# Homework 1

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# August 2022

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# 1 Using R: Vectors

### 1.1 (a)

Create a vector with 10 numbers (3, 12, 6, -5, 0, 8, 15, 1, -10, 7) and assign it to x.

```
x <- c(3, 12, 6, -5, 0, 8, 15, 1, -10, 7)
```

### 1.2 (b)

Using the commands seq, min, and max with one line of code create a new vector y with 10 elements ranging from the minimum value of x to the maximum value of x.

```
# Min, max of x vector
xMin = min(x)
xMax = max(x)
INCREMENT = (xMax - xMin) / (length(x) - 1) # size of each step

# Create the vector from min of x to max of x while maintaining length of 10
y = seq(min(x), max(x), INCREMENT)
y # display

[1] -10.000000 -7.222222 -4.444444 -1.666667 1.111111 3.888889
[7] 6.6666667 9.444444 12.2222222 15.000000

# Prove is 10 elements:
paste('The length of y is:', length(y) )
```

[1] "The length of y is: 10"

# 1.3 (c)

# 1.3.1 Compute the sum, mean, standard deviation, variance, mean absolute deviation, quartiles

```
# Combine x and y into single object
  xAndY <- cbind(x, y)</pre>
  # This is what x and y look like
  head(xAndY)
[1,] 3 -10.000000
[2,] 12 -7.222222
[3,] 6 -4.44444
[4,] -5 -1.666667
[5,] 0 1.111111
[6,] 8 3.888889
  # List of all the function to calculate on x and y (excluding quintile)
  vectorOfFuns <- c('sum', 'mean', 'sd', 'var', 'mad', 'quantile')</pre>
  vectorOfFunsNames <- c('sum', 'mean', 'standard deviation', 'variance', # function full na</pre>
                          'mean absolute deviation', 'quartiles')
  for (funIdx in 1:length(vectorOfFuns)) {
    \# Print result of the function calculation output, for x and y
    print( paste( vectorOfFunsNames[funIdx], 'of x and y:' ) )
    # Calulate using a function from `vectorOfFuns`
    output <- apply(xAndY,
                                                    # Using X and Y
                                                   # There are n cols in x & y
                    ncol(xAndY),
                    paste0(vectorOfFuns[funIdx]) ) # Retrieve/apply function
    print(output)
[1] "sum of x and y:"
х у
37 25
```

```
[1] "mean of x and y:"
 х у
3.7 2.5
[1] "standard deviation of x and y:"
      X
7.572611 8.410140
[1] "variance of x and y:"
      х
57.34444 70.73045
[1] "mean absolute deviation of x and y:"
      х
5.93040 10.29583
[1] "quartiles of x and y:"
         x y
0%
    -10.00 -10.00
25% 0.25 -3.75
50%
     4.50 2.50
75% 7.75 8.75
100% 15.00 15.00
```

# 1.3.2 Compute the Quintiles

```
# Quintile for x
  quantile(x, probs = seq(0, 1, 0.20))
  0% 20%
             40%
                   60% 80% 100%
-10.0 -1.0
             2.2
                   6.4 8.8 15.0
  # Quintile for y
  quantile(y, probs = seq(0, 1, 0.20))
          0%
                       20%
                                    40%
                                                 60%
                                                               80%
-1.000000e+01 -5.000000e+00 -1.665335e-15 5.000000e+00 1.000000e+01
        100%
1.500000e+01
```

#### 1.4 (d)

Use sample() to create a new 7 element vector z by using R to randomly sample from x with replacement.

```
z = sample(x, size=7)
z # show value

[1] 1 8 15 -10 0 -5 7
```

#### 1.5 (e)

Use t.test() to compute a statistical test for differences in means between the vectors x and y.

Are the differences in means significant?

No, they are not statistically significant (90%, 95%, or 99%), given

- t-value is  $< \sim 1.96$
- p-value is > 0.10, 0.05, 0.01

#### 1.6 (f)

To sort a data frame in R, use the order() function. Sort the vector x and re-run the t-test as a paired t-test.

# 1.7 (g)

Create a logical vector that identifies which numbers in x are negative.

```
xThatAreNeg = x < 0
x # reminder of what x looks like

[1] 3 12 6 -5 0 8 15 1 -10 7

xThatAreNeg # display</pre>
```

[1] FALSE FALSE FALSE TRUE FALSE FALSE FALSE TRUE FALSE

# 1.8 (h)

Use this logical vector to remove all entries with negative numbers from x. (Make sure to overwrite the vector x so that the new vector x has 8 elements!)

```
x = subset(x, !xThatAreNeg)
```

[1] 3 12 6 0 8 15 1 7

# 2 Using R: Some missing values

#### 2.1 (a)

#### 2.1.1 Use the code below to create the dataframe X and

```
col1 <- c(1,2,3,NA,5)

col2 <- c(4,5,6,89,101)

col3 <- c(45,NA,66,121,201)

col4 <- c(14,NA,13,NA,27)

X <- rbind (col1,col2,col3,col4)
```

#### 2.1.2 Then write code to display all rows in X with missing values.

```
# All rows of X that have NAs (e.g., not complete) X[!complete.cases(X),]
      [,1] [,2] [,3] [,4] [,5]
col1
                       3
                            NA
                                   5
col3
        45
               NA
                     66
                          121
                                 201
col4
        14
               NA
                     13
                           NA
                                  27
```

# 2.2 (b)

#### 2.2.1 Use the following vector y for this part:

```
y <- c(3,12,99,99,7,99,21)
```

- i. Some statistical applications and older systems sometimes code missing values with a number, e.g., 99. In order to let R know that is a missing value you need to recode it as 'NA'. Please write a line of code that will replace any 99's in the vector y with 'NA'.
- ii. With the updated vector y, write code that will count the number of missing values in it.