

# Question 1 - Basic Math

# Part a - Function to compute the log of a positive number

log(64)

# Part b - What is the default base for the log function?

log(x,base)

# Part c - the log of a negative number

# The real logarithm is undefined for negative real numbers produces a NAN

log(-15)

# Part d - the square-root of a positive number

sqrt(25)

# Question 2 - Random Number Generation

# Part a - Create a vector of 15 standard normal random

# variables calculate its mean and SD (Standard Deviation)

random\_vector <- rnorm(15)

random\_vector

mean(random\_vector)

sd(random\_vector)

# Part b - Change the mean to 10 and the SD to 2 to recalculate

# the vector of 15 random normal variables. Calculate its mean

# and SD

random\_vector <- rnorm(15, m = 10, sd = 2)

random\_vector

mean(random\_vector)

```
sd(random_vector)
```

```
# Because they are both random generated and your
```

```
# changing the mean of the 2nd one produced random numbers around
```

```
# that 2nd mean.
```

```
# Question 3 - Vector Operations
```

```
# Part a -- c - Create 2 vectors for heights and weights of 6 individuals
```

```
weights <- c(60, 72, 57, 90, 95, 72)
```

```
heights <- c(1.80, 1.85, 1.72, 1.90, 1.74, 1.91)
```

```
# Part d - Create a scatterplot of weight vs. height
```

```
# Interpret the scatterplot
```

```
plot(heights,
```

```
      weights,
```

```
      main = "Weight versus Height for 6 Individuals",
```

```
      xlab = "Height (m)",
```

```
      ylab = "Weight (kg)")
```

```
# height and weight are positively correlated, except
```

```
# for a shorter individual who weighs the most.
```

```
# Part e - Calculate the BMI for each individual
```

```
# BMI = weight in kg divided by the square of the height in m
```

```
BMI <- weights / (heights ^ 2)
```

```
BMI
```

```
# Part f - Calculate the mean for weight
```

```
mean_weight <- mean(weights)
```

```
mean_weight
```

```
# Part g - Subtract the mean for each value of weight
```

```
weight_difference <- weights - mean_weight
```

```
weight_difference
```

```
# part h - Sum the result
```

```
# Now you know why we square the deviations from the mean to calculate a
```

```
# standard deviation!
```

```
sum_of_weight_difference <- sum(weight_difference)
```

```
sum_of_weight_difference
```

```
# Question 4 - Enter your data science profile in R as a data frame
```

```
# Data frame consists of two columns - data science categories, ranking
```

```
# Categories - Computer Programming, math, statistics, machine learning, domain
```

```
# expertise, communication and presentation skills, and data visualization
```

```
# Ranking - 1 as worst, 5 as best for best ranking for each category
```

```
# Create a bargraph of your data science profile
```

```
Data_Science_Categories <- c("Computer Programming", "Math", "Statistics", "Machine Learning",  
"Domain Expertise", "Communication & Presentation Skills", "Data Visualization")
```

```
Ranking <- c(4,4,3,2,1,2,3)
```

```
che <- data.frame(Data_Science_Categories, Ranking)
```

```
#Setting margin settings
```

```
par(mar=c(11,4,4,4))
```

```
# make category names fit under the columns with cex.names and las
```

```
barplot(che$Ranking, main="Data Science Ranking", names.arg=che$Data_Science_Categories,  
ylab="Rankings", las=2, cex.names=.6)
```

