

## Mixed models to determine the relationship between urbanization and turtle behaviour

R code to determine the relationship between urbanization and turtle behaviour (active defensive behaviours (Aggression), shell emergence time (Shell), time of initial movement (Start), total time spent moving (Move)).

### Packages

```
library(Hmisc)
library(writexl)
library(ggplot2)
library(dplyr)
library(lmerTest)
library(optimx)
library(lme4)
library(PerformanceAnalytics)
library(effects)
library(ggeffects)
library(splines)
library(glmtoolbox)
library(afex)
library(nloptr)
library(dfoptim)
library(psych)
library(ordinal)
library(ggpubr)
library(terra)
library(AICcmodavg)
library(visreg)
library(MuMIn)
```

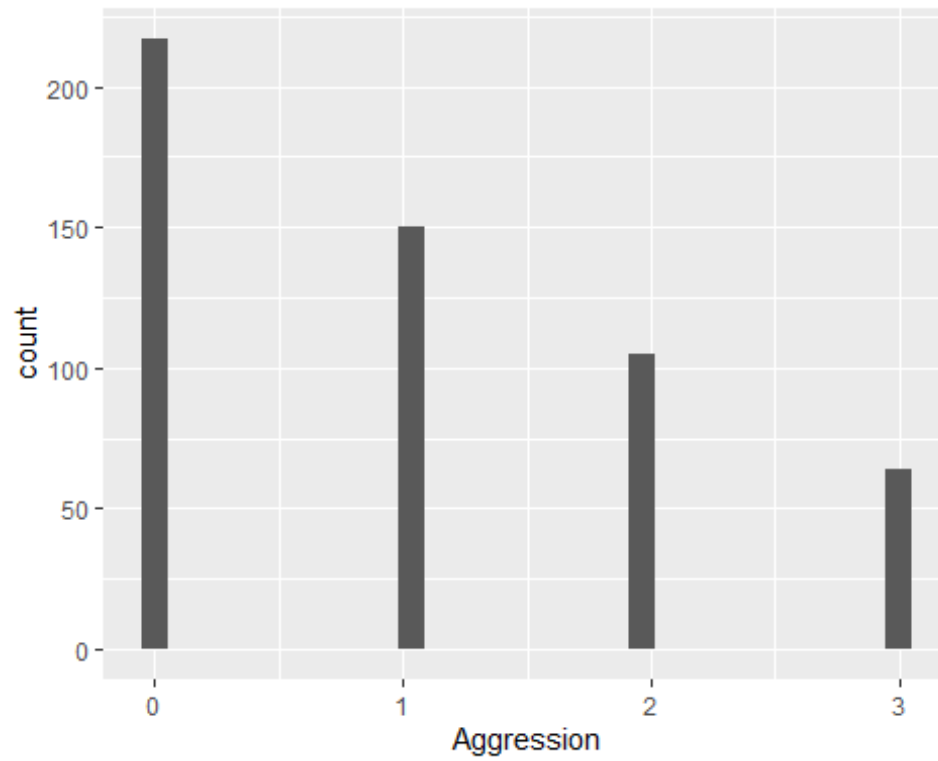
### Upload the dataset to use

```
MixedData<-read.csv("C:/Users/sebas/Desktop/Masters Work/Masters Work/Stats/B  
in.Shell.600/Mixed Model Correlation Data.600.csv")
```

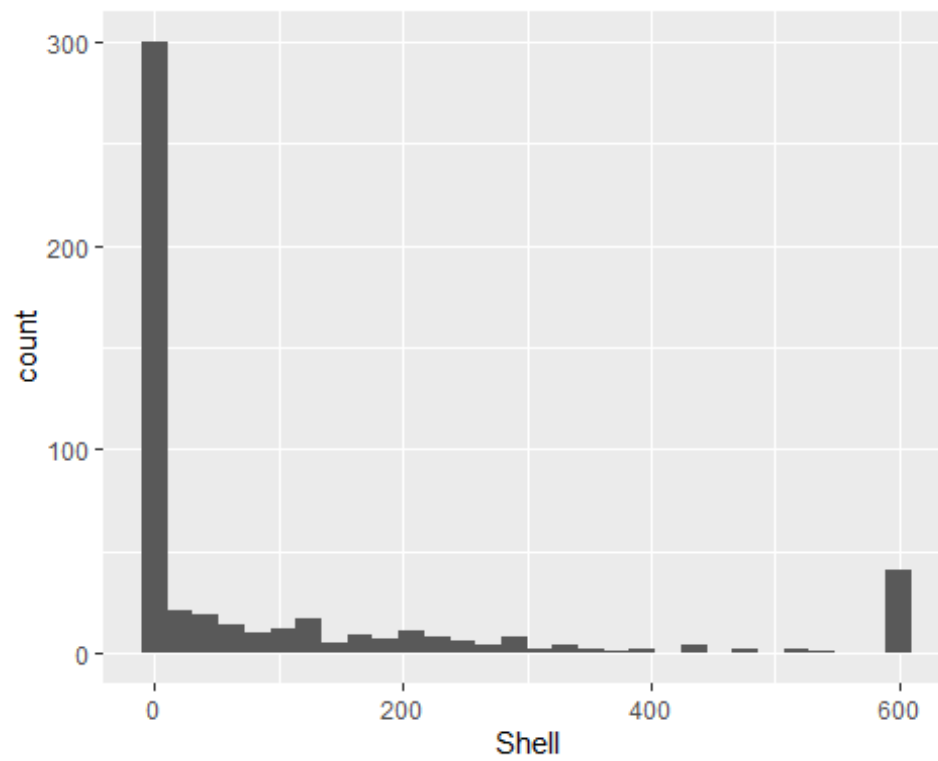
### Plot Data

I am doing this step to make sure I use the right data distribution for my models later on. If I do need to transform any of the data distributions, I will need to recalculate the distance at which each land cover class has a maximum impact on the transformed behaviour.

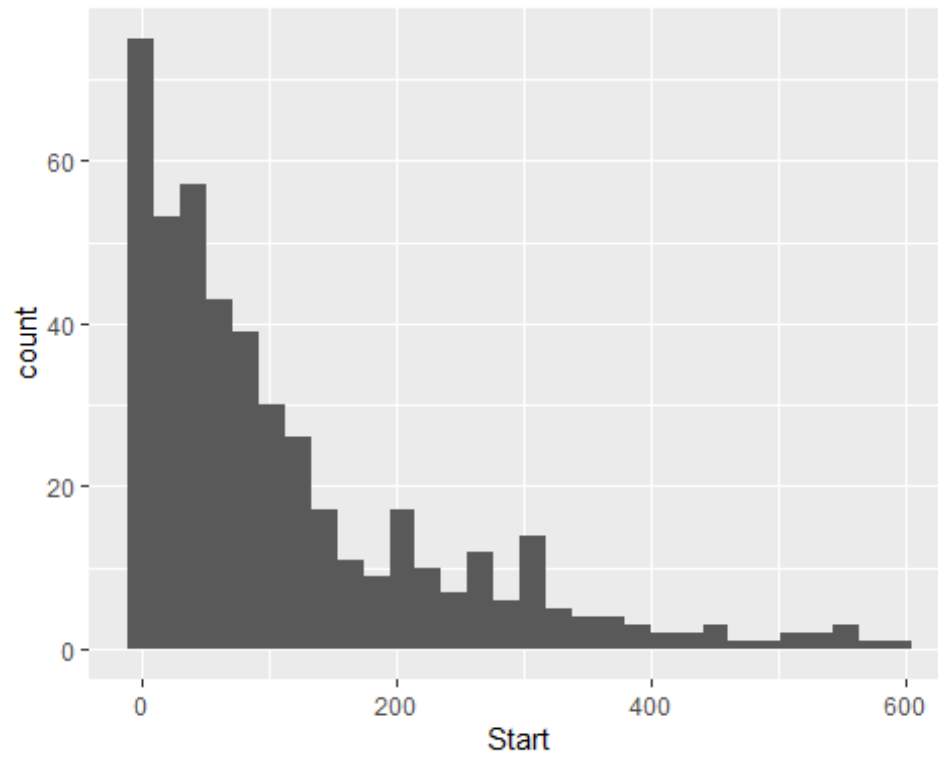
```
ggplot(MixedData, aes(x=Aggression)) + geom_histogram()
```



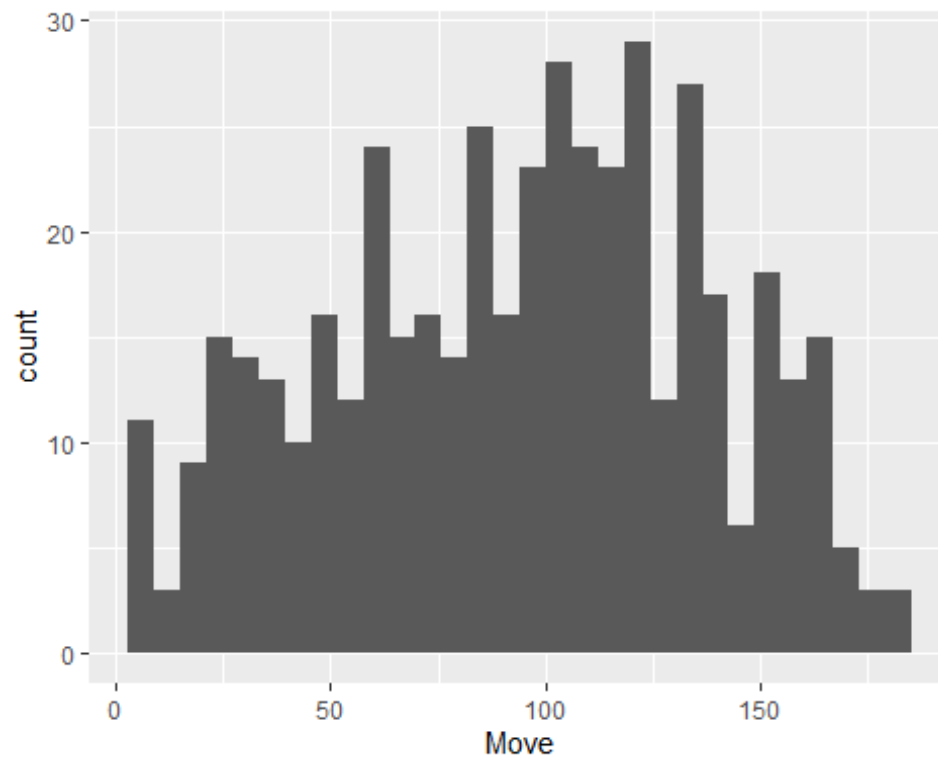
```
ggplot(MixedData, aes(x=Shell)) + geom_histogram()
```



```
ggplot(MixedData, aes(x=Start)) + geom_histogram()
```



```
ggplot(MixedData, aes(x=Move)) + geom_histogram()
```



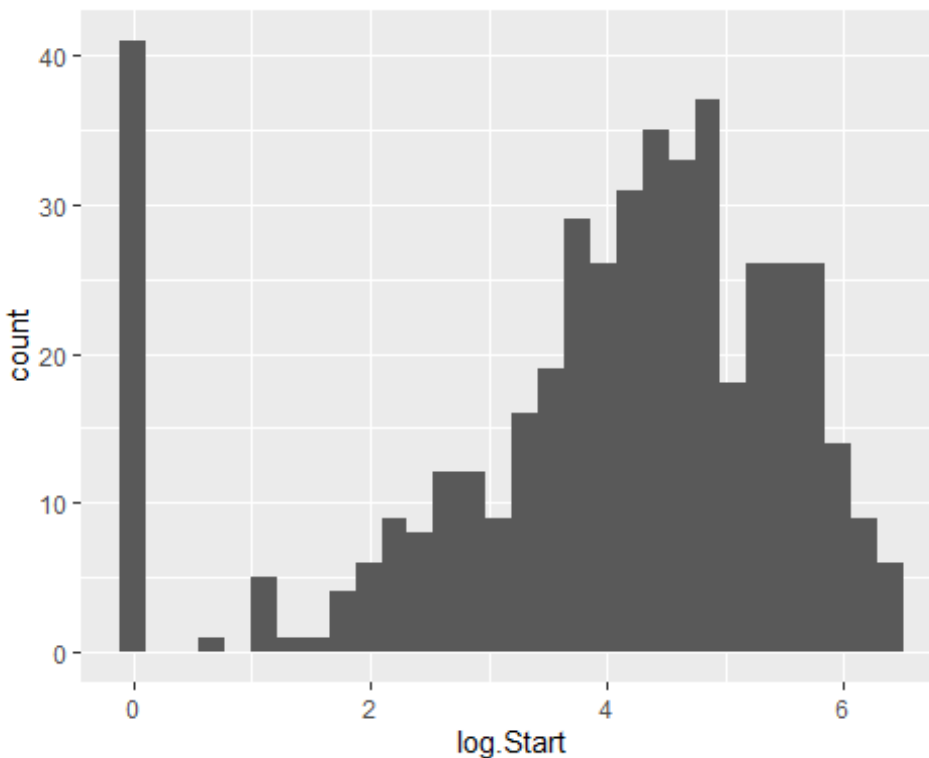
Time of initial movement has a skewed distribution so I will plot a log transformed version of it.

