Exercises Day 1

Your name

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Exercise 1

Suppose you are running a behavioural experiment, studying the mental speed of young and old people. You know that young people have an average reaction time of 550 with a standart deviation of 120.

a) What is the probability of getting a reaction time of 420 or lower from a single young person?

```
pnorm(420,mean=550,sd=120)
```

[1] 0.1393302

b) What is the probability of getting a reaction time of 560 or higher from a single young person?

```
1-pnorm(560, mean=550, sd=120)
```

[1] 0.4667932

c) For your study, you need 10 reaction time values for young people. How can you generate a sample with n=10?

```
x1 <- rnorm(10, mean=550, sd=120)
```

d) You get 15 values from a population of old people. What is the mean and the standart deviation of their underlying normal distribution?

```
x2 \leftarrow c(434.7, 671.4, 428.9, 454.4, 806.1, 483.3, 819.1, 630.4, 836.2, 661.4, 511.7, 507.2, 568.0, 707.5
mean(x2)
```

[1] 609.4933

sd(x2)

[1] 140.3252

e) What is the probability that the 15 values are actually drawn from the same distribution as in 1c

t.test(x2, mu=550)

```
##
## One Sample t-test
##
## data: x2
## t = 1.642, df = 14, p-value = 0.1228
## alternative hypothesis: true mean is not equal to 550
## 95 percent confidence interval:
## 531.7838 687.2028
## sample estimates:
## mean of x
## 609.4933
```

Exercise 2

You are a researcher studying the intelligence of dragons in the european mountains. You suspect that larger dragons are also smarter and you've collected various samples for intelligence (testscore) and size (bodylength) from different mountains.

a) Is there a significant relationship between intelligence and body size in dragons? Use linear regression.

```
load("dragons.RData")

Model_lm <- lm(testScore ~ bodyLength, data = dragons)
summary(Model_lm)</pre>
```

```
##
## Call:
## lm(formula = testScore ~ bodyLength, data = dragons)
##
## Residuals:
##
      Min
               1Q Median
                               3Q
                                      Max
## -68.070 -15.792 -0.852 16.786
                                   63.368
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) -63.10331
                           13.18621
                                    -4.786 2.28e-06 ***
                            0.06529
                                     9.377 < 2e-16 ***
## bodyLength
                0.61224
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 23.17 on 478 degrees of freedom
## Multiple R-squared: 0.1554, Adjusted R-squared: 0.1536
## F-statistic: 87.94 on 1 and 478 DF, p-value: < 2.2e-16
```

b) Maybe the location of each recording sample influences the results. Create a LME that accounts for differences between mountain ranges. What is the relationship between intelligence and body size now?

```
Model_Mountains <- lmer(testScore ~ bodyLength + (1|mountainRange), data = dragons)
summary(Model_Mountains)</pre>
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: testScore ~ bodyLength + (1 | mountainRange)
##
      Data: dragons
##
## REML criterion at convergence: 4187.3
##
## Scaled residuals:
       Min
                  10
                       Median
                                    30
## -2.69490 -0.68646 -0.01821 0.67354 2.64609
##
## Random effects:
## Groups
                  Name
                              Variance Std.Dev.
                                       17.07
## mountainRange (Intercept) 291.2
## Residual
                              340.1
                                       18.44
## Number of obs: 480, groups: mountainRange, 8
##
## Fixed effects:
##
               Estimate Std. Error t value
## (Intercept) 31.06599
                          20.25713
                                     1.534
## bodyLength
              0.14447
                           0.09597
                                     1.505
##
## Correlation of Fixed Effects:
              (Intr)
## bodyLength -0.954
```

c) You notice that dragons with different colors behave differently. Control in your LME for the color of each dragon. How do the results change?

```
Model_Color <- lmer(testScore ~ bodyLength + Color + (1|mountainRange), data = dragons)
summary(Model_Color)</pre>
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: testScore ~ bodyLength + Color + (1 | mountainRange)
##
      Data: dragons
## REML criterion at convergence: 4014.8
##
## Scaled residuals:
##
      Min
                1Q Median
                                3Q
                                       Max
## -3.6979 -0.6275 0.0068 0.6859 2.6846
## Random effects:
## Groups
                              Variance Std.Dev.
                  Name
## mountainRange (Intercept) 337.4
                                       18.37
                                       15.40
## Residual
                              237.1
## Number of obs: 480, groups: mountainRange, 8
##
## Fixed effects:
##
              Estimate Std. Error t value
```

```
## (Intercept) 34.80775 17.54353 1.984
## bodyLength 0.07632 0.08097 0.943
## ColorRed 20.38107 1.42452 14.307
##
## Correlation of Fixed Effects:
## (Intr) bdyLng
## bodyLength -0.927
## ColorRed 0.005 -0.048
```