

**Assignment #5— Due Monday, 6 Dec.**

\*This homework covers Chapter 9. Submit your homework on Canvas or send it to our TA, Mr. LYU Zhongyuan (zlyuab@connect.ust.hk).

\*No late homework will be accepted for credit.

\*Append the R codes you used to your submission. *If the problem does not need R or is not explicitly stated to complete in R, then you should just do it by hand with a calculator.*

\*In case of rounding error, keep 3 figures after the decimal point.

**Problem 1** (Use R) The purpose is to predicting whether a patient has breast cancer based on 9 predictors (features of the tumor) as follows:

*Cl.thickness*: Clump Thickness

*Cell.size*: Uniformity of Cell Size

*Cell.shape*: Uniformity of Cell Shape

*Marg.adhesion*: Marginal Adhesion

*Epith.c.size*: Single Epithelial Cell Size

*Bare.nuclei*: Bare Nuclei

*Bl.cromatin*: Bland Chromatin

*Normal.nucleoli*: Normal Nucleoli

*Mitoses*: Mitoses

The response is the variable *class* taking two possible values *benign* and *malignant*. The data was collected by Dr. Wolberg from his clinical cases, containing 683 observations, stored in the file *BreastCancer.txt*.

- (a) Fit a logistic regression model using all the predictor variables, report the summary and interpret your results. Is the model significant?
- (b) Based on part (a), construct the 95% confidence interval for  $\beta_{Cl.thickness}$  and test the hypothesis

$$H_0 : \beta_{Cell.shape} = 0 \quad \text{versus} \quad H_1 : \beta_{Cell.shape} \neq 0$$

at the significance level 0.10.

- (c) Use the predictors *Cl.thickness*, *Cell.shape*, *Marg.adhesion*, *Bare.nuclei*, *Bl.cromatin* to fit a logistic regression model, report the summary and compare it to the full model in part (a).
- (d) Given part (c) and a new patient with *Cl.thickness*=6, *Cell.shape*=3, *Marg.adhesion*=8, *Bare.nuclei*=2, *Bl.cromatin*=5, what is the probability of this patient's tumor being *benign*?
- (e) Compare the model in part (c) and all its subset models (i.e., models only using variables from *Cl.thickness*, *Cell.shape*, *Marg.adhesion*, *Bare.nuclei*, *Bl.cromatin*), which of them has the smallest AIC? Report your conclusions.