flexible as the probability density function is allowed to take any form
- One unit increase in  $x_{ij}$  corresponds to an increase in  $h(t|x_i)$  by a factor of  $\exp(\beta_j)$

# Regression models with a survival response

## Cox's proportional hazards model

- Makes it possible to estimate  $\beta$  without having to specify the form of  $h_0(t)$
- Assumptions
  - Each event occurs at a distinct time
  - $\delta_i = 1$ ,  $i$ th observation is uncensored
  - $y_i$  is its future time
- Hazard function for the  $i$ th observation at time  $y_i$

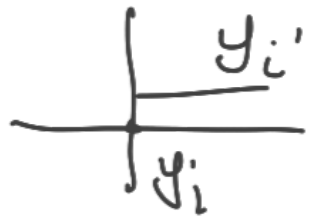
$$h(\underline{y_i} | \underline{x_i}) = \underline{h_0(y_i)} \exp \left( \sum_{j=1}^p \underline{x_{ij}} \beta_j \right)$$

$$= h_0(y_i) e^{\alpha(x_i, \beta)}$$

# Regression models with a survival response

## Cox's proportional hazards model

- Total hazard at time  $y_i$  for the at risk observations is



$$\sum_{i': y_{i'} \geq y_i} h_0(y_i) \exp \left( \sum_{j=1}^p x_{i'j} \beta_j \right)$$

- The probability that the  $i$ th observation is the one to fail at time  $y_i$  is given by

$$\frac{h_0(y_i) \exp \left( \sum_{j=1}^p x_{ij} \beta_j \right)}{\sum_{i': y_{i'} \geq y_i} h_0(y_i) \exp \left( \sum_{j=1}^p x_{i'j} \beta_j \right)} = \frac{\exp \left( \sum_{j=1}^p x_{ij} \beta_j \right)}{\sum_{i': y_{i'} \geq y_i} \exp \left( \sum_{j=1}^p x_{i'j} \beta_j \right)}$$

$$= \frac{e^{\alpha(x_i, \beta)}}{\sum_{i': y_{i'} \geq y_i} e^{\alpha(x_{i'}, \beta)}}$$

# Regression models with a survival response

## Cox's proportional hazards model

### Partial likelihood

- Valid regardless of the true value of  $h_0(t)$ , making the model very flexible and robust
- Does not correspond exactly to the probability of the data under assumption. However, it is a very good approximation

$\prod_{i: \delta_i=1} p_i(\beta)$  ←

$$PL(\beta) = \prod_{i: \delta_i=1} \frac{\exp\left(\sum_{j=1}^p x_{ij}\beta_j\right)}{\sum_{i': y_{i'} \geq y_i} \exp\left(\sum_{j=1}^p x_{i'j}\beta_j\right)}$$

→ Numerical optimization

- $\beta$  is estimated by maximizing the partial likelihood with respect to  $\beta$

# Regression models with a survival response

## Cox's proportional hazards model or log-rank test?

Case: For a single predictor case ( $p = 1$ ), which is assumed to be binary ( $x_i \in \{0, 1\}$ )

**In the case of a single binary covariate, the score test for  $H_0 : \beta = 0$  in Cox's proportional hazards model is equal to the log-rank test**

- Thus, it does not matter which approach is being used

$$y = \beta_0 + \beta_1 x$$

$H_0 : \beta_0 = 0, \beta_1 = 0$   
 $H_1 : \beta_0 \neq 0, \beta_1 \neq 0$

# Shrinkage for Cox model

- Similar to 'loss+penalty' formulation

$$\underset{\beta_0, \beta_1, \dots, \beta_p}{\text{minimize}} \{L(X, y, \beta) + \lambda P(\beta)\}$$

$L(X, y, \beta)$  – Loss function,  $P(\beta)$  – Penalty function,  $\lambda$  – Tuning parameter