### Log(x+1) transform time of initial movement (log.Start)

```
MixedData$log.Start <- log(MixedData$Start + 1)
```

### Plot log(x+1) transformed version of time of initial movement

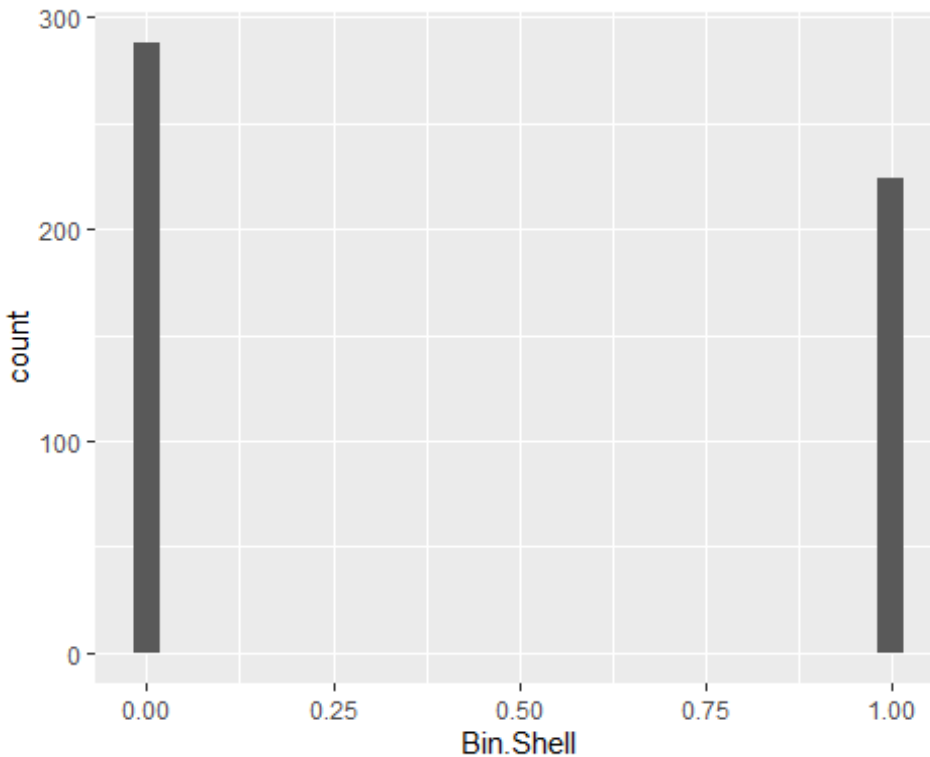
```
ggplot(MixedData, aes(x=log.Start)) + geom_histogram()
```



The log transformed version of time of initial movement is still somewhat skewed, so I will run my analysis on both versions for comparison. Shell emergence time is also skewed, so I will use a binary version of it instead, where “0” means a turtle emerged from its shell at 0 seconds, and “1” means a turtle did not emerge from its shell at 0 seconds.

### Plot binary version of shell emergence time data

```
ggplot(MixedData, aes(x=Bin.Shell)) + geom_histogram()
```



## Preparing Data

### *# Response variables*

```
MixedData$Aggression <- as.numeric(MixedData$Aggression)
MixedData$Bin.Shell <- as.numeric(MixedData$Bin.Shell)
MixedData$Start <- as.numeric(MixedData$Start)
MixedData$log.Start <- as.numeric(MixedData$log.Start)
MixedData$Move <- as.numeric(MixedData$Move)
```

### *# Predictor variables*

```
MixedData$Site <- as.factor(MixedData$Site)
MixedData$Sex <- as.factor(MixedData$Sex)
MixedData$Code <- as.factor(MixedData$Code)
MixedData$A.Temp <- as.numeric(MixedData$A.Temp)
MixedData$W.Temp <- as.numeric(MixedData$W.Temp)
MixedData$Time <- as.numeric(MixedData$Time)
MixedData$Mass <- as.numeric(MixedData$Mass)
MixedData$PL <- as.numeric(MixedData$PL)
MixedData$JulianDate <- as.numeric(MixedData$JulianDate)
MixedData$for.veg.200 <- as.numeric(MixedData$for.veg.200)
MixedData$for.veg.300 <- as.numeric(MixedData$for.veg.300)
MixedData$for.veg.1000 <- as.numeric(MixedData$for.veg.1000)
MixedData$wet.200 <- as.numeric(MixedData$wet.200)
MixedData$wet.300 <- as.numeric(MixedData$wet.300)
```

```

MixedData$wet.400 <- as.numeric(MixedData$wet.400)
MixedData$wet.600 <- as.numeric(MixedData$wet.600)
MixedData$wet.1000 <- as.numeric(MixedData$wet.1000)
MixedData$agri.100 <- as.numeric(MixedData$agri.100)
MixedData$agri.200 <- as.numeric(MixedData$agri.200)
MixedData$agri.400 <- as.numeric(MixedData$agri.400)
MixedData$agri.500 <- as.numeric(MixedData$agri.500)
MixedData$agri.600 <- as.numeric(MixedData$agri.600)
MixedData$agri.1000 <- as.numeric(MixedData$agri.1000)
MixedData$urban.1000 <- as.numeric(MixedData$urban.1000)
MixedData$water.100 <- as.numeric(MixedData$water.100)
MixedData$water.400 <- as.numeric(MixedData$water.400)
MixedData$water.500 <- as.numeric(MixedData$water.500)
MixedData$water.900 <- as.numeric(MixedData$water.900)
MixedData$water.1000 <- as.numeric(MixedData$water.1000)

```

## Standardization of the predictor variables

**### ALL continuous numeric predictor variables were standardized (mean zero, unit variance) before model selection.**

```

MixedData$A.Temp.Scaled <- scale(MixedData$A.Temp, center=TRUE, scale=TRUE)
MixedData$W.Temp.Scaled <- scale(MixedData$W.Temp, center=TRUE, scale=TRUE)
MixedData$Time.Scaled <- scale(MixedData$Time, center=TRUE, scale=TRUE)
MixedData$Mass.Scaled <- scale(MixedData$Mass, center=TRUE, scale=TRUE)
MixedData$PL.Scaled <- scale(MixedData$PL, center=TRUE, scale=TRUE)
MixedData$JulianDate.Scaled <- scale(MixedData$JulianDate, center=TRUE, scale=TRUE)
MixedData$for.veg.200.Scaled <- scale(MixedData$for.veg.200, center=TRUE, scale=TRUE)
MixedData$for.veg.300.Scaled <- scale(MixedData$for.veg.300, center=TRUE, scale=TRUE)
MixedData$for.veg.1000.Scaled <- scale(MixedData$for.veg.1000, center=TRUE, scale=TRUE)
MixedData$wet.200.Scaled <- scale(MixedData$wet.200, center=TRUE, scale=TRUE)
MixedData$wet.300.Scaled <- scale(MixedData$wet.300, center=TRUE, scale=TRUE)
MixedData$wet.400.Scaled <- scale(MixedData$wet.400, center=TRUE, scale=TRUE)
MixedData$wet.600.Scaled <- scale(MixedData$wet.600, center=TRUE, scale=TRUE)
MixedData$wet.1000.Scaled <- scale(MixedData$wet.1000, center=TRUE, scale=TRUE)
MixedData$agri.100.Scaled <- scale(MixedData$agri.100, center=TRUE, scale=TRUE)
MixedData$agri.200.Scaled <- scale(MixedData$agri.200, center=TRUE, scale=TRUE)
MixedData$agri.400.Scaled <- scale(MixedData$agri.400, center=TRUE, scale=TRUE)
MixedData$agri.500.Scaled <- scale(MixedData$agri.500, center=TRUE, scale=TRUE)
MixedData$agri.600.Scaled <- scale(MixedData$agri.600, center=TRUE, scale=TRUE)

```

```

E)
MixedData$agri.1000.Scaled <- scale(MixedData$agri.1000, center=TRUE, scale=T
RUE)
MixedData$urban.1000.Scaled <- scale(MixedData$urban.1000, center=TRUE, scale
=TRUE)
MixedData$water.100.Scaled <- scale(MixedData$water.100, center=TRUE, scale=T
RUE)
MixedData$water.400.Scaled <- scale(MixedData$water.400, center=TRUE, scale=T
RUE)
MixedData$water.500.Scaled <- scale(MixedData$water.500, center=TRUE, scale=T
RUE)
MixedData$water.900.Scaled <- scale(MixedData$water.900, center=TRUE, scale=T
RUE)
MixedData$water.1000.Scaled <- scale(MixedData$water.1000, center=TRUE, scale
=TRUE)

```

## Make scaled variables numeric

```

MixedData$A.Temp.Scaled <- as.numeric(MixedData$A.Temp.Scaled)
MixedData$W.Temp.Scaled <- as.numeric(MixedData$W.Temp.Scaled)
MixedData$Time.Scaled <- as.numeric(MixedData$Time.Scaled)
MixedData$Mass.Scaled <- as.numeric(MixedData$Mass.Scaled)
MixedData$PL.Scaled <- as.numeric(MixedData$PL.Scaled)
MixedData$JulianDate.Scaled <- as.numeric(MixedData$JulianDate.Scaled)
MixedData$for.veg.200.Scaled <- as.numeric(MixedData$for.veg.200.Scaled)
MixedData$for.veg.300.Scaled <- as.numeric(MixedData$for.veg.300.Scaled)
MixedData$for.veg.1000.Scaled <- as.numeric(MixedData$for.veg.1000.Scaled)
MixedData$wet.200.Scaled <- as.numeric(MixedData$wet.200.Scaled)
MixedData$wet.300.Scaled <- as.numeric(MixedData$wet.300.Scaled)
MixedData$wet.400.Scaled <- as.numeric(MixedData$wet.400.Scaled)
MixedData$wet.600.Scaled <- as.numeric(MixedData$wet.600.Scaled)
MixedData$wet.1000.Scaled <- as.numeric(MixedData$wet.1000.Scaled)
MixedData$agri.100.Scaled <- as.numeric(MixedData$agri.100.Scaled)
MixedData$agri.200.Scaled <- as.numeric(MixedData$agri.200.Scaled)
MixedData$agri.400.Scaled <- as.numeric(MixedData$agri.400.Scaled)
MixedData$agri.500.Scaled <- as.numeric(MixedData$agri.500.Scaled)
MixedData$agri.600.Scaled <- as.numeric(MixedData$agri.600.Scaled)
MixedData$agri.1000.Scaled <- as.numeric(MixedData$agri.1000.Scaled)
MixedData$urban.1000.Scaled <- as.numeric(MixedData$urban.1000.Scaled)
MixedData$water.100.Scaled <- as.numeric(MixedData$water.100.Scaled)
MixedData$water.400.Scaled <- as.numeric(MixedData$water.400.Scaled)
MixedData$water.500.Scaled <- as.numeric(MixedData$water.500.Scaled)
MixedData$water.900.Scaled <- as.numeric(MixedData$water.900.Scaled)
MixedData$water.1000.Scaled <- as.numeric(MixedData$water.1000.Scaled)

```

## Verification of model assumptions

### Active defensive behaviours (Aggression)

#### Multicollinearity: Generalized variance inflation factor ( $\text{GVIF}^{1/(2 \cdot \text{df})}$ )

$\text{GVIF}^{1/(2 \cdot \text{df})} > 2$  indicates the presence of multicollinearity, so I will remove variables with values over 2, starting with the highest value.

```
mod.Aggression.vif <- lm(Aggression ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled + Time.Scaled + Mass.Scaled + PL.Scaled + Sex + for.veg.1000.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.Aggression.vif)
```

	GVIF	df	$\text{GVIF}^{1/(2 \cdot \text{df})}$
## JulianDate.Scaled	3.0926	1	1.7586
## A.Temp.Scaled	3.8408	1	1.9598
## W.Temp.Scaled	5.7294	1	2.3936
## Time.Scaled	1.1397	1	1.0676
## Mass.Scaled	9.0761	1	3.0127
## PL.Scaled	8.2574	1	2.8736
## Sex	1.5100	1	1.2288
## for.veg.1000.Scaled	5.9695	1	2.4433
## wet.200.Scaled	1.3717	1	1.1712
## agri.100.Scaled	1.7661	1	1.3289
## urban.1000.Scaled	4.3219	1	2.0789
## water.400.Scaled	2.5555	1	1.5986

Turtle mass (Mass) has the highest  $\text{GVIF}^{1/(2 \cdot \text{df})} > 2$ , so I will remove it and recalculate the factors.

```
mod.Aggression.vif <- lm(Aggression ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.1000.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.Aggression.vif)
```

	GVIF	df	$\text{GVIF}^{1/(2 \cdot \text{df})}$
## JulianDate.Scaled	3.0921	1	1.7584
## A.Temp.Scaled	3.7862	1	1.9458
## W.Temp.Scaled	5.7100	1	2.3896
## Time.Scaled	1.1248	1	1.0606
## PL.Scaled	1.4024	1	1.1842
## Sex	1.4118	1	1.1882
## for.veg.1000.Scaled	5.9694	1	2.4432
## wet.200.Scaled	1.3707	1	1.1708
## agri.100.Scaled	1.7654	1	1.3287
## urban.1000.Scaled	4.2665	1	2.0656
## water.400.Scaled	2.5162	1	1.5863



Proportion of forest and vegetation area at 1000m (for.veg.1000) has the highest  $GVIF^{(1/(2*df))} > 2$ , so I will remove it and recalculate the factors.

```
mod.Aggression.vif <- lm(Aggression ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.Aggression.vif)

##              GVIF df GVIF^(1/(2*df))
## JulianDate.Scaled 2.7130 1      1.6471
## A.Temp.Scaled     3.7182 1      1.9283
## W.Temp.Scaled     5.6994 1      2.3873
## Time.Scaled       1.1062 1      1.0518
## PL.Scaled         1.3879 1      1.1781
## Sex               1.4090 1      1.1870
## wet.200.Scaled    1.3592 1      1.1658
## agri.100.Scaled   1.2301 1      1.1091
## urban.1000.Scaled 1.2841 1      1.1332
## water.400.Scaled  1.3704 1      1.1706
```

Water temperature (W.Temp) has the highest  $GVIF^{(1/(2*df))} > 2$ , so I will remove it and recalculate the factors.

```
mod.Aggression.vif <- lm(Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.Aggression.vif)

##              GVIF df GVIF^(1/(2*df))
## JulianDate.Scaled 1.7005 1      1.3040
## A.Temp.Scaled     1.5191 1      1.2325
## Time.Scaled       1.0828 1      1.0406
## PL.Scaled         1.4046 1      1.1852
## Sex               1.4112 1      1.1879
## wet.200.Scaled    1.3285 1      1.1526
## agri.100.Scaled   1.1437 1      1.0694
## urban.1000.Scaled 1.2163 1      1.1029
## water.400.Scaled  1.1445 1      1.0698
```

