Biostatistics 276 Assignment 3

PROJECT SUBMISSION INSTRUCTIONS

Please complete and save your project in pdf file. Before submission, please use the following naming convention:

firstname-lastname-P2.pdf

Your pdf file should be submitted to the e-mail address below:

BIOS276.31r8w8ndztx2a19n@u.box.com

Please do not use my personal e-mail address. After submission you should receive confirmation of successful uploading. I will let you know if I am missing your project by the day following the due date.

Please describe all your work in clear terms, before implementing R code. Each question should include a description of your approach with clear indication of where I can find the associated source code. Your code should be attached to your assignment or uploaded online to a repository I can freely access.

1. Consider the data-set diabetes data available from lars, (use the matrix of predictors x). The outcome is indexed with y and predictors are indexed by x a matrix of 10 predictors.

Add a intercept column in the data design matrix X, and consider the following regression model

$$Y = X\beta + \epsilon$$

with $X: n \times p$, $\beta: p \times 1$ and $\epsilon \sim N(0, \sigma^2 I_n)$.

We are interested in model selection by including only a subset of the original p predictors. The number of possible models is 2^p .

Let $\gamma \in \{0,1\}^p$ define a p-dimensional vector of predictor indicators, such that $\gamma_j = 1$ if X_j is included in the regression, otherwise $\gamma_j = 0$, (j = 1, ..., p). Also, let X_{γ} and β_{γ} denote the subset of predictors and regression coefficients associated with components of γ that are equal to 1.

Conditional on γ the sampling model is

$$Y \mid \gamma, \beta, \sigma^2 \sim N(X_{\gamma}\beta_{\gamma}, \sigma^2 I_n),$$

with improper prior:

$$p(\beta_{\gamma}, \sigma^2 \mid \gamma) \propto (\sigma^2)^{-\frac{q_{\gamma}}{2} - 1} \exp\left[-\frac{1}{2c\sigma^2}\beta_{\gamma}'(X_{\gamma}'X_{\gamma})\beta_{\gamma}\right]$$

where $q_{\gamma} = \sum_{j=1}^{p} \gamma_{j}$ is the model size, given a realization of γ and c = n, the sample size. Finally, assume

$$p(\gamma \mid \alpha) = \prod_{i=1}^{p} \alpha^{\gamma_i} (1 - \alpha)^{(1 - \gamma_i)}, \text{ with } \alpha \sim Beta(0.5, 0.5)$$

- **a.** Develop and implement a Reversible Jumps MCMC algorithm exploring the model space $p(\gamma \mid Y)$. Report the estimated inclusion probabilities $p(\gamma_j = 1 \mid Y)$, for all j = 1, 2, ..., p.
- **b.** Compare your results in (a) with results obtained sampling directly from $p(\gamma \mid Y)$. You should be able to implement this without the need for RJ Monte Carlo methods.

Due: 02/28/2023