Introduction to Survival Analysis

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Seminar Tutorial at Apex Lab



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Introduction to Survival Analysis

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Outline

Background

- Probability
- Censored Data
- Challenges

Methodology

- Non-parametric Models
 - Kaplan Meier Estimator
 - Survival Tree
- Parametric Model
 - Cox Hazard Proportional Model
 - Deep Survival Analysis

Evaluation

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Probability

Probability

Probability Density Function (P.D.F.):

$$p_t(t) = Pr(T = t) . \tag{1}$$

Cumulative distribution function (C.D.F.):

$$w_t(t) = Pr(T < t) = \int_0^t p_t(v) dv$$
 . (2)



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Censored Data

Censored Data

Right Censored Data

The event happens after the observation time.

• *E*: Event; *t*_{obsv}: The observe time;

•
$$\{(\mathbf{x}, t_{obsv}, e = \mathsf{True}/\mathsf{False})\};$$

• $\{(x, T_E)\}, T_E$ is the event happening log.

Example

- Patient's survival time.
- The true winning price of a bidding auction.
- The next visit time of the user.



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Challenges

Challenges



Right Censorship

- Partially data usage: discard large data for learning.
- Right Censorship: only know that the event happening time is greater than the observing time window.
- Evaluation: proper evaluation metric is needed.



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Modeling Right Censored Data in Display Ads

Losing and Winning in 2nd-price Auction



Modeling Right Censored Data

Right Censored

Right Censorship

As in 2nd price auction, if you *lose*, you only know that the *market price* is higher than your bidding price, which result in *right censorship*.



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Kaplan Meier Estimator

Preliminaries

- $S(t) = Pr(t < T_E)$: Survival rate
- F(t) = 1 S(t): Failing rate.

Algorithm

The estimator for an individual is given by

$$S(t) = \prod_{i:t_i \le t} \left(1 - \frac{d_i}{n_i} \right) , \qquad (3)$$

where d_i is the number of events and n_i is the total individuals at risk at time *i*.

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APEX

Survival Tree with Kaplan Meier Methods



Cons of KM

- Corse grained, the same for all individuals.
- Statistcal method, cannot apply personalized forecasting.



Survival Tree with Kaplan Meier Methods



Cons of KM

- Corse grained, the same for all individuals.
- Statistcal method, cannot apply personalized forecasting.

Question

How to apply an appropriate *clustering* method for one individual?

Tree-based Mapping

Goal

Given the auction feature x, forecast the market price distribution $p_x(z)^a$.

^aYuchen Wang, Kan Ren, Weinan Zhang, Yong Yu. Functional Bid Landscape Forecasting for Display Advertising. ECML-PKDD, 2016.



Tree-based Mapping

Methodology



Node Splitting



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Node Splitting KLD and Clustering

Kullback-Leibler Divergence (KLD)

A measure of the difference between two probability distributions P and Q.



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Node Splitting KLD and Clustering

Kullback-Leibler Divergence (KLD)

A measure of the difference between two probability distributions P and Q.

Node Splitting (one step)

Divide all the category (including in this node) values into two sets, maximizing KLD between the resulted two sets.



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Node Splitting KLD and Clustering

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Node Splitting (one step)

Divide all the category (including in this node) values into two sets, maximizing KLD between the resulted two sets.

Algorithm

Using K-Means Clustering according to KLD values.



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Node Splitting KLD and Clustering



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Handling Censorship

Survival Model

- For winning auctions: We have the true market price value.
- For lost auctions: We only know our proposed bid price and know that the true market price is higher than that.

Intuition

Most related works focus only on the winning auctions without considering the lost auction, which contains the information to infer the true distribution.

$$(b_i, w_i, m_i)_{i=1,2,\cdots,M} \longrightarrow (b_j, d_j, n_j)_{j=1,2,\cdots,N}$$

 $b_j < b_{j+1}$, d_j is number of winning auctions by $b_j - 1$, n_j is number of lost auctions by $b_j - 1$. So

$$w(b_x) = 1 - \prod_{b_j < b_x} \frac{n_j - d_j}{n_j}, \quad p(z) = w(z+1) - w(z).$$

Survival Model



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Cox Hazard Proportional Model

Hazard Rate The rate of the event happening given *not happened* before. Hazard Function The function $\lambda(t|\mathbf{x})$ to predict the hazard rate w.r.t. the covariate input \mathbf{x} .

Hazard Proportional Model The hazard function which models with the proportional relationship with the input covariate, where $\lambda(t|\mathbf{x}) = \lambda_0(t) \exp(h(\mathbf{x})).$

Example

Linear Cox Hazard Model: $h(\mathbf{x}) = \beta \mathbf{x}$. Question: What if $h(\mathbf{x})$ is non-linear?