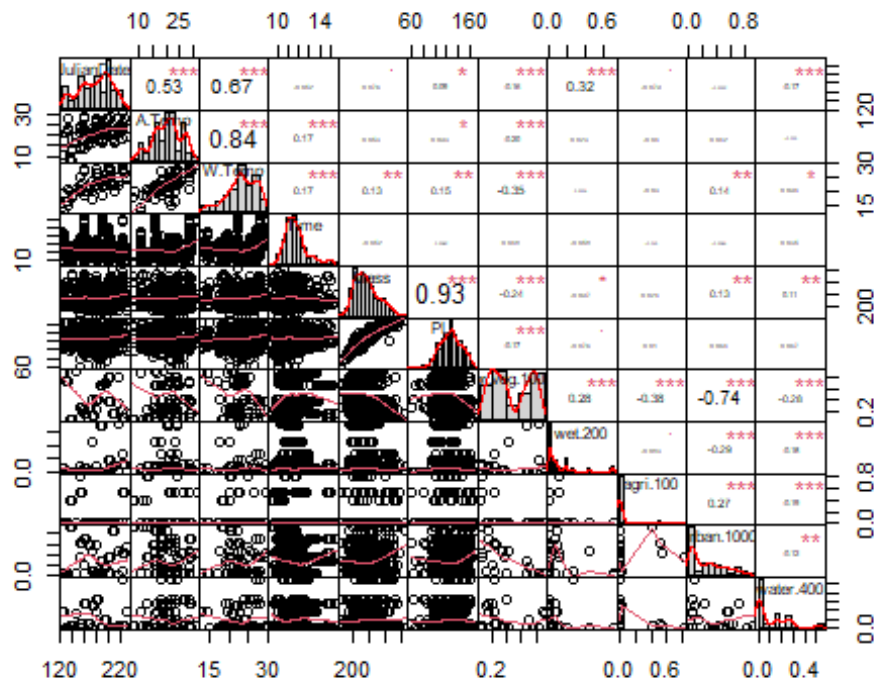
All of the  $GVIF^{(1/(2*df))} < 2$ , so I will not remove any more variables.

## Calculation of Pearson and Spearman correlation coefficients

### Pearson correlation coefficients

#### # Visualization of the correlations

```
cor.pearson.Aggression <- MixedData[, c(3,4,5,6,7,8,26,28,37,56,60)]
chart.Correlation(cor.pearson.Aggression, histogram=TRUE, pch=19)
```



*# Creation of the correlation table*

```
table.corr.pearson.Agression <- rcorr(as.matrix(MixedData[,c(3,4,5,6,7,8,26,
28,37,56,60)]), type="pearson")
```

```
table.cor.pearson.Agression.r <- table.corr.pearson.Agression$r # Pearson c
correlation coefficients
```

```
table.cor.pearson.Agression.p <- table.corr.pearson.Agression$P # P values
of the correlations
```

```
table.cor.pearson.Agression.r
```

##	JulianDate	A.Temp	W.Temp	Time	Mas
## JulianDate	1.00000000	0.52662360	0.668683098	-0.051688682	0.07560208
## A.Temp	0.526623600	1.00000000	0.838871081	0.171967476	0.05252964
## W.Temp	0.668683098	0.83887108	1.00000000	0.166126059	0.13221505
## Time	-0.051688682	0.17196748	0.166126059	1.00000000	-0.05245171
## Mass	0.075602082	0.05252964	0.132215053	-0.052451709	1.00000000
## PL	0.089516177	0.09250283	0.146064749	0.009461487	0.93311455
## for.veg.1000	-0.159757526	-0.20286665	-0.354421886	0.048125471	-0.23967087
## wet.200	0.319118112	0.02344567	0.002124113	-0.057823266	-0.09719429

```

## agri.100      -0.078714526 -0.03995312 -0.030225708 -0.013077109  0.0240670
4
## urban.1000   -0.003462429  0.04205111  0.143942429 -0.006102219  0.1273897
4
## water.400    -0.169078991 -0.02278033  0.093636191  0.035069081  0.1126047
0
##              PL for.veg.1000      wet.200      agri.100      urban.10
00
## JulianDate    0.089516177  -0.15975753  0.319118112 -0.07871453 -0.0034624
29
## A.Temp        0.092502829  -0.20286665  0.023445668 -0.03995312  0.0420511
10
## W.Temp        0.146064749  -0.35442189  0.002124113 -0.03022571  0.1439424
29
## Time          0.009461487  0.04812547 -0.057823266 -0.01307711 -0.0061022
19
## Mass          0.933114554  -0.23967087 -0.097194285  0.02406704  0.1273897
43
## PL            1.000000000  -0.17433634 -0.076087500  0.01037365  0.0639451
49
## for.veg.1000 -0.174336336  1.00000000  0.276458046 -0.38318444 -0.7437982
23
## wet.200       -0.076087500  0.27645805  1.000000000 -0.08318312 -0.2948294
46
## agri.100      0.010373653  -0.38318444 -0.083183118  1.00000000  0.2719946
55
## urban.1000    0.063945149  -0.74379822 -0.294829446  0.27199465  1.0000000
00
## water.400     0.067406397  -0.28142967 -0.183447101 -0.19070486 -0.1173627
38
##              water.400
## JulianDate    -0.16907899
## A.Temp        -0.02278033
## W.Temp        0.09363619
## Time          0.03506908
## Mass          0.11260470
## PL            0.06740640
## for.veg.1000 -0.28142967
## wet.200       -0.18344710
## agri.100      -0.19070486
## urban.1000    -0.11736274
## water.400     1.00000000

```

table.cor.pearson.Aggression.p

```

##              JulianDate      A.Temp      W.Temp      Time      M
ass
## JulianDate      NA 0.000000e+00 0.000000e+00 2.322103e-01 8.033738e
-02
## A.Temp          0.000000e+00      NA 0.000000e+00 6.283828e-05 2.246911e

```

```

-01
## W.Temp      0.000000e+00 0.000000e+00          NA 2.346306e-04 3.499660e
-03
## Time        2.322103e-01 6.283828e-05 2.346306e-04          NA 2.253805e
-01
## Mass        8.033738e-02 2.246911e-01 3.499660e-03 2.253805e-01
NA
## PL          3.846835e-02 3.242105e-02 1.256393e-03 8.271659e-01 0.000000e
+00
## for.veg.1000 2.040904e-04 2.189891e-06 8.881784e-16 2.660399e-01 1.932793e
-08
## wet.200      3.730349e-14 5.880847e-01 9.627472e-01 1.813210e-01 2.443073e
-02
## agri.100     6.861526e-02 3.559077e-01 5.061950e-01 7.626040e-01 5.782300e
-01
## urban.1000   9.362578e-01 3.311946e-01 1.464223e-03 8.879110e-01 3.132312e
-03
## water.400    8.365543e-05 5.987216e-01 3.906820e-02 4.177898e-01 9.075308e
-03
##              PL for.veg.1000      wet.200      agri.100      urban.1
000
## JulianDate   3.846835e-02 2.040904e-04 3.730349e-14 6.861526e-02 9.362578e
-01
## A.Temp       3.242105e-02 2.189891e-06 5.880847e-01 3.559077e-01 3.311946e
-01
## W.Temp       1.256393e-03 8.881784e-16 9.627472e-01 5.061950e-01 1.464223e
-03
## Time         8.271659e-01 2.660399e-01 1.813210e-01 7.626040e-01 8.879110e
-01
## Mass         0.000000e+00 1.932793e-08 2.443073e-02 5.782300e-01 3.132312e
-03
## PL           NA 5.033503e-05 7.868930e-02 8.108043e-01 1.396467e
-01
## for.veg.1000 5.033503e-05          NA 7.372303e-11 0.000000e+00 0.000000e
+00
## wet.200      7.868930e-02 7.372303e-11          NA 5.427077e-02 3.277600e
-12
## agri.100     8.108043e-01 0.000000e+00 5.427077e-02          NA 1.518086e
-10
## urban.1000   1.396467e-01 0.000000e+00 3.277600e-12 1.518086e-10
NA
## water.400    1.194137e-01 3.246692e-11 1.924186e-05 8.760883e-06 6.524110e
-03
##              water.400
## JulianDate   8.365543e-05
## A.Temp       5.987216e-01
## W.Temp       3.906820e-02
## Time         4.177898e-01
## Mass         9.075308e-03
## PL           1.194137e-01

```

```
## for.veg.1000 3.246692e-11
## wet.200      1.924186e-05
## agri.100     8.760883e-06
## urban.1000   6.524110e-03
## water.400    NA
```

### *Spearman correlation coefficients*

#### *# Creation of the correlation table*

```
table.corr.spearman.Aggression <- rcorr(as.matrix(MixedData[,c(3,4,5,6,7,8,26,28,37,56,60)]), type="spearman")
table.cor.spearman.Aggression.r <- table.corr.spearman.Aggression$r # Pearson correlation coefficients
table.cor.spearman.Aggression.p <- table.corr.spearman.Aggression$P # P values of the correlations
table.cor.spearman.Aggression.r
```

```
##          JulianDate      A.Temp      W.Temp      Time      Mass
## JulianDate  1.00000000  0.45842169  0.56884913 -0.098920865  0.08038929
## A.Temp      0.45842169  1.00000000  0.83718368  0.138498905  0.06589666
## W.Temp      0.56884913  0.83718368  1.00000000  0.173659029  0.13270468
## Time        -0.09892087  0.13849890  0.17365903  1.000000000  -0.04025948
## Mass        0.08038929  0.06589666  0.13270468 -0.040259479  1.00000000
## PL          0.08898378  0.10442546  0.15193967 -0.013644546  0.94856246
## for.veg.1000 -0.13614271 -0.17001898 -0.28928407  0.119570436 -0.19217208
## wet.200     0.38993834  0.04537204  0.12470905  0.022468674 -0.04181263
## agri.100    -0.08934127 -0.02430981 -0.06768594 -0.063059962  0.04983615
## urban.1000  0.05728527  0.13104537  0.21782810 -0.071452190  0.10868203
## water.400   -0.30298704 -0.10238175 -0.01252963  0.007177042  0.09129109
##          PL for.veg.1000      wet.200      agri.100      urban.1000
## JulianDate  0.08898378  -0.1361427  0.38993834 -0.08934127  0.05728527
## A.Temp      0.10442546  -0.1700190  0.04537204 -0.02430981  0.13104537
## W.Temp      0.15193967  -0.2892841  0.12470905 -0.06768594  0.21782810
## Time        -0.01364455  0.1195704  0.02246867 -0.06305996 -0.07145219
## Mass        0.94856246  -0.1921721 -0.04181263  0.04983615  0.10868203
## PL          1.00000000  -0.1623624 -0.06444220  0.05678992  0.07998962
## for.veg.1000 -0.16236237  1.0000000  0.16925793 -0.47395684 -0.82572013
## wet.200     -0.06444220  0.1692579  1.00000000 -0.06033822 -0.23397742
## agri.100    0.05678992  -0.4739568 -0.06033822  1.00000000  0.27938984
## urban.1000  0.07998962  -0.8257201 -0.23397742  0.27938984  1.00000000
## water.400   0.06318439  -0.2190942 -0.19135539 -0.03550963  0.02511729
##          water.400
## JulianDate  -0.302987043
## A.Temp      -0.102381748
## W.Temp      -0.012529631
## Time        0.007177042
## Mass        0.091291089
## PL          0.063184387
## for.veg.1000 -0.219094175
## wet.200     -0.191355393
## agri.100    -0.035509628
```

```
## urban.1000    0.025117287
## water.400     1.000000000
```

```
table.cor.spearman.Aggression.p
```

```
##          JulianDate      A.Temp      W.Temp      Time      M
ass
## JulianDate      NA 0.000000e+00 0.000000e+00 0.0219948308 6.291206e
-02
## A.Temp      0.000000e+00      NA 0.000000e+00 0.0013067179 1.275805e
-01
## W.Temp      0.000000e+00 0.000000e+00      NA 0.0001191832 3.379071e
-03
## Time      2.199483e-02 1.306718e-03 1.191832e-04      NA 3.522285e
-01
## Mass      6.291206e-02 1.275805e-01 3.379071e-03 0.3522284632
NA
## PL      3.964107e-02 1.567793e-02 7.879312e-04 0.7528559899 0.000000e
+00
## for.veg.1000 1.581516e-03 7.625616e-05 8.007350e-11 0.0055757019 7.445223e
-06
## wet.200      0.000000e+00 2.943962e-01 5.906328e-03 0.6037338847 3.339467e
-01
## agri.100      3.866754e-02 5.744009e-01 1.362161e-01 0.1448442211 2.493981e
-01
## urban.1000 1.854229e-01 2.366128e-03 1.246624e-06 0.0984350178 1.181043e
-02
## water.400      7.642775e-13 1.773963e-02 7.829165e-01 0.8683342163 3.459895e
-02
##          PL for.veg.1000      wet.200      agri.100      urban.1
000
## JulianDate 0.0396410665 1.581516e-03 0.000000e+00 3.866754e-02 1.854229e
-01
## A.Temp      0.0156779333 7.625616e-05 2.943962e-01 5.744009e-01 2.366128e
-03
## W.Temp      0.0007879312 8.007350e-11 5.906328e-03 1.362161e-01 1.246624e
-06
## Time      0.7528559899 5.575702e-03 6.037339e-01 1.448442e-01 9.843502e
-02
## Mass      0.0000000000 7.445223e-06 3.339467e-01 2.493981e-01 1.181043e
-02
## PL      NA 1.621239e-04 1.365892e-01 1.896750e-01 6.448653e
-02
## for.veg.1000 0.0001621239      NA 8.219673e-05 0.000000e+00 0.000000e
+00
## wet.200      0.1365891515 8.219673e-05      NA 1.630337e-01 4.239393e
-08
## agri.100      0.1896750465 0.000000e+00 1.630337e-01      NA 4.554446e
-11
## urban.1000 0.0644865292 0.000000e+00 4.239393e-08 4.554446e-11
```

```

NA
## water.400      0.1444276999 3.008301e-07 8.152307e-06 4.119614e-01 5.617520e
-01
##              water.400
## JulianDate    7.642775e-13
## A.Temp        1.773963e-02
## W.Temp        7.829165e-01
## Time          8.683342e-01
## Mass          3.459895e-02
## PL            1.444277e-01
## for.veg.1000  3.008301e-07
## wet.200       8.152307e-06
## agri.100      4.119614e-01
## urban.1000    5.617520e-01
## water.400     NA

```

The Pearson correlation coefficient of 0.93 between Mass and PL, 0.84 between W.Temp and A.Temp and -0.74 between urban.for.veg.1000 and urban.1000 confirmed the deletion of Mass, W.Temp, and for.veg.1000 with the calculation of the  $GVIF^{(1/(2*df))}$ .

