Homework 1

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1. The slope of a linear regression line is 2.5 and the intercept is -1. What is the equation of the line?

$$y = -1 + 2.5x$$

What would be the predicted value of Y when x = 4.3?
For $x = 4$, $\hat{y} = -1 + 2.5(4.3) = 9.75$

2. A researcher is interested in determining if there is a difference in yield among three different cultivars. Perform the appropriate test to determine if there is a significant difference in yield among them. Use the following dataset:

```
data <- data.frame(
  Crop_Variety = c("Wheat1", "Wheat1", "Wheat2", "Wheat2", "Wheat3", "Wheat3"),
  Yield = c(3200, 2900, 2600, 2400, 2200, 2000)
)
data
```

```
##   Crop_Variety Yield
## 1       Wheat1  3200
## 2       Wheat1  2900
## 3       Wheat2  2600
## 4       Wheat2  2400
## 5       Wheat3  2200
## 6       Wheat3  2000
```

We can perform an one-way ANOVA test to check whether the difference in cultivars yield is significant. These are the null and alternative hypothesis:

$$H_0 : \text{the means are equal}$$
$$H_1 : \text{at least one mean differs}$$

```
result <- aov(Yield ~ Crop_Variety, data = data)
print(summary(result))
```

```
##              Df Sum Sq Mean Sq F value Pr(>F)
## Crop_Variety  2 910000  455000   16.06  0.025 *
## Residuals     3  85000   28333
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

For a significance level of $\alpha = 0.05$, we reject the null hypothesis that the mean yield of the three cultivars are equal, because p-value $= 0.025 < 0.05$, which means that at least one mean yield differs.

3. A researcher wants to determine if there is a significant difference in growth rate before and after a treatment. He collected data on growth rate for 6 individuals before and after the treatment. Perform the appropriate test to determine if there is a significant difference in growth rate before and after the treatment. Use the following dataset:

```
data <- data.frame(
  Plant = c(1, 2, 3, 4, 5, 6),
  Treatment = c("A", "A", "A", "B", "B", "B"),
  Before = c(10, 8, 12, 11, 9, 15),
  After = c(12, 9, 14, 15, 11, 17)
)
data
```

```
##   Plant Treatment Before After
## 1     1         A     10    12
## 2     2         A      8     9
## 3     3         A     12    14
## 4     4         B     11    15
## 5     5         B      9    11
## 6     6         B     15    17
```

We can use a paired two-sided t-test. We have to used a paired version because the sample is the same, and we have matches before and after a treatment. The two-sided over another alternative is preferred because we are only interested to check whether there's a difference between the two means (not whether one is greater or lesser then the other). These are the null and alternative hypothesis:

$$H_0 : \text{the difference between the means is } 0$$
$$H_1 : \text{the difference between the means is not equal to } 0$$

```
result <- t.test(data$Before, data$After, paired = T, alternative = "two.sided")
print(result)
```

```
##
##  Paired t-test
##
## data:  data$Before and data$After
## t = -5.398, df = 5, p-value = 0.002947
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
##  -3.198463 -1.134870
## sample estimates:
## mean difference
##       -2.166667
```

As the p-value $= 0.002947 < 0.05$, using a significance level of $\alpha = 0.05$, we reject the null hypothesis that the difference between the means is 0, i.e., the growth rate changed after applying the treatment.

4. Standardize (make it a standard normal distribution) the following data and include the new values below.

```
x <- c(9.58,  6.64,  6.75,  5.26, 19.07, 10.32, 17.62,  6.32, 14.75, 18.36,
       11.97, 18.87,  6.01, 10.51, 12.41, 17.09, 18.12, 11.84, 7.63, 14.26)
z <- (x - mean(x)) / sd(x)
print(z)
```

```
##  [1] -0.53513178 -1.14281330 -1.12007691 -1.42805156  1.42639800 -0.38217793
##  [7]  1.12669113 -1.20895550  0.53347823  1.27964498 -0.04113218  1.38505912
## [13] -1.27303076 -0.34290600  0.04981335  1.01714311  1.23003833 -0.06800246
## [19] -0.93818585  0.43219797
```

5. Explain what each term is and give an example:

a) Pseudo-replication
   Is when you use the same group of people or things repeated times to collect information, rather
   then collecting data from different groups each time. It inflates the sample size, leading to incorrect
   inferential statistical tests.
   Example: a researcher randomly selects 4 plots of land, applies a new fertilizer to all the plots, and
   measures the yield of each plot. This is a pseudo-replication, because all the plots used the same
   fertilizer (treatment), which means that the measurements are not independent from each other.

b) True replication
   Is the smallest experimental unit to which a treatment is independently applied.
   Example: a researcher randomly selects 4 plots of land, applies a different fertilizer (treatment) to each
   plot, and measures the yield of each plot. This is the case of a true replication because the researcher
   accounted for independent treatments for each group of samples.

c) Experimental unit
   Is the smallest entity that can be randomly assigned to a different treatment condition.
   Example: if a researcher randomly selects 4 plots of land, applies a different fertilizer (treatment) to
   each plot, and measures the yield of each plot, then each plot would be a experimental unit, because
   in this case is the smallest entity in which you would apply a different treatment.

d) Observational unit
   Is the unit observed (measured) by the researcher in the study.
   Example: if a researcher is studying the impact of a new fertilizer on the crop yield, the observational
   unit could be a group of plants.

e) Dependent variable
   It is the variable of interest being studied in the experiment, that might be affected by other variables
   (the independent variables).
   Example: a researcher is studying how a fertilizer (a independent variable X_1) and the quantity of
   nitrogen in the soil (another independent variable X_2) affects the crop yield (the dependent variable
   Y).

6. A researcher is interested in determining if there is a significant difference in yield between an ex-
   perimental fertilizer and a control fertilizer. He collected data on yield for 10 plots treated with the
   experimental fertilizer and 10 plots treated with the control fertilizer. The mean yield for the exper-
   imental fertilizer is 8 with a standard deviation of 2 and the mean yield for the control fertilizer is 7
   with a standard deviation also 2. Determine the 95% confidence interval for the difference in means
   and interpret the results.

We can use a t-statistic to build a confidence interval for the difference in means.
Denote A as the experimental fertilizer and B as the control fertilizer:

3

