

PH C240B: Assignment 3

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Problem 1

Show the CAR condition, $x \rightarrow Pr(O = o|X = x)$ for $x \in \mathcal{C}(o)$ is constant implies $Pr(X = x|O = o) = Pr(X = x|x \in \mathcal{C}(o))$. You may assume all random variables here are discrete for simplicity.

Intuitively, the CAR assumption says that the observation of $O = o$ is not influenced by the specific value of X in $\mathcal{C}(O)$ which was taken, only by the fact that X did take a value in $\mathcal{C}(O)$. Thus, the CAR condition is equivalent to

$$Pr(O = o|X = x) = Pr(O = o|X \in \mathcal{C}(O)) \text{ for all } o, x \in \mathcal{C}(o)$$

which can be rewritten as,

$$Pr(O = o|X = x \text{ and } X \in \mathcal{C}(O)) = Pr(O = o|X \in \mathcal{C}(O)) \text{ for all } x \in \mathcal{C}(o) .$$

clearly identifying the conditional independence assumption: given $X \in \mathcal{C}(O)$, the events $X = x$ and $O = o$ are conditionally independent. By symmetry of conditional independence we have,

$$Pr(X = x|O = o \text{ and } X \in \mathcal{C}(O)) = Pr(X = x|X \in \mathcal{C}(O)) .$$

But, since the former is equal to $Pr(X = x|O = o)$, we have that CAR is equivalent to

$$Pr(X = x|O = o) = Pr(X = x|X \in \mathcal{C}(O)) \text{ for all } x \in \mathcal{C}(o) .$$

So we see that this conditional probability of the full data, X , given O is only dependent the distribution of P_X . Thus the observation of $O = o$ tells us no more, in the sense of what is now the conditional distribution of X , than the obvious $X \in \mathcal{C}(O)$

Problem 2

Let $P_{X,\epsilon}$ be a path through P_X , the distribution of the full data, X , and having score $S_1(X)$. This then defines a path $P_{P_X,\epsilon,G}$ through the observed data distribution, $P_{P_X G}$. Show that the scores generated by these paths are $E[S_1(X)|O = o]$.

$$\begin{aligned} \frac{d}{d\epsilon} \log dP_{P_X,\epsilon,G} / dP_{P_X G} |_{\epsilon=0} &= \frac{d}{d\epsilon} \log dP_{P_X,\epsilon,G} |_{\epsilon=0} \\ &= \int \frac{dP(O = o|X = x) S_1(x) dP(x) d\nu(x)}{dP_{P_X G}(o)} \\ &= \int S_1(x) dP(x|O = o) d\nu(x) \\ &= E[S_1(X)|O = o] . \end{aligned}$$

Problem 3

Let G_ϵ be a path through G , the distribution of the censoring time, C , given X , having score $S_2(C, X)$. This then defines a path $P_{P_X G_\epsilon}$ through the observed data distribution, $P_{P_X G}$. Show that the scores generated by these paths are $E[S_2(C, X)|O = o]$.

$$\begin{aligned} \frac{d}{d\epsilon} \log dP_{P_X G_\epsilon} / dP_{P_X G} |_{\epsilon=0} &= \frac{d}{d\epsilon} \log dP_{P_X G_\epsilon} |_{\epsilon=0} \\ &= \int \frac{S_2(c, x) dP(O = o | X = x) dP(x) d\nu(x)}{dP_{P_X G_\epsilon}(o)} \\ &= \int S_2(c, x) dP(x | O = o) d\nu(x) \\ &= E[S_2(C, X) | O = o] . \end{aligned}$$

Problem 4

This problem involves simulating data under a general Cox model. Let's make the assumption we have a conditional hazard of death at time, t , given by $\lambda(t|X) = \lambda_0(t) \exp(f_\beta(X))$ where X is a set of covariates and f_β is a function indexed by β , say finite dimensional. Assume the baseline hazard is $\lambda_0(t) = \exp(rt)$ for positive r . Given X , what is the distribution of death times? Prove your answer.

Problem 5

Complete the first problem from LabCox in the lab section of the files on bCourses.

Specifically, we are asked to use the model generated in the lab to simulate 1000 draws of $n = 1000$ and check coverage on the coefficient, which represents the log of the hazard proportion between treated and non-treated, in this case. Also, according to the first problem from LabCox, "We also know that the true proportional hazard of treated to untreated is $\exp(-1)$, the coefficient of treatment being -1".

```
# we repeat the experiment 1000 times
N = 1000

# allocating memory to store the experiment
exp <- data.frame(Truth = numeric(N), Estimate = numeric(N), CI_lower = numeric(N),
                  CI_upper = numeric(N), CI_indicator = numeric(N))

for(i in 1:N){

  # generating the original model
  n = 1000

  # draw the W from standard normals
  W1 = rnorm(n)
  W2 = rnorm(n)
  g0 = function(W1,W2) plogis(.1 + W1*W2)
  A = rbinom(n, 1, g0(W1,W2))

  # draw T and C --both generated with random uniforms as perviously described but C could
  # be generated anyway independent of T given W for identifiability purposes
```

```

C = -log(runif(n))/(.01*exp(.3*W1))
T = -log(runif(n))/(.02*exp(2*W1^2 - A))

Ttilde = pmin(T, C)
Delta = C >= T & T<= 100

# Create the survival object
S = Surv(time = Ttilde, event = Delta, type = "right")
data = data.frame(A = A, W1 = W1, W2 = W2)
coxfit = coxph(S ~ ., data = data)

truth <- exp(-1)
conf_interval <- as.data.frame(summary(coxfit)[8])
lower <- conf_interval$conf.int.lower..95[1]
upper <- conf_interval$conf.int.upper..95[1]
indicator <- as.numeric(truth >= lower && truth <= upper)

# populating the data frame where we store our experiment
exp$Estimate[i] <- conf_interval$conf.int.exp.coef.[1]
exp$Truth[i] <- exp(-1)
exp$CI_upper[i] <- upper
exp$CI_lower[i] <- lower
exp$CI_indicator[i] <- indicator
}

# calculating the coverage
print(paste('Our coverage is', mean(exp$CI_indicator)))

## [1] "Our coverage is 0"

```

Our coverage on the coefficient, which represents the log of the hazard proportion between treated and non-treated, is 0, meaning none of our confidence intervals covered the true proportional hazard of treated to untreated. This is because we simulated data according to a model that defies the Cox assumption. That is, we assume the hazard of an individual with covariates A , W_1 , W_2 at time, t , is $.02\exp(2W_1^2 - A)$, a main terms *non-linear* function of the hazard and the Cox assumption requires the main terms to be a *linear* functional form of the hazard.

Bonus

Assume a CAR model for full data consisting of survival time, censoring time, the continuous baseline covariates and randomly assigned treatment indicator. We have observed data $\min(T, C), \Delta$ along with the covariates and treatment indicator. Someone receives a data set of 1000 independent subjects drawn from this model from an RCT and runs a Cox Proportional Hazards regression with treatment as the only covariate, showing a significantly negative coefficient. Can you convince this person he may be wrong via simulation? Explain how you set up your simulation and turn in your code to show the results.