#### Verification of the assumptions with the initial model

```

mod.Aggression.full <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scaled +
Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000
.Scaled + water.400.Scaled + (1|Code) + (1|Site), data = MixedData, REML = TR
UE, na.action=na.exclude)

```

```

summary(mod.Aggression.full)

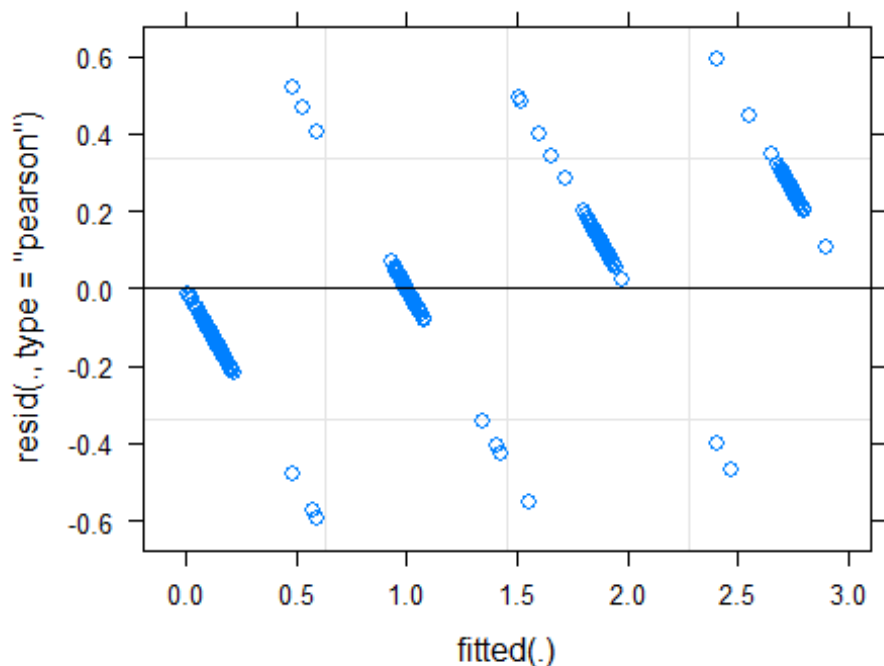
```

```

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
##          PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled
+
##          water.400.Scaled + (1 | Code) + (1 | Site)
## Data: MixedData
##
## REML criterion at convergence: 1500.1
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.55498 -0.31577 -0.06114  0.28952  1.55232
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
##  Code     (Intercept) 0.89340   0.9452
##  Site     (Intercept) 0.03487   0.1867
##  Residual                    0.14606   0.3822
## Number of obs: 522, groups: Code, 492; Site, 24
##
## Fixed effects:

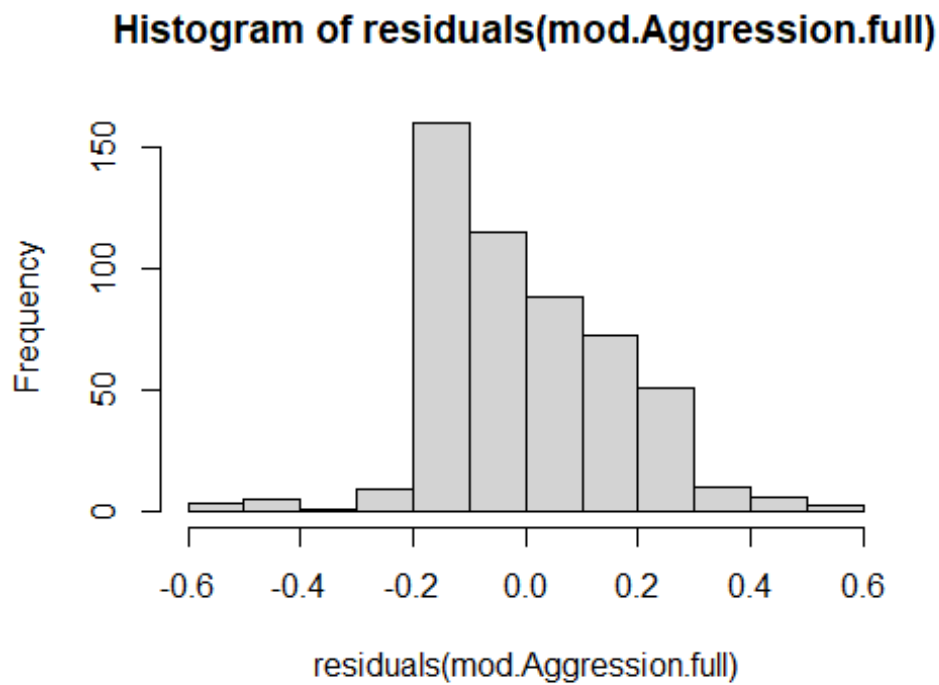
```

```
##               Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)      1.00148    0.08420   31.50789   11.895 3.46e-13 ***
## JulianDate.Scaled -0.05395    0.07452   17.14086    -0.724   0.4788
## A.Temp.Scaled      0.14378    0.06375   45.47667    2.255   0.0290 *
## Time.Scaled        -0.04752    0.04417  253.92511   -1.076   0.2831
## PL.Scaled          -0.05307    0.05647  474.75813   -0.940   0.3478
## SexM               0.04213    0.09554  247.28036    0.441   0.6596
## wet.200.Scaled     -0.03924    0.06700   12.36223   -0.586   0.5686
## agri.100.Scaled    -0.05992    0.05884   23.67696   -1.018   0.3188
## urban.1000.Scaled  0.09694    0.06763   10.82073    1.433   0.1800
## water.400.Scaled   0.13923    0.06576   10.79946    2.117   0.0583 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##      (Intr) JlnD.S A.Tm.S Tm.Scl PL.Scl SexM  w.200. a.100. u.1000
## JulnDt.Scl -0.048
## A.Temp.Scl  0.073 -0.467
## Time.Scaled 0.025  0.135 -0.253
## PL.Scaled   -0.336 -0.051 -0.046 -0.021
## SexM         -0.670  0.044 -0.066 -0.022  0.455
## wt.200.Scl  -0.066 -0.326  0.122 -0.012  0.084  0.022
## agr.100.Scl  0.001  0.114 -0.023  0.016 -0.067 -0.086  0.024
## urbn.1000.S -0.098 -0.126  0.065  0.014  0.017  0.067  0.349 -0.228
## wtr.400.Scl -0.011  0.038 -0.015 -0.012 -0.101 -0.050  0.234  0.202  0.194
plot(mod.Aggression.full)
```

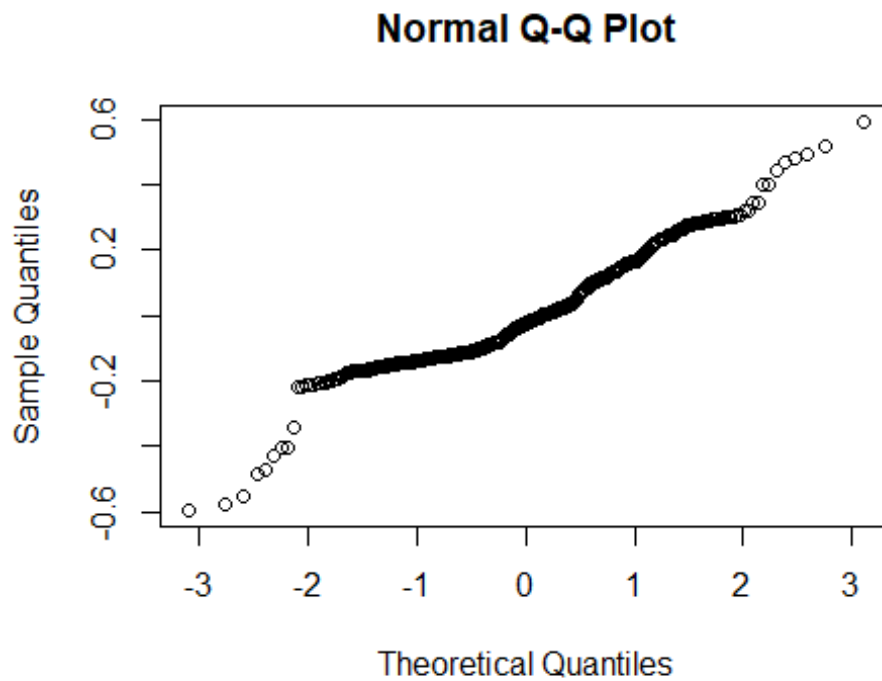




```
hist(residuals(mod.Aggression.full))
```



```
qqnorm(resid(mod.Aggression.full))
```



## Time of initial movement (Start)

### Multicollinearity: Generalized variance inflation factor ( $GVIF^{1/(2*df)}$ )

$GVIF^{1/(2*df)} > 2$  indicates the presence of multicollinearity, so I will remove variables with values over 2, starting with the highest value.

```
mod.Start.vif <- lm(Start ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled
+ Time.Scaled + Mass.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.
Scaled + agri.600.Scaled + urban.1000.Scaled + water.500.Scaled, data = Mixed
Data, na.action=na.exclude)
gvif(mod.Start.vif)
```

##		GVIF	df	$GVIF^{1/(2*df)}$
##	JulianDate.Scaled	2.2956	1	1.5151
##	A.Temp.Scaled	4.2591	1	2.0638
##	W.Temp.Scaled	5.6476	1	2.3765
##	Time.Scaled	1.1199	1	1.0583
##	Mass.Scaled	8.8715	1	2.9785
##	PL.Scaled	7.9423	1	2.8182
##	Sex	1.4811	1	1.2170
##	for.veg.200.Scaled	3.3965	1	1.8430
##	wet.400.Scaled	2.6633	1	1.6320
##	agri.600.Scaled	1.3300	1	1.1533
##	urban.1000.Scaled	2.9008	1	1.7032
##	water.500.Scaled	2.0301	1	1.4248

Turtle mass (Mass) has the highest  $GVIF^{1/(2*df)} > 2$ , so I will remove it and recalculate the factors.

```
mod.Start.vif <- lm(Start ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled
+ Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.
600.Scaled + urban.1000.Scaled + water.500.Scaled, data = MixedData, na.actio
n=na.exclude)
gvif(mod.Start.vif)
```

##		GVIF	df	$GVIF^{1/(2*df)}$
##	JulianDate.Scaled	2.2953	1	1.5150
##	A.Temp.Scaled	4.2167	1	2.0535
##	W.Temp.Scaled	5.6371	1	2.3743
##	Time.Scaled	1.1054	1	1.0514
##	PL.Scaled	1.3687	1	1.1699
##	Sex	1.3664	1	1.1689
##	for.veg.200.Scaled	3.3965	1	1.8430
##	wet.400.Scaled	2.6633	1	1.6320
##	agri.600.Scaled	1.3279	1	1.1523
##	urban.1000.Scaled	2.8466	1	1.6872
##	water.500.Scaled	1.9759	1	1.4057

Water temperature (W.Temp) has the highest  $GVIF^{1/(2*df)} > 2$ , so I will remove it and recalculate the factors.

```
mod.Start.vif <- lm(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urb
an.1000.Scaled + water.500.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.Start.vif)
```

```
##          GVIF df GVIF^(1/(2*df))
## JulianDate.Scaled  1.6219  1      1.2735
## A.Temp.Scaled      1.7465  1      1.3216
## Time.Scaled        1.0943  1      1.0461
## PL.Scaled          1.3825  1      1.1758
## Sex                1.3683  1      1.1697
## for.veg.200.Scaled  3.2244  1      1.7957
## wet.400.Scaled     2.5927  1      1.6102
## agri.600.Scaled    1.4225  1      1.1927
## urban.1000.Scaled  2.6813  1      1.6375
## water.500.Scaled   1.7734  1      1.3317
```

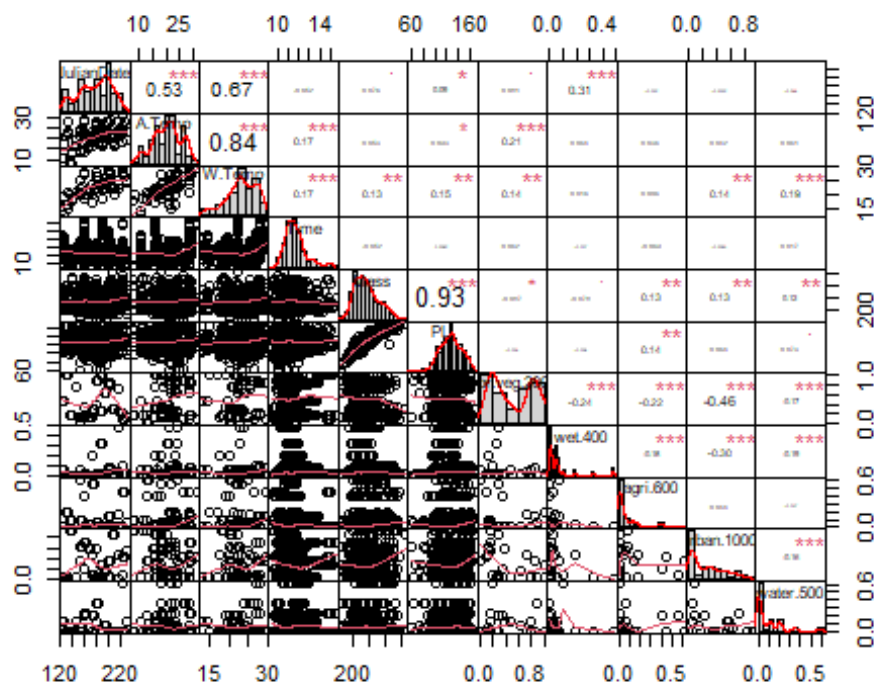
All of the  $GVIF^{1/(2*df)} < 2$ , so I will not remove any more variables.

