Assignment 4 STAT 315-463: Multivariable Statistical Methods and Applications

```
library(here)
library(lattice)
library(lme4)
library(ggplot2)
library(tidyverse)
# Read in data file
tern14 <- read.table("Terns2014.csv", header = TRUE, sep = ',', na.strings = "na")
# Only using simple Linear Regression
Model <- lm(Age ~ Wing * ID, data = tern14)</pre>
summary(Model)
##
## lm(formula = Age ~ Wing * ID, data = tern14)
## Residuals:
               1Q Median
                               3Q
                                      Max
## -3.6826 -1.6299 -0.5009 1.5109 6.3868
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 2.6194915 0.4952646 5.289 2.06e-07 ***
## Wing
              0.1530470 0.0050336 30.405 < 2e-16 ***
## ID
              0.0101452 0.0159284 0.637
                                               0.525
## Wing:ID
             -0.0002035 0.0001494 -1.363
                                               0.174
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 2.094 on 387 degrees of freedom
## Multiple R-squared: 0.8963, Adjusted R-squared: 0.8955
## F-statistic: 1115 on 3 and 387 DF, p-value: < 2.2e-16
AIC(Model)
## [1] 1693.598
```

Model 1: Random intercepts

```
Model.1 <- lmer(Age ~ Wing + (1|ID), data = tern14)
AIC(Model.1)</pre>
```

```
## [1] 1143.745
```

In Model 1, we are looking at the model with the random intercepts, which assume that some terns have more and some have less wing length.

The AIC value of this mixed linear model indicates that it fits better than the model that ignores individual effects (The AIC score of random intercepts model is smaller than the simple linear regression model).

```
summary(Model.1)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: Age ~ Wing + (1 | ID)
##
      Data: tern14
##
## REML criterion at convergence: 1135.7
##
## Scaled residuals:
##
       Min
                1Q Median
                                        Max
  -3.4005 -0.4709 -0.0712 0.4367 5.6895
##
##
## Random effects:
    Groups
                         Variance Std.Dev.
                                   2.0669
##
             (Intercept) 4.2721
                         0.5619
                                   0.7496
    Residual
## Number of obs: 391, groups: ID, 65
##
## Fixed effects:
##
               Estimate Std. Error t value
## (Intercept) 2.976157
                          0.283128
                                      10.51
## Wing
               0.147138
                          0.001146
                                    128.34
##
## Correlation of Fixed Effects:
##
        (Intr)
## Wing -0.395
```

Random Effects: Around 88.4% of variability is explained by the random intercepts. The variance of residual shows that there are still some factors from the simple body measurements of terns which affect age and cannot be explained by the model. This value is pretty close to zero, which

Fixed Effects: The intercept in the summary above shows that the average tern's wing length when they are born is 2.97 mm and the slope suggests that in average, each increase in wing length by 1 mm takes 0.14 days.

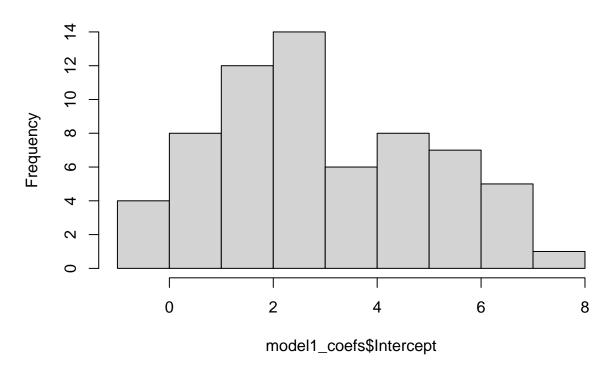
```
model1_coefs <- coef(Model.1)$ID %>%
  rename(Intercept = '(Intercept)', Slope = Wing) %>%
  rownames_to_column("ID")

# See coefficients
summary(model1_coefs$Intercept)
```

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## -0.4788 1.3941 2.6807 2.9762 4.6634 7.3462
```

hist(model1_coefs\$Intercept)

Histogram of model1_coefs\$Intercept



Model 2: Random slopes

In this case, we are expecting all terns to start off at around the same point, but the effects of wing length on age differs across each tern.

```
Model.2 <- lmer(Age ~ Wing + (0 + Wing|ID), data = tern14)
AIC(Model.2)</pre>
```

[1] 1174.457

The AIC value suggests this model outperforms the simple linear regression model, which does not take the different growth rate into consideration.