Discussion

Relationship among hazard rate λ , P.D.F. function p(z), C.D.F. function S(b)

$$\lambda(b) = \lim_{db \to 0} \frac{\Pr(b \le z \le b + db|z > b)}{db}$$

$$= \lim_{db \to 0} \frac{\Pr(b \le z \le b + db)/\Pr(z > b)}{db}$$
(5)
$$= \lim_{db \to 0} \frac{(w_z(b + db) - w_z(b))/S(b)}{db} = \frac{p_z(b)}{S(b)} = -\frac{S'(b)}{S(b)} .$$

$$p_t(t|\mathbf{x}) = \frac{\partial w_t(t|\mathbf{x})}{\partial t} = \frac{-\partial S(t|\mathbf{x})}{\partial t}$$

$$= \frac{\partial \exp(\int_0^t \lambda(v|\mathbf{x})dv)}{\partial t}$$
(6)
$$= \exp\left(\int_0^t \lambda(v|\mathbf{x})dv\right) \lambda(t|\mathbf{x}) .$$

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Cost Function: Partial Likelihood

$$\begin{aligned} \mathsf{Likelihood}_{i} &= \frac{\lambda(t_{i}|\mathbf{x}_{i})}{\sum_{j:t_{j}>t_{i}}\lambda(t_{i}|\mathbf{x}_{j})} \\ &= \frac{\lambda_{0}(t_{i})e^{h(\mathbf{x}_{i})}}{\sum_{j:t_{j}>t_{i}}\lambda_{0}(t_{i})e^{h(\mathbf{x}_{j})}} \\ &= \frac{e^{h(\mathbf{x}_{i})}}{\sum_{j:t_{j}>t_{i}}e^{h(\mathbf{x}_{j})}} \\ \mathcal{L}_{\mathsf{PL}} &= -\log \prod_{i:(\mathbf{x}_{i},t_{i})}\mathsf{Likelihood}_{i} \\ &= -\sum_{i:(\mathbf{x}_{i},t_{i})} \left(h(\mathbf{x}_{i}) - \log \sum_{j:t_{j}>t_{i}}e^{h(\mathbf{x}_{j})}\right) \end{aligned}$$
(8)

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Parametric Model

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Base Hazard Function

Example

Weibull Distribution:
$$\lambda_0(t) = rac{k}{\eta} \left(rac{t}{\eta}
ight)^{k-1} \cdot e^{-(t/\eta)^k}$$
.
Question: formulation assumption; without considering **x**



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Deep Survival Analysis

NN-based Cox Model

Using deep neural network to model $h(\mathbf{x})$.^{*a*} ^{*b*} ^{*c*}

^aFaraggi D, Simon R. A neural network model for survival data[J]. Statistics in medicine, 1995.

^bRanganath R, Perotte A, Elhadad N, et al. Deep Survival Analysis[C]//Machine Learning for Healthcare Conference. 2016.

^cLuck M, Sylvain T, Cardinal H, et al. Deep Learning for Patient-Specific Kidney Graft Survival Analysis[J]. arXiv preprint arXiv:1705.10245, 2017.



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Deep Survival Analysis

Generative NN-based Survival Time Estimation^a

^aDeep Multi-task Gaussian Processes for Survival Analysis with Competing Risks, **NIPS 2017**

$$\begin{split} & f_Z \sim \mathcal{GP}(0, \mathbb{K}_{\Theta_Z}), \quad f_T \sim \mathcal{GP}(0, \mathbb{K}_{\Theta_T}) \\ & \mathbb{Z}_i \sim \mathcal{N}(f_Z(\mathbb{X}_i), \sigma_Z^2 \mathbb{I}), \quad \mathbb{T}_i \sim \mathcal{N}(f_T(\mathbb{X}_i), \sigma_T^2 \mathbb{I}) \\ & T_i = \min(T_i^1, \dots, T_i^K) \;. \end{split}$$

(9)



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Deep Survival Analysis DeepHit (Lee et al. AAAI 2018)

Output (softmax) Layer



DeepHit: A Deep Learning Approach to Survival Analysis with Competing Risks. Lee et al. AAAI 2018

Deep Survival Analysis DeepHit (Lee et al. AAAI 2018)



$$S(b) = P(z \le b | \mathbf{x})$$

= $\sum_{j=0}^{b} P(z = z_j | \mathbf{x})$ (10)

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Evaluation

Log-Likelihood

$$\bar{P} = -\frac{1}{N} \sum_{(\boldsymbol{x}_i, z_i) \in D^{\text{test}}} \log p'_z(z_i | \boldsymbol{x}_i) , \qquad (11)$$

where $N = |D^{\text{test}}|$ is the number of the test dataset and $p'_t(t|\mathbf{x})$ is the learned P.D.F.



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Evaluation

Relationship between hazard and P.D.F.

$$\lambda(b) = \lim_{db \to 0} \frac{\Pr(b \le z \le b + db|z > b)}{db}$$

$$= \lim_{db \to 0} \frac{\Pr(b \le z \le b + db)/\Pr(z > b)}{db} \qquad (12)$$

$$= \lim_{db \to 0} \frac{(w_z(b + db) - w_z(b))/S(b)}{db} = \frac{p_z(b)}{S(b)} = -\frac{S'(b)}{S(b)} .$$

$$p_t(t|\mathbf{x}) = \frac{\partial w_t(t|\mathbf{x})}{\partial t} = \frac{-\partial S(t|\mathbf{x})}{\partial t}$$

$$= \frac{\partial \exp(\int_0^t \lambda(v|\mathbf{x})dv)}{\partial t} \qquad (13)$$

$$= \exp\left(\int_0^t \lambda(v|\mathbf{x})dv\right) \lambda(t|\mathbf{x}) .$$

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Evaluation

Concordance Index (C-index)

Considering all possible pairs $(T_i, E_i), (T_j, E_j)$ for $i \le j$, the C-index is calculated by considering the number of pairs correctly ordered by the model divided by the total number of admissible pairs. *admissible*: can be ordered in a meaningful way. (uncensored, uncensored); (uncensored, right-censored). admissible pairs.



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