## Calculation of Pearson and Spearman correlation coefficients

### Pearson correlation coefficients

*# Visualization of the correlations*

```
cor.pearson.Start <- MixedData[, c(3,4,5,6,7,8,18,30,42,56,61)]
chart.Correlation(cor.pearson.Start, histogram=TRUE, pch=19)
```



```
# Creation of the correlation table
```

```
table.corr.pearson.Start <- rcorr(as.matrix(MixedData[,c(3,4,5,6,7,8,18,30,42,56,61)]), type="pearson")
```

```
table.cor.pearson.Start.r <- table.corr.pearson.Start$r # Pearson correlation coefficients
```

```
table.cor.pearson.Start.p <- table.corr.pearson.Start$P # P values of the correlations
```

```
table.cor.pearson.Start.r
```

```
##          JulianDate      A.Temp      W.Temp      Time      Mass
## JulianDate  1.000000000  0.52662360  0.668683098 -0.051688682  0.07560208
## A.Temp      0.526623600  1.000000000  0.838871081  0.171967476  0.05252964
## W.Temp      0.668683098  0.838871081  1.000000000  0.166126059  0.13221505
## Time        -0.051688682  0.17196748  0.166126059  1.000000000 -0.05245171
## Mass        0.075602082  0.05252964  0.132215053 -0.052451709  1.000000000
## PL          0.089516177  0.09250283  0.146064749  0.009461487  0.93311455
## for.veg.200 0.081109497  0.21284818  0.135718201  0.062119056 -0.08740432
## wet.400     0.306093497  0.06431535  0.015608535 -0.036551862 -0.07810264
## agri.600    -0.035110853  0.03604656  0.006005415 -0.069402894  0.13163938
## urban.1000 -0.003462429  0.04205111  0.143942429 -0.006102219  0.12738974
## water.500   -0.044004133  0.06070427  0.192147258  0.016514392  0.11851616
##          PL for.veg.200      wet.400      agri.600      urban.1000
## JulianDate  0.089516177  0.08110950  0.30609350 -0.035110853 -0.003462429
## A.Temp      0.092502829  0.21284818  0.06431535  0.036046563  0.042051110
## W.Temp      0.146064749  0.13571820  0.01560853  0.006005415  0.143942429
## Time        0.009461487  0.06211906 -0.03655186 -0.069402894 -0.006102219
## Mass        0.933114554 -0.08740432 -0.07810264  0.131639380  0.127389743
## PL          1.000000000 -0.03366684 -0.03941912  0.137973397  0.063945149
## for.veg.200 -0.033666836  1.00000000 -0.24135779 -0.219409133 -0.460382656
## wet.400     -0.039419118 -0.24135779  1.00000000 -0.178579433 -0.297525027
## agri.600    0.137973397 -0.21940913 -0.17857943  1.000000000  0.043890365
## urban.1000  0.063945149 -0.46038266 -0.29752503  0.043890365  1.000000000
## water.500   0.072939414 -0.17138084 -0.18871224 -0.017252283 -0.155373576
##          water.500
## JulianDate -0.04400413
## A.Temp      0.06070427
## W.Temp      0.19214726
## Time        0.01651439
## Mass        0.11851616
## PL          0.07293941
## for.veg.200 -0.17138084
## wet.400     -0.18871224
## agri.600    -0.01725228
## urban.1000 -0.15537358
## water.500   1.00000000
```

```
table.cor.pearson.Start.p
```

```
##          JulianDate      A.Temp      W.Temp      Time      Mas
s
```

```

## JulianDate          NA 0.000000e+00 0.000000e+00 2.322103e-01 0.08033738
4
## A.Temp              0.000000e+00          NA 0.000000e+00 6.283828e-05 0.22469108
8
## W.Temp              0.000000e+00 0.000000e+00          NA 2.346306e-04 0.00349966
0
## Time                2.322103e-01 6.283828e-05 2.346306e-04          NA 0.22538053
4
## Mass                8.033738e-02 2.246911e-01 3.499660e-03 2.253805e-01          N
A
## PL                  3.846835e-02 3.242105e-02 1.256393e-03 8.271659e-01 0.00000000
0
## for.veg.200         6.058262e-02 6.579274e-07 2.716312e-03 1.509483e-01 0.04310363
0
## wet.400             4.334311e-13 1.369959e-01 7.314244e-01 3.983658e-01 0.07080188
5
## agri.600           4.172351e-01 4.049233e-01 8.949431e-01 1.084987e-01 0.00225920
1
## urban.1000         9.362578e-01 3.311946e-01 1.464223e-03 8.879110e-01 0.00313231
2
## water.500          3.092088e-01 1.604920e-01 2.000264e-05 7.028551e-01 0.00601193
5
##                      PL   for.veg.200      wet.400      agri.600      urban.100
0
## JulianDate          0.038468345 6.058262e-02 4.334311e-13 4.172351e-01 9.362578e-0
1
## A.Temp              0.032421052 6.579274e-07 1.369959e-01 4.049233e-01 3.311946e-0
1
## W.Temp              0.001256393 2.716312e-03 7.314244e-01 8.949431e-01 1.464223e-0
3
## Time                0.827165890 1.509483e-01 3.983658e-01 1.084987e-01 8.879110e-0
1
## Mass                0.000000000 4.310363e-02 7.080188e-02 2.259201e-03 3.132312e-0
3
## PL                  NA 4.370909e-01 3.628289e-01 1.378178e-03 1.396467e-0
1
## for.veg.200         0.437090868          NA 1.525600e-08 2.890116e-07 0.000000e+0
0
## wet.400             0.362828857 1.525600e-08          NA 3.207206e-05 2.036149e-1
2
## agri.600           0.001378178 2.890116e-07 3.207206e-05          NA 3.104625e-0
1
## urban.1000         0.139646656 0.000000e+00 2.036149e-12 3.104625e-01          N
A
## water.500          0.091912781 6.662331e-05 1.090597e-05 6.902496e-01 3.053590e-0
4
##                      water.500
## JulianDate          3.092088e-01
## A.Temp              1.604920e-01
## W.Temp              2.000264e-05

```

```
## Time          7.028551e-01
## Mass          6.011935e-03
## PL            9.191278e-02
## for.veg.200   6.662331e-05
## wet.400       1.090597e-05
## agri.600      6.902496e-01
## urban.1000    3.053590e-04
## water.500     NA
```

### *Spearman correlation coefficients*

*# Creation of the correlation table*

```
table.corr.spearman.Start <- rcorr(as.matrix(MixedData[,c(3,4,5,6,7,8,18,30,42,56,61)]), type="spearman")
```

```
table.cor.spearman.Start.r <- table.corr.spearman.Start$r # Pearson correlation coefficients
```

```
table.cor.spearman.Start.p <- table.corr.spearman.Start$p # P values of the correlations
```

```
table.cor.spearman.Start.r
```

```
##          JulianDate      A.Temp      W.Temp      Time      Mass
## JulianDate  1.00000000  0.45842169  0.56884913 -0.098920865  0.08038929
## A.Temp      0.45842169  1.00000000  0.83718368  0.138498905  0.06589666
## W.Temp      0.56884913  0.83718368  1.00000000  0.173659029  0.13270468
## Time        -0.09892087  0.13849890  0.17365903  1.000000000  -0.04025948
## Mass        0.08038929  0.06589666  0.13270468  -0.040259479  1.00000000
## PL          0.08898378  0.10442546  0.15193967 -0.013644546  0.94856246
## for.veg.200  0.07590530  0.18670931  0.17243577  0.072682965  -0.05303332
## wet.400     0.41005248  0.11243977  0.19326625  0.146391742  -0.04217241
## agri.600    -0.06694664  0.22598407  0.11700410 -0.093444945  0.16767037
## urban.1000  0.05728527  0.13104537  0.21782810 -0.071452190  0.10868203
## water.500   -0.27348685 -0.05671475  0.03613419  0.002018679  0.09506451
##          PL for.veg.200      wet.400      agri.600      urban.1000
## JulianDate  0.08898378  0.07590530  0.41005248 -0.066946640  0.05728527
## A.Temp      0.10442546  0.18670931  0.11243977  0.225984071  0.13104537
## W.Temp      0.15193967  0.17243577  0.19326625  0.117004100  0.21782810
## Time        -0.01364455  0.07268297  0.14639174 -0.093444945  -0.07145219
## Mass        0.94856246 -0.05303332 -0.04217241  0.167670371  0.10868203
## PL          1.00000000 -0.01916834 -0.04821271  0.181622684  0.07998962
## for.veg.200 -0.01916834  1.00000000 -0.14248989 -0.184907521  -0.41380576
## wet.400     -0.04821271 -0.14248989  1.00000000 -0.300029610  -0.23226774
## agri.600    0.18162268 -0.18490752 -0.30002961  1.000000000  0.48970284
## urban.1000  0.07998962 -0.41380576 -0.23226774  0.489702837  1.00000000
## water.500   0.06924256 -0.27799035 -0.42187319  0.005477243  0.05503142
##          water.500
## JulianDate -0.273486853
## A.Temp      -0.056714750
## W.Temp      0.036134191
## Time        0.002018679
## Mass        0.095064505
## PL          0.069242556
```

```
## for.veg.200 -0.277990353
## wet.400      -0.421873192
## agri.600     0.005477243
## urban.1000   0.055031416
## water.500    1.000000000
```

```
table.cor.spearman.Start.p
```

```
##          JulianDate      A.Temp      W.Temp      Time      Ma
ss
## JulianDate      NA 0.000000e+00 0.000000e+00 0.0219948308 6.291206e-
02
## A.Temp      0.000000e+00      NA 0.000000e+00 0.0013067179 1.275805e-
01
## W.Temp      0.000000e+00 0.000000e+00      NA 0.0001191832 3.379071e-
03
## Time      2.199483e-02 1.306718e-03 1.191832e-04      NA 3.522285e-
01
## Mass      6.291206e-02 1.275805e-01 3.379071e-03 0.3522284632
NA
## PL      3.964107e-02 1.567793e-02 7.879312e-04 0.7528559899 0.000000e+
00
## for.veg.200 7.912840e-02 1.356056e-05 1.332995e-04 0.0927575303 2.202715e-
01
## wet.400      0.000000e+00 9.177827e-03 1.785318e-05 0.0006747625 3.298003e-
01
## agri.600      1.216114e-01 1.234880e-07 9.833391e-03 0.0305341790 9.602068e-
05
## urban.1000 1.854229e-01 2.366128e-03 1.246624e-06 0.0984350178 1.181043e-
02
## water.500 1.194147e-10 1.898472e-01 4.267291e-01 0.9628107149 2.775370e-
02
##          PL  for.veg.200      wet.400      agri.600      urban.10
00
## JulianDate 3.964107e-02 7.912840e-02 0.000000e+00 1.216114e-01 1.854229e-
01
## A.Temp      1.567793e-02 1.356056e-05 9.177827e-03 1.234880e-07 2.366128e-
03
## W.Temp      7.879312e-04 1.332995e-04 1.785318e-05 9.833391e-03 1.246624e-
06
## Time      7.528560e-01 9.275753e-02 6.747625e-04 3.053418e-02 9.843502e-
02
## Mass      0.000000e+00 2.202715e-01 3.298003e-01 9.602068e-05 1.181043e-
02
## PL      NA 6.582225e-01 2.656216e-01 2.375451e-05 6.448653e-
02
## for.veg.200 6.582225e-01      NA 9.394054e-04 1.646414e-05 0.000000e+
00
## wet.400      2.656216e-01 9.394054e-04      NA 1.302514e-12 5.346181e-
08
```

```
## agri.600      2.375451e-05 1.646414e-05 1.302514e-12      NA 0.000000e+
00
## urban.1000   6.448653e-02 0.000000e+00 5.346181e-08 0.000000e+00
NA
## water.500    1.096520e-01 5.735767e-11 0.000000e+00 8.993265e-01 2.033521e-
01
##              water.500
## JulianDate   1.194147e-10
## A.Temp       1.898472e-01
## W.Temp       4.267291e-01
## Time         9.628107e-01
## Mass         2.775370e-02
## PL           1.096520e-01
## for.veg.200  5.735767e-11
## wet.400      0.000000e+00
## agri.600     8.993265e-01
## urban.1000   2.033521e-01
## water.500    NA
```

The Pearson correlation coefficient of 0.93 between Mass and PL and 0.84 between W.Temp and A.Temp confirmed the deletion of Mass and W.Temp with the calculation of the  $GVIF^{(1/(2*df))}$ .

