Q_5

Exercise 18 on page 571.

```
\mathbf{a}
```

```
code:
```

```
m1 <- c(13, 17, 7, 14);
                              h1 <- rep("1", length(m1));
                              h2 <- rep("2", length(m2));
 m2 < -c(21, 13, 20, 17);
                              h3 <- rep("3", length(m3));
 m3 <- c(18, 15, 20, 17);
 m4 <- c( 7, 11, 18, 10);
                              h4 <- rep("4", length(m4));
                               h5 <- rep("5", length(m5));
 m5 < -c(6, 11, 15, 8);
 measure \leftarrow c(d1, d2, d3, d4, d5)
 hormone <- c(h1, h2, h3, h4, h5)
  vals <- data.frame(measure, hormone)</pre>
  anova_result <- anova(lm(measure~ hormone, data=vals))</pre>
  print(anova_result)
output:
  Response: measure
            Df Sum Sq Mean Sq F value Pr(>F)
             4 200.3 50.075 3.4855 0.03336
 hormone
  Residuals 15
               215.5 14.367
```

The F value is 3.4855 which which gives a P value of 0.03336, so the null hypothesis can be rejected.

b)

```
\bar{x}_1 = 12.75
\bar{x}_2 = 17.75
\bar{x}_3 = 17.5
\bar{x}_4 = 11.5
\bar{x}_5 = 10
w = Q_{\alpha,I,I(J-1)} = Q_{0.05,5,5\cdot(4-1)}\sqrt{14.367/4} = 8.276
```

Q5 b) contined from previous page

 $\frac{1}{10}$ $\frac{1}{10}$

△x5, x4 = 1,5 ew △x5, x1 = 2,75 ew △x5, x3 = 7,5 ew △x5, x2 = 7,75 ew

Using to Key procedure, none of the sample means differ signifigantly.