```
xbar_A <- 8
xbar_B <- 7
s_A <- s_B <- 2
n_A <- n_B <- 10

# pooled estimate of the standard deviation is used because both s_B and s_B are equal
sP <- sqrt(((((n_A - 1) * s_A ^ 2) + ((n_B - 1) * s_B ^ 2)) / (n_A + n_B - 2))

# standard error between difference of means
se_diff <- sP * sqrt((1 / n_A) + (1 / n_B))

# use the t-statistic with P(T <= alpha / 2) and "n_a + n_B - 2" degrees of freedom
ic <- c()
ic[1] <- (xbar_A - xbar_B) + (qt(p = 0.025, df = n_A + n_B - 2) * se_diff)
ic[2] <- (xbar_A - xbar_B) - (qt(p = 0.025, df = n_A + n_B - 2) * se_diff)
ic
```

```
## [1] -0.8791218  2.8791218
```

7. Answer the following:

   a) What are the dimensions of D and B in the following equation? $\underset{?\times?}{D} = \underset{3\times4}{A} \times \underset{?\times?}{B} \times \underset{10\times10}{C}$

```
# we should use conformable matrices
A <- matrix(nrow = 3, ncol = 4)
B <- matrix(nrow = 4, ncol = 10)
C <- matrix(nrow = 10, ncol = 10)
D <- A %*% B %*% C
cat('D:', paste0(dim(D), collapse = 'x'))
```

```
## D: 3x10
```

   b) Which row and column would the item $b_{1,3} = 24$ go on matrix **B**?
      In the row 1 and the column 3.

   c) What are the dimensions of A'A and AA'?

```
AtA <- t(A) %*% A
cat("A'A:", paste0(dim(AtA), collapse = 'x'), '\n')
```

```
## A'A: 4x4
```

```
AAt <- A %*% t(A)
cat("AA':", paste0(dim(AAt), collapse = 'x'))
```

```
## AA': 3x3
```

8. Define the following matrices:

a) diagonal matrix
Is a square matrix whose off-diagonal entries are all equal to zero (the diagonal from the top left corner to the bottom right corner of a square matrix is called the main diagonal or only diagonal).

b) identity matrix (What R function do we use to create it?)
Is a square matrix whose main diagonal elements equal to 1 and off-diagonal elements are zeros.

```
# create 3x3 identity matrix
diag(3)
```

```
##      [,1] [,2] [,3]
## [1,]    1    0    0
## [2,]    0    1    0
## [3,]    0    0    1
```

c) upper or lower triangular matrix
Upper triangular matrix: a square matrix whose elements below the main diagonal are zeros.
Lower triangular matrix: a square matrix whose elements above the main diagonal are zeros.

d) square matrix
Is a matrix whose quantity of rows and columns are the same.

9. Given the following matrices:

```
A = [1 2 3]
    [4 5 6]

B = [7   8]
    [9  10]
    [11 12]

C = [4 7 2]
    [5 8 4]
    [6 9 5]

k = 10
```

Calculate:
a) $A \times B$

```
A <- matrix(1:6, nrow = 2, byrow = T)
B <- matrix(7:12, nrow = 3, byrow = T)
A %*% B
```

```
##      [,1] [,2]
## [1,]   58   64
## [2,]  139  154
```

b) Inverse of C

```
C <- matrix(c(4, 7, 2, 5, 8, 4, 6, 9, 5), nrow = 3, byrow = T)
solve(C)
```

```
##              [,1]        [,2] [,3]
## [1,]   1.3333333 -5.666667    4
## [2,] -0.3333333  2.666667   -2
## [3,] -1.0000000  2.000000   -1
```

c) $A \otimes B$ (Kronecker product of A and B)

```
kronecker(A, B)
```

```
##       [,1] [,2] [,3] [,4] [,5] [,6]
## [1,]    7    8   14   16   21   24
## [2,]    9   10   18   20   27   30
## [3,]   11   12   22   24   33   36
## [4,]   28   32   35   40   42   48
## [5,]   36   40   45   50   54   60
## [6,]   44   48   55   60   66   72
```

d) Transpose of B

```
t(B)
```

```
##       [,1] [,2] [,3]
## [1,]    7    9   11
## [2,]    8   10   12
```

e) kA

```
k <- 10
k * A
```

```
##       [,1] [,2] [,3]
## [1,]   10   20   30
## [2,]   40   50   60
```

10. Represent the following system of equations with matrix notation. Also show how to create the resulting matrix with R.

$$3x + 2y = 12$$
$$4x - y = 2$$

Writing as $Av = r$:

$$\begin{bmatrix} 3 & 2 \\ 4 & -1 \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix} = \begin{bmatrix} 12 \\ 2 \end{bmatrix}$$

```
A <- matrix(c(3, 2, 4, -1), nrow = 2, byrow = T)
v <- c(12, 2)
r <- solve(A, v)
cat('x =', r[1], '\ny =', r[2])
```

```
## x = 1.454545
## y = 3.818182
```