#### Verification of the assumptions with the initial model

```
mod.Start.full <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scale
d + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled +
urban.1000.Scaled + water.500.Scaled + (1|Code) + (1|Site), data = MixedData,
REML = TRUE, na.action=na.exclude)
```

```
summary(mod.Start.full)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scal
ed +
##      Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled +
##      urban.1000.Scaled + water.500.Scaled + (1 | Code) + (1 |      Site)
##      Data: MixedData
##
## REML criterion at convergence: 5522.9
##
## Scaled residuals:
##      Min      1Q  Median      3Q      Max
## -1.7051 -0.5974 -0.2511  0.4462  4.0698
##
## Random effects:
##      Groups      Name      Variance Std.Dev.
##      Code      (Intercept)  1711      41.37
##      Site      (Intercept)  2377      48.75
##      Residual              10853     104.18
```

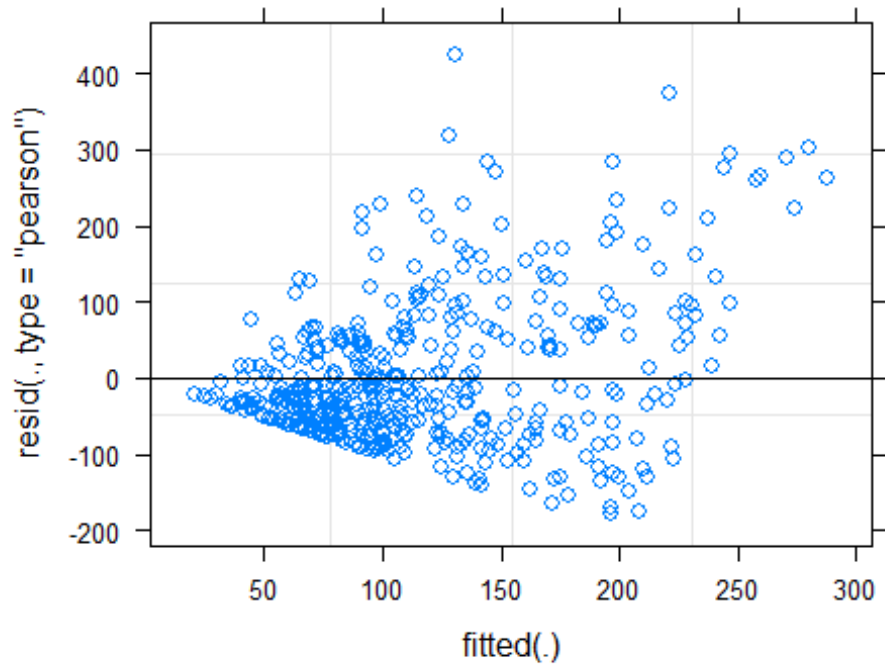


```

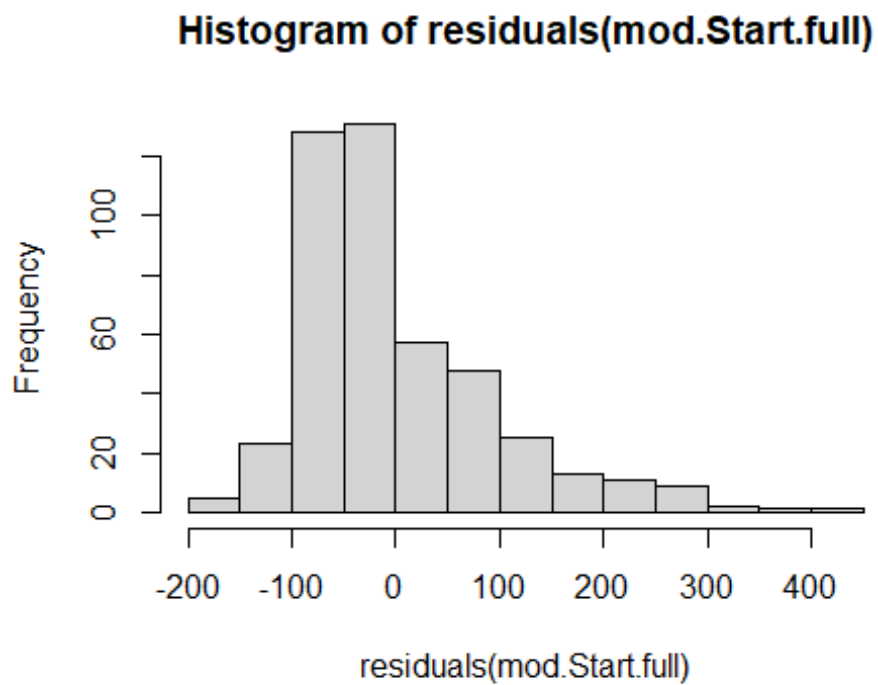
## Number of obs: 454, groups: Code, 430; Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    121.855    14.207    25.474   8.577 5.53e-09 ***
## JulianDate.Scaled -19.886    13.450    18.438  -1.478  0.15616
## A.Temp.Scaled   -4.217    10.057   148.565  -0.419  0.67559
## Time.Scaled     -4.557     5.925   440.328  -0.769  0.44221
## PL.Scaled       18.995     6.949   418.253   2.734  0.00653 **
## SexM            5.277    12.823   439.387   0.412  0.68089
## for.veg.200.Scaled -15.564    20.757    14.567  -0.750  0.46531
## wet.400.Scaled   17.301    18.906    13.880   0.915  0.37575
## agri.600.Scaled   1.036    12.655    15.764   0.082  0.93575
## urban.1000.Scaled -15.886    19.774    13.777  -0.803  0.43539
## water.500.Scaled -12.429    15.241    13.489  -0.816  0.42895
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S Tm.Sc1 PL.Sc1 SexM   f..200 w.400. a.600.
## JulnDt.Scld -0.057
## A.Temp.Scld  0.100 -0.358
## Time.Scaled  0.045  0.086 -0.235
## PL.Scaled    -0.264 -0.021 -0.031 -0.039
## SexM         -0.522  0.042 -0.116 -0.050  0.469
## fr.vg.200.S  0.009 -0.122 -0.219  0.044 -0.015  0.051
## wt.400.Scld -0.058 -0.323 -0.093  0.030  0.010  0.039  0.704
## agr.600.Sc1 -0.106 -0.070 -0.146  0.056 -0.048  0.009  0.591  0.553
## urbn.1000.S -0.071 -0.230 -0.090  0.049 -0.014  0.057  0.751  0.730  0.512
## wtr.500.Sc1 -0.043 -0.204 -0.117  0.017 -0.059  0.011  0.575  0.599  0.426
##              u.1000
## JulnDt.Scld
## A.Temp.Scld
## Time.Scaled
## PL.Scaled
## SexM
## fr.vg.200.S
## wt.400.Scld
## agr.600.Sc1
## urbn.1000.S
## wtr.500.Sc1  0.624

plot(mod.Start.full)

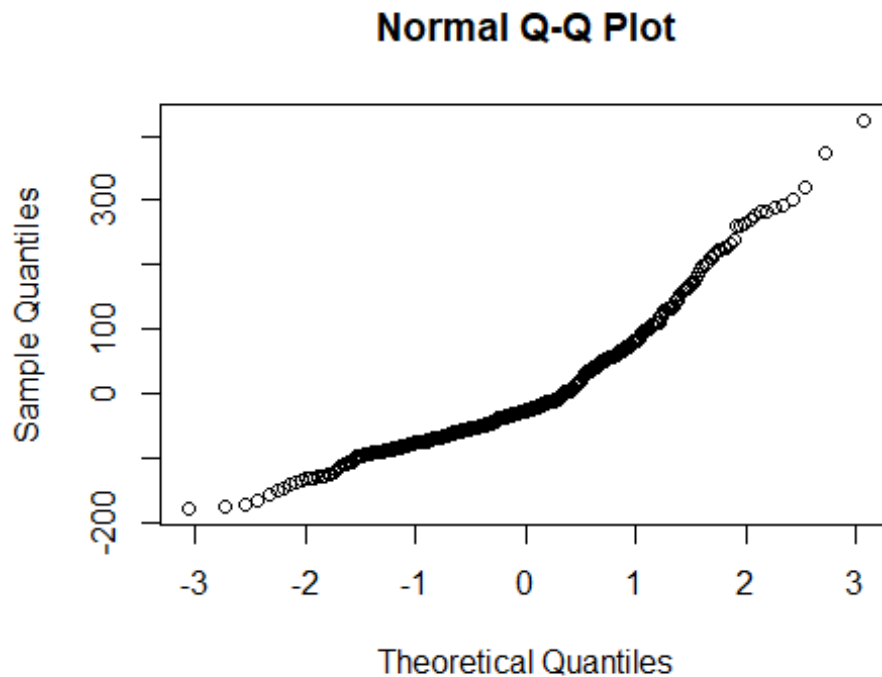
```



```
hist(residuals(mod.Start.full))
```



```
qqnorm(resid(mod.Start.full))
```



### Log(x+1) transformed version of time of initial movement (log.Start)

```
MixedData$log.Start <- as.numeric(MixedData$log.Start)
```

### Multicollinearity: Generalized variance inflation factor ( $\text{GVIF}^{1/(2 \cdot \text{df})}$ )

$\text{GVIF}^{1/(2 \cdot \text{df})} > 2$  indicates the presence of multicollinearity, so I will remove variables with values over 2, starting with the highest value.

```
mod.log.Start.vif <- lm(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled + Time.Scaled + Mass.Scaled + PL.Scaled + Sex + for.veg.1000.Scaled + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.log.Start.vif)
```

	GVIF	df	$\text{GVIF}^{1/(2 \cdot \text{df})}$
JulianDate.Scaled	2.6565	1	1.6299
A.Temp.Scaled	4.0073	1	2.0018
W.Temp.Scaled	5.7361	1	2.3950
Time.Scaled	1.1459	1	1.0705
Mass.Scaled	8.8880	1	2.9813
PL.Scaled	7.9226	1	2.8147
Sex	1.4672	1	1.2113
for.veg.1000.Scaled	6.6011	1	2.5693
wet.400.Scaled	1.2942	1	1.1376
agri.400.Scaled	2.3605	1	1.5364
urban.1000.Scaled	3.8120	1	1.9524
water.100.Scaled	1.7370	1	1.3180

Turtle mass (Mass) has the highest  $GVIF^{(1/(2*df))} > 2$ , so I will remove it and recalculate the factors.

```
mod.log.Start.vif <- lm(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.1000.Scaled + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.log.Start.vif)
```

##		GVIF	df	$GVIF^{(1/(2*df))}$
##	JulianDate.Scaled	2.6556	1	1.6296
##	A.Temp.Scaled	3.9633	1	1.9908
##	W.Temp.Scaled	5.7275	1	2.3932
##	Time.Scaled	1.1307	1	1.0633
##	PL.Scaled	1.3764	1	1.1732
##	Sex	1.3577	1	1.1652
##	for.veg.1000.Scaled	6.4136	1	2.5325
##	wet.400.Scaled	1.2936	1	1.1374
##	agri.400.Scaled	2.2981	1	1.5159
##	urban.1000.Scaled	3.8005	1	1.9495
##	water.100.Scaled	1.7329	1	1.3164

Proportion of forest and vegetation area at 1000m (for.veg.1000) has the highest  $GVIF^{(1/(2*df))} > 2$ , so I will remove it and recalculate the factors.

```
mod.log.Start.vif <- lm(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.log.Start.vif)
```

##		GVIF	df	$GVIF^{(1/(2*df))}$
##	JulianDate.Scaled	2.5793	1	1.6060
##	A.Temp.Scaled	3.9474	1	1.9868
##	W.Temp.Scaled	5.2164	1	2.2839
##	Time.Scaled	1.1188	1	1.0577
##	PL.Scaled	1.3362	1	1.1559
##	Sex	1.3547	1	1.1639
##	wet.400.Scaled	1.2931	1	1.1371
##	agri.400.Scaled	1.0550	1	1.0271
##	urban.1000.Scaled	1.1520	1	1.0733
##	water.100.Scaled	1.5273	1	1.2358

Water temperature (W.Temp) has the highest  $GVIF^{(1/(2*df))} > 2$ , so I will remove it and recalculate the factors.

```
mod.log.Start.vif <- lm(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.log.Start.vif)
```

```
##          GVIF df GVIF^(1/(2*df))
## JulianDate.Scaled 1.8283 1 1.3521
## A.Temp.Scaled 1.5302 1 1.2370
## Time.Scaled 1.1089 1 1.0530
## PL.Scaled 1.3620 1 1.1670
## Sex 1.3661 1 1.1688
## wet.400.Scaled 1.2598 1 1.1224
## agri.400.Scaled 1.0579 1 1.0285
## urban.1000.Scaled 1.1253 1 1.0608
## water.100.Scaled 1.4303 1 1.1960
```

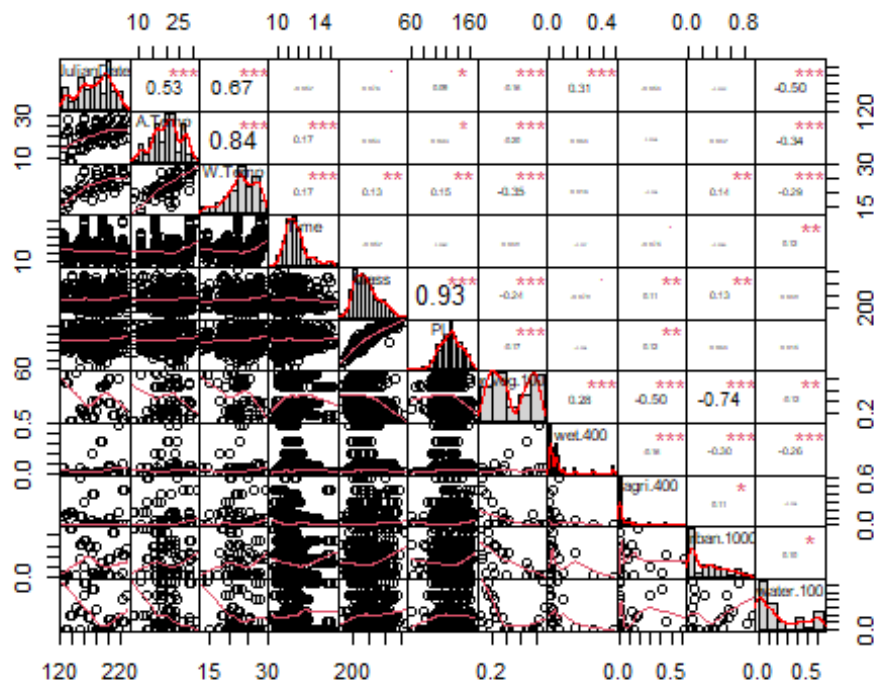
All of the  $GVIF^{1/(2*df)} < 2$ , so I will not remove any more variables.