```
summary(Model.2)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: Age ~ Wing + (0 + Wing | ID)
## Data: tern14
```

```
## REML criterion at convergence: 1166.5
## Scaled residuals:
      Min 1Q Median
                               3Q
## -3.0715 -0.4559 -0.0779 0.4010 4.2163
## Random effects:
## Groups Name Variance Std.Dev.
            Wing 0.0005676 0.02382
## Residual
               0.5962503 0.77217
## Number of obs: 391, groups: ID, 65
## Fixed effects:
              Estimate Std. Error t value
## (Intercept) 2.527025 0.122949
                                    20.55
## Wing
              0.153688
                       0.003315
                                   46.35
##
## Correlation of Fixed Effects:
       (Intr)
## Wing -0.405
```

Random Effects:

Fixed Effects:

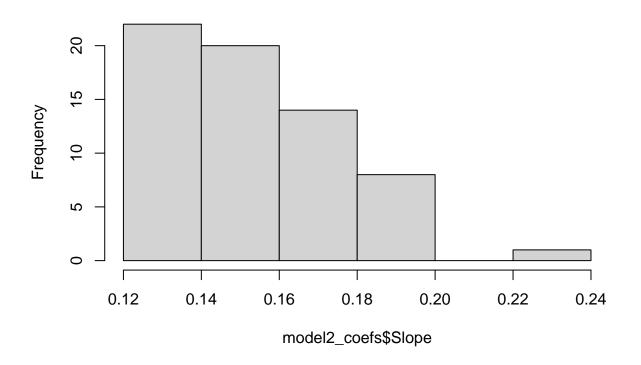
```
model2_coefs <- coef(Model.2)$ID %>%
  rename(Intercept = '(Intercept)', Slope=Wing) %>%
  rownames_to_column("ID")

summary(model2_coefs)
```

```
##
                      Intercept
                                      Slope
        ID
## Length:65
                    Min. :2.527
                                   Min. :0.1208
## Class:character 1st Qu.:2.527
                                   1st Qu.:0.1370
## Mode :character Median :2.527
                                   Median :0.1486
##
                         :2.527
                                        :0.1537
                    Mean
                                   Mean
##
                    3rd Qu.:2.527
                                   3rd Qu.:0.1710
##
                    Max. :2.527
                                   Max. :0.2388
```

hist(model2_coefs\$Slope)

Histogram of model2_coefs\$Slope



coef(Model.2)

```
## $ID
      (Intercept)
##
                        Wing
## 1
         2.527025 0.1296118
## 2
         2.527025 0.1266603
## 3
         2.527025 0.1441504
## 4
         2.527025 0.1283708
## 5
         2.527025 0.1664200
## 6
         2.527025 0.1352117
## 7
         2.527025 0.1397402
## 8
         2.527025 0.1387303
## 9
         2.527025 0.1816841
## 10
         2.527025 0.1647057
## 11
         2.527025 0.1910185
## 12
         2.527025 0.1718231
## 13
         2.527025 0.1725178
         2.527025 0.1304463
## 14
## 15
         2.527025 0.1707286
## 16
         2.527025 0.1621836
## 17
         2.527025 0.1238346
## 18
         2.527025 0.1445179
## 19
         2.527025 0.1577769
## 20
         2.527025 0.1469140
## 21
         2.527025 0.1873774
         2.527025 0.1885235
## 22
```

```
## 23
         2.527025 0.1388217
## 24
         2.527025 0.1799855
         2.527025 0.1739090
## 25
## 26
         2.527025 0.2388141
## 27
         2.527025 0.1556988
## 28
         2.527025 0.1982779
## 29
         2.527025 0.1408648
## 30
         2.527025 0.1738029
## 31
         2.527025 0.1462256
## 32
         2.527025 0.1276246
## 33
         2.527025 0.1529132
         2.527025 0.1370442
## 34
## 35
         2.527025 0.1508789
## 36
         2.527025 0.1676289
## 37
         2.527025 0.1709695
## 38
         2.527025 0.1383305
## 39
         2.527025 0.1391249
## 40
         2.527025 0.1517927
## 41
         2.527025 0.1485587
## 42
         2.527025 0.1720056
## 43
         2.527025 0.1464499
## 44
         2.527025 0.1618591
         2.527025 0.1821830
## 45
         2.527025 0.1417453
## 46
## 47
         2.527025 0.1576291
## 48
         2.527025 0.1300611
## 49
         2.527025 0.1767589
## 50
         2.527025 0.1207979
## 51
         2.527025 0.1833605
## 52
         2.527025 0.1438838
## 53
         2.527025 0.1307264
## 54
         2.527025 0.1551169
## 55
         2.527025 0.1451454
         2.527025 0.1361845
## 56
## 57
         2.527025 0.1235031
## 58
         2.527025 0.1246739
## 59
         2.527025 0.1955957
## 60
         2.527025 0.1401833
## 61
         2.527025 0.1303502
         2.527025 0.1509885
## 62
         2.527025 0.1273769
## 63
## 64
         2.527025 0.1520855
         2.527025 0.1268382
## 65
##
## attr(,"class")
## [1] "coef.mer"
```

Model 3: Random slopes and intercepts

