DS-GA 3001.009 Applied Statistics: Homework #4

Due on Thursday, October 19, 2023

Please hand in your homework via Gradescope (entry code: RKXJN2) before 11:59 PM.

- 1. In class we talked about how to estimate β in the Cox model. This problem investigates the estimation of the baseline survival function S(t) (i.e. the survival function for an individual with x = 0).
 - (a) Based on the lecture note, explain why the following is a reasonable estimator:

$$\widehat{S}(t) = \exp\left(-\sum_{i:t_i \le t} \frac{\mathbb{1}(\Delta_i = 1)}{\sum_{k \in R_i} \exp(x_k^{\top} \widehat{\beta})}\right).$$

Here R_i is the risk set at time t_i , and $\widehat{\beta}$ is the estimate of β from the Cox model.

- (b) If there is no feature (i.e. $\beta = \hat{\beta} = 0$), comment on the similarities and differences between the above estimator and the Kaplan-Meier estimator for S(t).
- 2. A dataset consists of n observations $(x_1, y_1), \dots, (x_n, y_n)$, with $x_i \in \mathbb{R}^p, y_i \in \mathbb{N}$, following a multinomial model $(y_1, \dots, y_n) \sim \text{Multi}(N; (p_1, \dots, p_n))$ with

$$p_i = \frac{\exp(x_i^\top \beta)}{\sum_{j=1}^n \exp(x_j^\top \beta)}.$$

(a) Show that the log-likelihood under this model is given by $\ell_{\rm M}(\beta) + c$, where

$$\ell_{\mathrm{M}}(\beta) = \sum_{i=1}^{n} y_i \left(x_i^{\top} \beta - \log \left(\sum_{j=1}^{n} \exp(x_j^{\top} \beta) \right) \right),$$

and $c \in \mathbb{R}$ is independent of β .

(b) The Poissonization trick introduces an additional parameter $\phi \in \mathbb{R}$ and the following log-likelihood

$$\ell_{\mathrm{P}}(\beta, \phi) = \sum_{i=1}^{n} \left(y_i(x_i^{\top} \beta + \phi) - e^{x_i^{\top} \beta + \phi} \right).$$

Show that ℓ_{M} is the profile likelihood of ℓ_{P} , i.e. $\ell_{\mathrm{M}}(\beta) = \max_{\phi \in \mathbb{R}} \ell_{\mathrm{P}}(\beta, \phi) + c'$ for some constant $c' \in \mathbb{R}$ independent of β .

- (c) How does the result in (b) justify the use of Poissonization in Lindsey's method? You may assume $\Delta_k \equiv \Delta$ and $h(z_k) \equiv 1$ in your discussion.
- 3. Coding: we will explore an AIDS dataset and understand the effects of different treatments on the survival curves for different patients. Based on the inline instructions, fill in the missing codes in https://tinyurl.com/4bdcyy7c. Be sure to submit a pdf with your codes, outputs, and colab link.

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