- Consider minimizing a penalized version of the negative log partial likelihood

$$-\log \left( \prod_{i: \delta_i=1} \frac{\exp \left( \sum_{j=1}^p x_{ij} \beta_j \right)}{\sum_{i': y_{i'} \geq y_i} \exp \left( \sum_{j=1}^p x_{i'j} \beta_j \right)} \right) + \lambda P(\beta)$$

Where (i)  $P(\beta) = \sum_{j=1}^p \beta_j^2$  for ridge penalty

(ii)  $P(\beta) = \sum_{j=1}^p |\beta_j|$  for lasso penalty

- When  $\lambda = 0$ , minimization is equivalent to maximizing the usual Cox partial likelihood
- When  $\lambda > 0$ , minimizing yields a shrunken version of the coefficient estimates



# Shrinkage for Cox model

- When  $\lambda$  is large, then using a ridge penalty will give small coefficients that are not equal to zero
- When  $\lambda$  is sufficiently large, using a lasso penalty will give some coefficients exactly equal to zero

Partial likelihood fn.

$$PL = \prod_{i, \delta_i=1} \frac{e^{\alpha(x_i, \beta)}}{\sum_{i', y_{i'} \geq y_i} e^{\alpha(x_{i'}, \beta)}}$$

$$-\log PL = -\frac{1}{N} \sum_{\substack{L \\ \delta_i=1}} \left( \alpha(x_i, \beta) - \log \left( \sum_{i', y_{i'} \geq y_i} e^{\alpha(x_{i'}, \beta)} \right) \right)$$

Decision tree  $X_i = [x_{i1}, \dots, x_{pi}]^T$  and

Deep  
NN.  
ML

→  $\alpha(X_i, \beta)$

Chap. 11: An Introduction to  
statistical learning,

Second edition

hastie / Tibshirani.

# AUC for survival analysis

- Area under the curve is a way to quantify the performance of a two-class classifier
- Generalizing the notion to survival analysis:
  - Estimated risk score is calculated using the Cox model coefficients

$$\text{Estimated risk score} = \hat{\eta}_i = \hat{\beta}_1 x_{i1} + \dots + \hat{\beta}_p x_{ip} \quad \text{for } i = 1, \dots, m$$

- If  $\hat{\eta}_{i'} > \hat{\eta}_i$ , the model predicts that  $i'$ 'th observation has a larger hazard than the  $i$ th observation
- Thus survival time  $t_i > t_{i'}$
- Harrell's concordance index (C-index) computes the proportion of observation pairs for which  $\hat{\eta}_{i'} > \hat{\eta}_i$  and  $y_i > y_{i'}$

$$C = \frac{\sum_{i,i': y_i > y_{i'}} I(\hat{\eta}_{i'} > \hat{\eta}_i) \delta_{i'}}{\sum_{i,i': y_i > y_{i'}} \delta_{i'}}$$

$$\begin{aligned} I(\hat{\eta}_{i'} > \hat{\eta}_i) &= 1 \text{ if } \hat{\eta}_{i'} > \hat{\eta}_i \\ I(\hat{\eta}_{i'} > \hat{\eta}_i) &= 0 \text{ otherwise} \end{aligned}$$

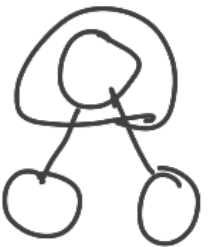
# Time – dependent covariates

- Time –dependent covariates: Predictors whose value may change over time
- Ex: Patient's blood pressure  $x_i \rightarrow x_i(t)$
- Proportional hazards model has the ability to handle time-dependent covariates
- For the example: the blood pressure,  $x_{ij}$  and  $x_{i,j}$  is replaced with  $\underline{x_{ij}(y_i)}$  and  $\underline{x_{i,j}(y_i)}$  respectively

$$\boxed{x_{ij}} \beta_j \rightarrow x_{ij}(t_i)$$

## Survival Trees

- Survival trees are a modification of classification and regression trees that use a split criterion
- It maximizes the difference between the survival curves in the resulting daughter nodes.
- Survival trees can then be used to create random survival forests



Tutorial for: RUL, Survival Analysis.  
Q - states & Discrete analysis