## Calculation of Pearson and Spearman correlation coefficients

### Pearson correlation coefficients

#### # Visualization of the correlations

```
cor.pearson.log.Start <- MixedData[, c(3,4,5,6,7,8,26,30,40,56,57)]
chart.Correlation(cor.pearson.log.Start, histogram=TRUE, pch=19)
```



#### # Creation of the correlation table

```
table.corr.pearson.log.Start <- rcorr(as.matrix(MixedData[,c(3,4,5,6,7,8,26,30,40,56,57)]), type="pearson")
table.corr.pearson.log.Start.r <- table.corr.pearson.log.Start$r # Pearson correlation coefficients
table.corr.pearson.log.Start.p <- table.corr.pearson.log.Start$P # P values of
```

# the correlations

table.cor.pearson.log.Start.r

##	JulianDate	A.Temp	W.Temp	Time	Mas
S					
## JulianDate	1.000000000	0.526623600	0.66868310	-0.051688682	0.0756020
8					
## A.Temp	0.526623600	1.000000000	0.83887108	0.171967476	0.0525296
4					
## W.Temp	0.668683098	0.838871081	1.00000000	0.166126059	0.1322150
5					
## Time	-0.051688682	0.171967476	0.16612606	1.000000000	-0.0524517
1					
## Mass	0.075602082	0.052529643	0.13221505	-0.052451709	1.0000000
0					
## PL	0.089516177	0.092502829	0.14606475	0.009461487	0.9331145
5					
## for.veg.1000	-0.159757526	-0.202866655	-0.35442189	0.048125471	-0.2396708
7					
## wet.400	0.306093497	0.064315348	0.01560853	-0.036551862	-0.0781026
4					
## agri.400	-0.053734482	0.001830737	-0.02805362	-0.074905763	0.1149580
0					
## urban.1000	-0.003462429	0.042051110	0.14394243	-0.006102219	0.1273897
4					
## water.100	-0.496260565	-0.338053897	-0.29046345	0.117991743	0.0478384
1					
##	PL	for.veg.1000	wet.400	agri.400	urban.10
00					
## JulianDate	0.089516177	-0.15975753	0.30609350	-0.053734482	-0.0034624
29					
## A.Temp	0.092502829	-0.20286665	0.06431535	0.001830737	0.0420511
10					
## W.Temp	0.146064749	-0.35442189	0.01560853	-0.028053619	0.1439424
29					
## Time	0.009461487	0.04812547	-0.03655186	-0.074905763	-0.0061022
19					
## Mass	0.933114554	-0.23967087	-0.07810264	0.114958001	0.1273897
43					
## PL	1.000000000	-0.17433634	-0.03941912	0.116911690	0.0639451
49					
## for.veg.1000	-0.174336336	1.00000000	0.28348111	-0.495333110	-0.7437982
23					
## wet.400	-0.039419118	0.28348111	1.00000000	-0.161677421	-0.2975250
27					
## agri.400	0.116911690	-0.49533311	-0.16167742	1.000000000	0.1086598
50					
## urban.1000	0.063945149	-0.74379822	-0.29752503	0.108659850	1.0000000
00					
## water.100	0.015430905	-0.12313807	-0.26458013	-0.028765465	0.1035319

```

38
##                water.100
## JulianDate    -0.49626056
## A.Temp        -0.33805390
## W.Temp        -0.29046345
## Time          0.11799174
## Mass          0.04783841
## PL            0.01543090
## for.veg.1000 -0.12313807
## wet.400       -0.26458013
## agri.400      -0.02876546
## urban.1000    0.10353194
## water.100     1.00000000

```

table.cor.pearson.log.Start.p

```

##                JulianDate      A.Temp      W.Temp      Time      M
ass
## JulianDate      NA 0.000000e+00 0.000000e+00 2.322103e-01 8.033738e
-02
## A.Temp          0.000000e+00      NA 0.000000e+00 6.283828e-05 2.246911e
-01
## W.Temp          0.000000e+00 0.000000e+00      NA 2.346306e-04 3.499660e
-03
## Time            2.322103e-01 6.283828e-05 2.346306e-04      NA 2.253805e
-01
## Mass            8.033738e-02 2.246911e-01 3.499660e-03 2.253805e-01
NA
## PL              3.846835e-02 3.242105e-02 1.256393e-03 8.271659e-01 0.000000e
+00
## for.veg.1000    2.040904e-04 2.189891e-06 8.881784e-16 2.660399e-01 1.932793e
-08
## wet.400         4.334311e-13 1.369959e-01 7.314244e-01 3.983658e-01 7.080188e
-02
## agri.400        2.142232e-01 9.662709e-01 5.372464e-01 8.317038e-02 7.719600e
-03
## urban.1000      9.362578e-01 3.311946e-01 1.464223e-03 8.879110e-01 3.132312e
-03
## water.100       0.000000e+00 8.881784e-16 6.658052e-11 6.240147e-03 2.689054e
-01
##                PL for.veg.1000      wet.400      agri.400      urban.1
000
## JulianDate      3.846835e-02 2.040904e-04 4.334311e-13 0.2142231652 9.362578e
-01
## A.Temp          3.242105e-02 2.189891e-06 1.369959e-01 0.9662709322 3.311946e
-01
## W.Temp          1.256393e-03 8.881784e-16 7.314244e-01 0.5372463677 1.464223e
-03
## Time            8.271659e-01 2.660399e-01 3.983658e-01 0.0831703770 8.879110e
-01

```

```
## Mass          0.000000e+00 1.932793e-08 7.080188e-02 0.0077195996 3.132312e
-03
## PL              NA 5.033503e-05 3.628289e-01 0.0067865758 1.396467e
-01
## for.veg.1000 5.033503e-05              NA 2.303513e-11 0.0000000000 0.000000e
+00
## wet.400        3.628289e-01 2.303513e-11              NA 0.0001705096 2.036149e
-12
## agri.400       6.786576e-03 0.000000e+00 1.705096e-04              NA 1.182777e
-02
## urban.1000     1.396467e-01 0.000000e+00 2.036149e-12 0.0118277719
NA
## water.100      7.217625e-01 4.302738e-03 4.896181e-10 0.5063374531 1.649439e
-02
##              water.100
## JulianDate     0.000000e+00
## A.Temp          8.881784e-16
## W.Temp          6.658052e-11
## Time            6.240147e-03
## Mass            2.689054e-01
## PL              7.217625e-01
## for.veg.1000    4.302738e-03
## wet.400         4.896181e-10
## agri.400        5.063375e-01
## urban.1000     1.649439e-02
## water.100      NA
```

### *Spearman correlation coefficients*

*# Creation of the correlation table*

```
table.corr.spearman.log.Start <- rcorr(as.matrix(MixedData[,c(3,4,5,6,7,8,26,
30,40,56,57)]), type="spearman")
table.cor.spearman.log.Start.r <- table.corr.spearman.log.Start$r # Pearson c
orrelation coefficients
table.cor.spearman.log.Start.p <- table.corr.spearman.log.Start$P # P values
of the correlations
table.cor.spearman.log.Start.r
```

```
##          JulianDate      A.Temp      W.Temp      Time      Mass
## JulianDate    1.00000000  0.45842169  0.56884913 -0.09892087  0.08038929
## A.Temp        0.45842169  1.00000000  0.83718368  0.13849890  0.06589666
## W.Temp        0.56884913  0.83718368  1.00000000  0.17365903  0.13270468
## Time          -0.09892087  0.13849890  0.17365903  1.00000000 -0.04025948
## Mass          0.08038929  0.06589666  0.13270468 -0.04025948  1.00000000
## PL            0.08898378  0.10442546  0.15193967 -0.01364455  0.94856246
## for.veg.1000 -0.13614271 -0.17001898 -0.28928407  0.11957044 -0.19217208
## wet.400       0.41005248  0.11243977  0.19326625  0.14639174 -0.04217241
## agri.400      -0.13443891  0.19719332  0.07363333 -0.09493393  0.14999256
## urban.1000    0.05728527  0.13104537  0.21782810 -0.07145219  0.10868203
## water.100     -0.40716475 -0.37402320 -0.28326055  0.05606805  0.07784216
##              PL for.veg.1000      wet.400      agri.400      urban.1000
```



```

## JulianDate      0.08898378  -0.1361427  0.41005248 -0.13443891  0.05728527
## A.Temp          0.10442546  -0.1700190  0.11243977  0.19719332  0.13104537
## W.Temp          0.15193967  -0.2892841  0.19326625  0.07363333  0.21782810
## Time           -0.01364455   0.1195704  0.14639174 -0.09493393 -0.07145219
## Mass           0.94856246  -0.1921721 -0.04217241  0.14999256  0.10868203
## PL             1.00000000  -0.1623624 -0.04821271  0.16970352  0.07998962
## for.veg.1000 -0.16236237   1.0000000  0.24464327 -0.62367762 -0.82572013
## wet.400        -0.04821271   0.2446433  1.00000000 -0.30522895 -0.23226774
## agri.400       0.16970352  -0.6236776 -0.30522895  1.00000000  0.46271885
## urban.1000     0.07998962  -0.8257201 -0.23226774  0.46271885  1.00000000
## water.100      0.03022562  -0.1316108 -0.14172001 -0.05751472  0.08587093
##               water.100
## JulianDate     -0.40716475
## A.Temp         -0.37402320
## W.Temp         -0.28326055
## Time           0.05606805
## Mass           0.07784216
## PL             0.03022562
## for.veg.1000 -0.13161079
## wet.400        -0.14172001
## agri.400       -0.05751472
## urban.1000     0.08587093
## water.100      1.00000000

```

table.cor.spearman.log.Start.p

```

##               JulianDate      A.Temp      W.Temp      Time      Ma
ss
## JulianDate      NA 0.000000e+00 0.000000e+00 0.0219948308 6.291206e-
02
## A.Temp          0.000000000      NA 0.000000e+00 0.0013067179 1.275805e-
01
## W.Temp          0.000000000 0.000000e+00      NA 0.0001191832 3.379071e-
03
## Time           0.021994831 1.306718e-03 1.191832e-04      NA 3.522285e-
01
## Mass           0.062912061 1.275805e-01 3.379071e-03 0.3522284632
NA
## PL             0.039641066 1.567793e-02 7.879312e-04 0.7528559899 0.000000e+
00
## for.veg.1000 0.001581516 7.625616e-05 8.007350e-11 0.0055757019 7.445223e-
06
## wet.400        0.000000000 9.177827e-03 1.785318e-05 0.0006747625 3.298003e-
01
## agri.400       0.001812250 4.226134e-06 1.049537e-01 0.0279695118 4.936074e-
04
## urban.1000     0.185422901 2.366128e-03 1.246624e-06 0.0984350178 1.181043e-
02
## water.100      0.000000000 0.000000e+00 2.027840e-10 0.1949553499 7.174987e-
02

```

```

##                                PL for.veg.1000      wet.400      agri.400      urban.1
000
## JulianDate      3.964107e-02 1.581516e-03 0.000000e+00 1.812250e-03 1.854229e
-01
## A.Temp          1.567793e-02 7.625616e-05 9.177827e-03 4.226134e-06 2.366128e
-03
## W.Temp          7.879312e-04 8.007350e-11 1.785318e-05 1.049537e-01 1.246624e
-06
## Time            7.528560e-01 5.575702e-03 6.747625e-04 2.796951e-02 9.843502e
-02
## Mass            0.000000e+00 7.445223e-06 3.298003e-01 4.936074e-04 1.181043e
-02
## PL              NA 1.621239e-04 2.656216e-01 7.989296e-05 6.448653e
-02
## for.veg.1000    1.621239e-04      NA 9.574610e-09 0.000000e+00 0.000000e
+00
## wet.400         2.656216e-01 9.574610e-09      NA 5.080381e-13 5.346181e
-08
## agri.400        7.989296e-05 0.000000e+00 5.080381e-13      NA 0.000000e
+00
## urban.1000      6.448653e-02 0.000000e+00 5.346181e-08 0.000000e+00
NA
## water.100       4.853994e-01 2.264245e-03 1.001819e-03 1.836652e-01 4.691287e
-02
##                                water.100
## JulianDate      0.000000e+00
## A.Temp          0.000000e+00
## W.Temp          2.027840e-10
## Time            1.949553e-01
## Mass            7.174987e-02
## PL              4.853994e-01
## for.veg.1000    2.264245e-03
## wet.400         1.001819e-03
## agri.400        1.836652e-01
## urban.1000      4.691287e-02
## water.100       NA

```

The Pearson correlation coefficient of 0.93 between Mass and PL, -0.74 between for.veg.1000 and urban.1000, and 0.84 between W.Temp and A.Temp confirmed the deletion of Mass, for.veg.1000, and W.Temp with the calculation of the  $GVIF^{(1/(2*df))}$ .

