

Assignment 1 – Due February 8, 2023

- 1) Find the full iris data set called *iris.csv* on Brightspace.
 - (a) Read the data into an R variable called *dat* and label all missing data (-99) as R's *NA*.
 - (b) Compute the sample means and variances of the widths and lengths of both petals and sepals. Use variable names *mdat* for the means and *vdat* for the variances.
 - (c) Add a new column called “Bsepal” to *dat* which is an indicator for sepal width ≥ 3.1 (1 if larger 0 otherwise).
 - (d) Add a new versicolor iris with sepal length, sepal width, petal length and petal width of 6.1, 3.2, 4.1, 1.3 respectively.
 - (e) Break the data for the three types of irises (setosa, versicolor, virginica) into variables named *datse*, *datve* and *datvi* respectively.
 - (f) Repeat (b) for the three types of irises. Use a similar naming convention so add an *m* for the sample means and *v* for the sample variances.

Submit your R code as *q1.R*. Also, please label each part (a)-(f) with a comment header (i.e. `# (a)`). On any line R ignores anything after `#`.

- 2) Write R code that will compute the root of the following function

$$f(x) = \frac{1}{1 + e^{-(x-1)}} - \frac{1}{2}.$$

Submit your R code as *q2.R*.

- 3) The probability density function for a normal distribution (“bell curve”) with mean μ and variance σ^2 is given by

$$f(x) = \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{(x-\mu)^2}{2\sigma^2}}, \quad x \in \mathbb{R}, \mu \in \mathbb{R}, \sigma^2 > 0.$$

- (a) Write an R function called *mdnorm1* that computes $f(x)$ for any value of x , μ and σ .
- (b) Copy the function in (a) and call it *mdnorm2*. Set the defaults for μ and σ to be 0 and 1 respectively.
- (c) Copy the function in (b) and call it *mdnorm3*. Add a new input called *log* that defaults to *FALSE*. If *FALSE* return $f(x)$ otherwise return $\ln f(x)$. Make sure you first simplify the expression returned when *TRUE*.

Submit your R code as *q3.R*. Also, please label each part (a)-(c) with a comment like you did for question 1).