Exercises 9 - SOLUTIONS

1.

(a) Blocks form a 'restriction on the randomization': thus we need to apply each dye to a particular brand of denim in random order; however, the order in which each brand of denim is considered can be taken arbitrarily.

```
(b) > strength <- c(60, 55, 61, 58, 54, 60, 54, 62, 59, 57, 62, 55, 65, 60, 55,
     60, 58, 62, 62, 56)
 > d <- rep(1:4, rep(5, 4))
> b <- rep(c("A", "B", "C", "D", "E"), 4)
> dye <- factor(d)
> brand <- factor(b)</pre>
> denim <- data.frame(strength, dye, brand)
> denim.aov <- aov(strength ~ dye + brand, data = denim)
> summary(denim.aov)
             Df Sum Sq Mean Sq F value
                                         Pr(>F)
 dye
              3 12.95
                        4.32
                                2.376
                                          0.121
              4 157.00
                         39.25
                                21.606 2.06e-05 ***
 brand
Residuals
             12 21.80
                          1.82
Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1
```

The relevant F-statistic of 2.376 is not very significant on the $F_{3,12}$ distribution: indeed the p-value is 0.121. Conclusion: no evidence that a difference exists in the effects of the 4 dyes.

2.

(a) The use of 4 different cows in the experiment may contribute in some way to the overall variability in the infectivity readings. We can take this into account by taking the cows as blocks.

```
(b) > infect <- c(20, 29, 25, 46, 23, 31, 24, 51, 12, 11, 8, 29)
 > t \leftarrow rep(c("T1", "T2", "T3"), rep(4, 3))
 > c <- rep(1:4, 3)
> test <- factor(t)
> cow <- factor(c)
> bse <- data.frame(infect, test, cow)
> bse.aov <- aov(infect ~ test + cow, data = bse)
> summary(bse.aov)
             Df Sum Sq Mean Sq F value
                                          Pr(>F)
              2 703.5
                         351.7
                                  40.72 0.000323 ***
 test
              3 1106.9
                         369.0
                                  42.71 0.000192 ***
 COW
Residuals
              6
                  51.8
                           8.6
Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1
```

The relevant F-statistic is 40.72, which is significant on the $F_{2,6}$ distribution, as evidenced by the p-value of 0.000323. Thus, there are differences in the amounts of infectivity that the 3 tests can detect, i.e. they are **not** equally effective.