```
Model.3 <- lmer(Age ~ Wing + (1 + Wing|ID), data = tern14)
## Warning in checkConv(attr(opt, "derivs"), opt$par, ctrl = control$checkConv, :</pre>
```

```
## Model failed to converge with max|grad| = 0.439389 (tol = 0.002, component 1)
```

```
summary(Model.3)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: Age ~ Wing + (1 + Wing | ID)
##
      Data: tern14
##
## REML criterion at convergence: 932.9
##
## Scaled residuals:
##
       Min
                1Q Median
                                3Q
                                       Max
  -3.2764 -0.4639 -0.0748 0.4586
##
                                    4.9026
##
## Random effects:
##
   Groups
                         Variance Std.Dev. Corr
##
             (Intercept) 3.4346907 1.85329
                         0.0003303 0.01817
             Wing
                                             -0.38
## Residual
                         0.2256019 0.47498
## Number of obs: 391, groups: ID, 65
##
## Fixed effects:
##
               Estimate Std. Error t value
## (Intercept) 2.697838
                          0.262709
                                     10.27
                                      56.54
## Wing
               0.150908
                          0.002669
##
## Correlation of Fixed Effects:
##
        (Intr)
## Wing -0.510
## optimizer (nloptwrap) convergence code: 0 (OK)
## Model failed to converge with max|grad| = 0.439389 (tol = 0.002, component 1)
```

AIC(Model.3)

[1] 944.8728

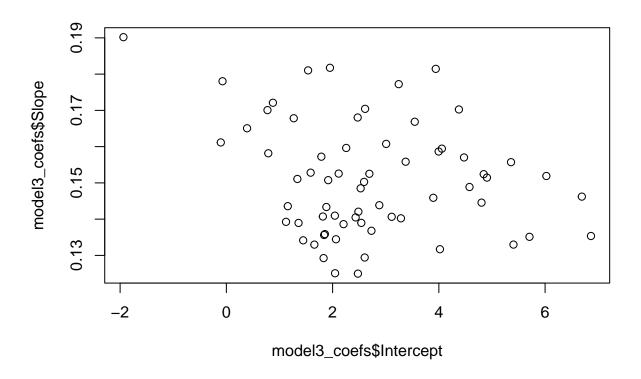
From the AIC result above, we can see that this model with random slopes and random intercepts has the best fit (the smallest AIC value among all four models). It does make sense because each fern has different growth rate and their wing lengths also differ when they were just born.

Random Effects:

Fixed Effects:

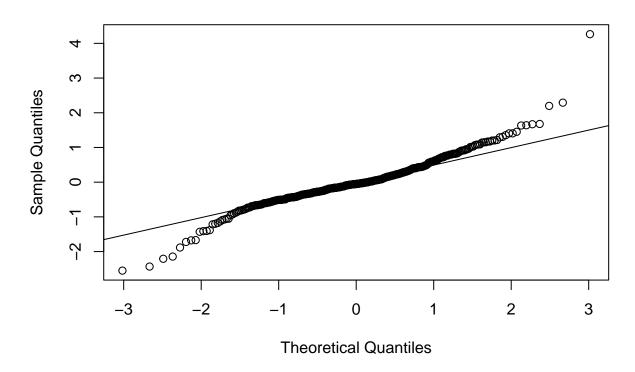
```
model3_coefs <- coef(Model.3)$ID %>%
    rename(Intercept = '(Intercept)', Slope=Wing) %>%
    rownames_to_column("ID")

plot(model3_coefs$Intercept, model3_coefs$Slope)
```



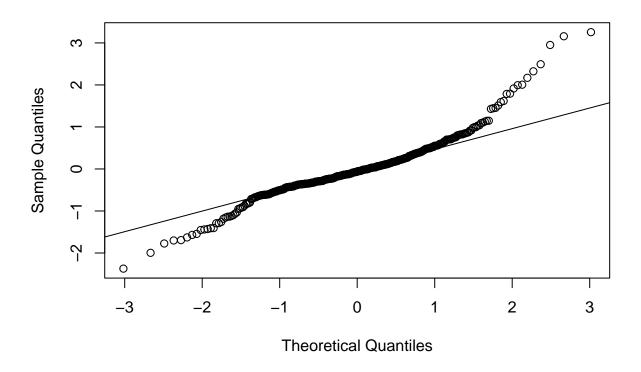
```
qqnorm(resid(Model.1))
qqline(resid(Model.1))
```

Normal Q-Q Plot



```
qqnorm(resid(Model.2))
qqline(resid(Model.2))
```

Normal Q-Q Plot



```
qqnorm(resid(Model.3))
qqline(resid(Model.3))
```

Normal Q-Q Plot