#### Verification of the assumptions with the initial model

```

mod.log.Start.full <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Ti
me.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.S
caled + water.100.Scaled + (1|Code) + (1|Site), data = MixedData, REML = TRUE
, na.action=na.exclude)

```

```

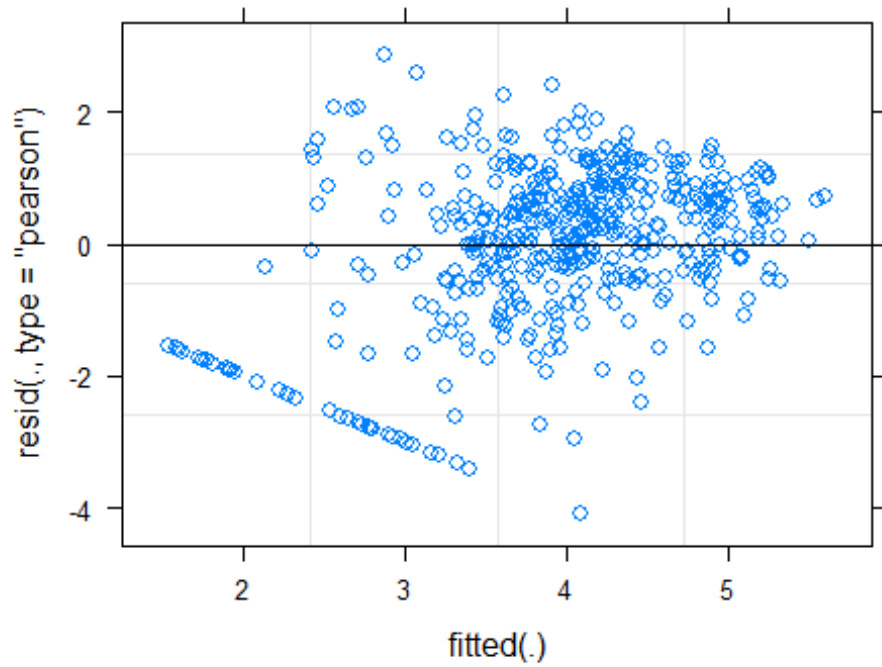
summary(mod.log.Start.full)

```

```

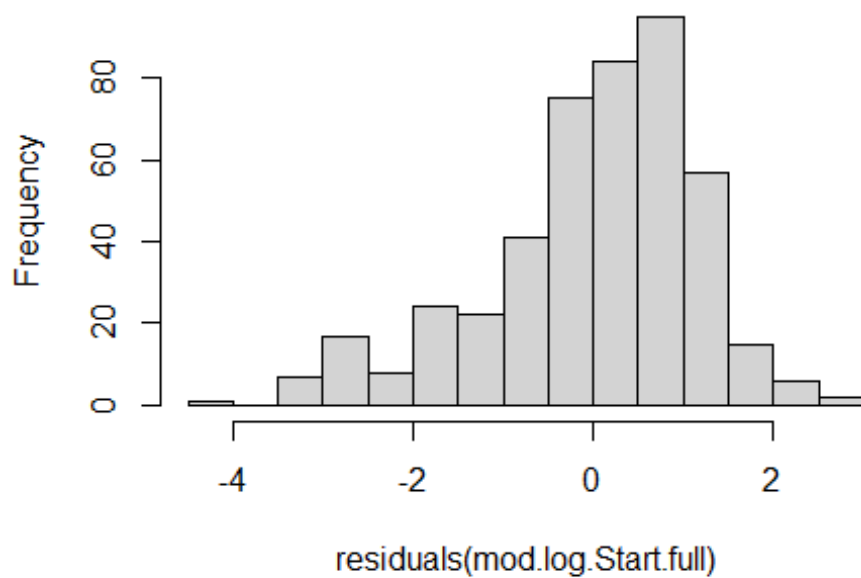
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
##      PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled
##      +
##      water.100.Scaled + (1 | Code) + (1 | Site)
##      Data: MixedData
##
## REML criterion at convergence: 1684
##
## Scaled residuals:
##      Min      1Q  Median      3Q      Max
## -3.0644 -0.3958  0.1451  0.6178  2.1520
##
## Random effects:
##      Groups      Name      Variance Std.Dev.
##      Code      (Intercept) 0.4099   0.6402
##      Site      (Intercept) 0.3127   0.5592
##      Residual              1.7741   1.3320
## Number of obs: 454, groups:  Code, 430; Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    4.19998    0.17362   29.34991   24.191   <2e-16 ***
## JulianDate.Scaled -0.19806    0.16170   21.16973   -1.225    0.2341
## A.Temp.Scaled   -0.19874    0.12576  100.73408   -1.580    0.1172
## Time.Scaled      0.07622    0.07781  439.62923    0.980    0.3278
## PL.Scaled        0.21939    0.09145  410.59234    2.399    0.0169 *
## SexM             -0.24429    0.16850  441.25735   -1.450    0.1478
## wet.400.Scaled   0.22219    0.15373   14.33551    1.445    0.1699
## agri.400.Scaled  0.14719    0.11940   18.93332    1.233    0.2327
## urban.1000.Scaled -0.11470    0.14520   15.38532   -0.790    0.4416
## water.100.Scaled -0.25738    0.16186   14.62996   -1.590    0.1332
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnDt.S A.Tm.S Tm.Scl PL.Scl SexM   w.400. a.400. u.1000
## JulnDt.Scl -0.070
## A.Temp.Scl  0.112 -0.395
## Time.Scaled 0.044  0.084 -0.237
## PL.Scaled   -0.290 -0.046 -0.047 -0.034
## SexM        -0.564  0.038 -0.110 -0.053  0.472
## wt.400.Scl -0.056 -0.251  0.125 -0.016  0.045  0.004
## agr.400.Scl -0.109  0.069 -0.006  0.041 -0.044 -0.034  0.163
## urbn.1000.S -0.089 -0.165  0.122  0.028  0.024  0.046  0.329 -0.014
## wtr.100.Scl  0.030  0.166  0.123 -0.075 -0.056 -0.047  0.243  0.083  0.011
plot(mod.log.Start.full)

```

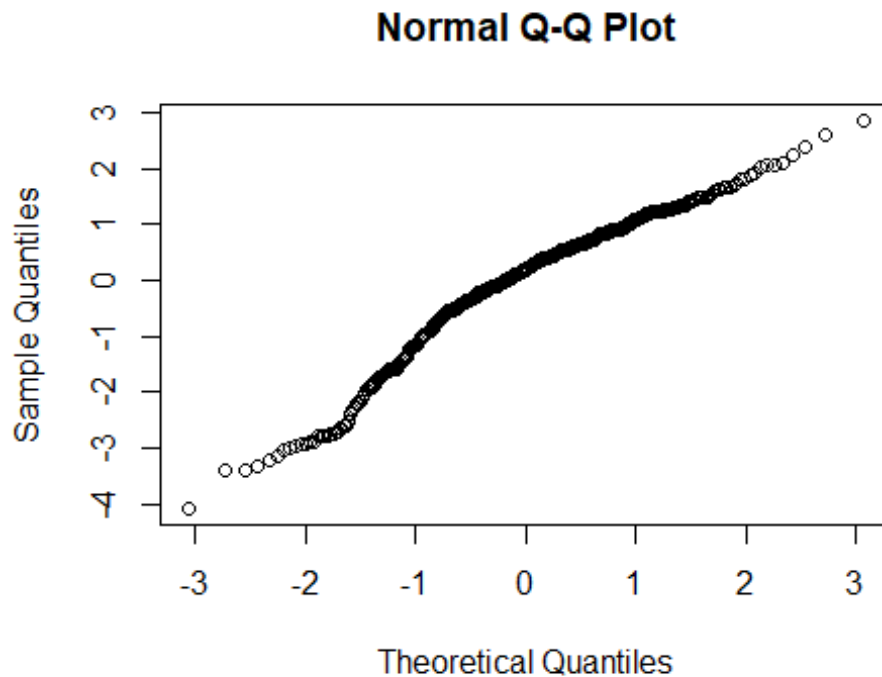


```
hist(residuals(mod.log.Start.full))
```

**Histogram of residuals(mod.log.Start.full)**



```
qqnorm(resid(mod.log.Start.full))
```



## Time of shell emergence (Bin.Shell)

### Multicollinearity: Generalized variance inflation factor ( $\text{GVIF}^{1/(2 \cdot \text{df})}$ )

$\text{GVIF}^{1/(2 \cdot \text{df})} > 2$  indicates the presence of multicollinearity, so I will remove variables with values over 2, starting with the highest value.

```
mod.Bin.Shell.vif <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled + Time.Scaled + Mass.Scaled + PL.Scaled + Sex + for.veg.1000.Scaled + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled, data = MixedData, family = binomial, na.action=na.exclude)
gvif(mod.Bin.Shell.vif)
```

##		GVIF	df	$\text{GVIF}^{1/(2 \cdot \text{df})}$
##	JulianDate.Scaled	2.2335	1	1.4945
##	A.Temp.Scaled	3.2566	1	1.8046
##	W.Temp.Scaled	4.6305	1	2.1519
##	Time.Scaled	1.0978	1	1.0478
##	Mass.Scaled	9.4050	1	3.0668
##	PL.Scaled	8.4056	1	2.8992
##	Sex	1.4693	1	1.2121
##	for.veg.1000.Scaled	76.3505	1	8.7379
##	wet.300.Scaled	3.7945	1	1.9479
##	agri.1000.Scaled	24.1413	1	4.9134
##	urban.1000.Scaled	60.1982	1	7.7587
##	water.900.Scaled	12.7235	1	3.5670

Proportion of forest and vegetation area at 1000m (for.veg.1000) has the highest  $GVIF^{1/(2*df)} > 2$ , so I will remove it and recalculate the factors.

```
mod.Bin.Shell.vif <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled + Time.Scaled + Mass.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled, data = MixedData, family = binomial, na.action=na.exclude)
gvif(mod.Bin.Shell.vif)
```

##		GVIF	df	$GVIF^{1/(2*df)}$
##	JulianDate.Scaled	2.2084	1	1.4861
##	A.Temp.Scaled	3.2449	1	1.8014
##	W.Temp.Scaled	4.5909	1	2.1426
##	Time.Scaled	1.0962	1	1.0470
##	Mass.Scaled	9.3543	1	3.0585
##	PL.Scaled	8.3520	1	2.8900
##	Sex	1.4499	1	1.2041
##	wet.300.Scaled	1.4939	1	1.2223
##	agri.1000.Scaled	1.1970	1	1.0941
##	urban.1000.Scaled	1.3446	1	1.1596
##	water.900.Scaled	1.2665	1	1.1254

Turtle mass (Mass) has the highest  $GVIF^{1/(2*df)} > 2$ , so I will remove it and recalculate the factors.

```
mod.Bin.Shell.vif <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled, data = MixedData, family = binomial, na.action=na.exclude)
gvif(mod.Bin.Shell.vif)
```

##		GVIF	df	$GVIF^{1/(2*df)}$
##	JulianDate.Scaled	2.1874	1	1.4790
##	A.Temp.Scaled	3.2392	1	1.7998
##	W.Temp.Scaled	4.5708	1	2.1379
##	Time.Scaled	1.0857	1	1.0420
##	PL.Scaled	1.4000	1	1.1832
##	Sex	1.3525	1	1.1630
##	wet.300.Scaled	1.4926	1	1.2217
##	agri.1000.Scaled	1.1971	1	1.0941
##	urban.1000.Scaled	1.2621	1	1.1234
##	water.900.Scaled	1.2191	1	1.1041

Water temperature (W.Temp) has the highest  $GVIF^{1/(2*df)} > 2$ , so I will remove it and recalculate the factors.

```
mod.Bin.Shell.vif <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled, data = MixedData, family = binomial, na.action=na.exclude)
gvif(mod.Bin.Shell.vif)
```

```
##          GVIF df GVIF^(1/(2*df))
## JulianDate.Scaled 1.6023 1 1.2658
## A.Temp.Scaled 1.4137 1 1.1890
## Time.Scaled 1.0726 1 1.0357
## PL.Scaled 1.4134 1 1.1889
## Sex 1.3611 1 1.1667
## wet.300.Scaled 1.4687 1 1.2119
## agri.1000.Scaled 1.2172 1 1.1033
## urban.1000.Scaled 1.2138 1 1.1017
## water.900.Scaled 1.1263 1 1.0613
```

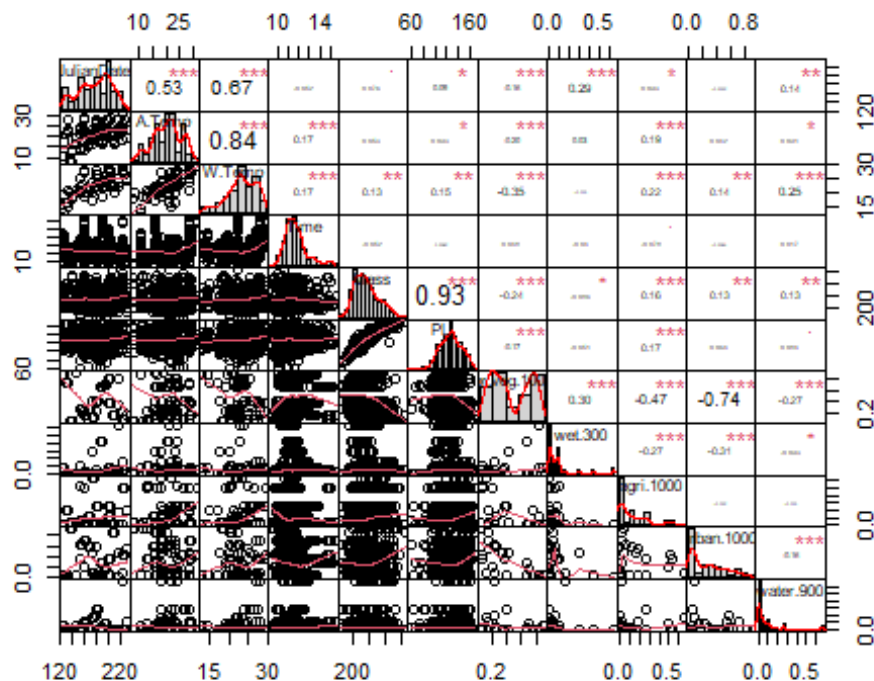
All of the  $GVIF^{1/(2*df)} < 2$ , so I will not remove any more variables.

## Calculation of Pearson and Spearman correlation coefficients

### Pearson correlation coefficients

#### # Visualization of the correlations

```
cor.pearson.Bin.Shell <- MixedData[, c(3,4,5,6,7,8,26,29,46,56,65)]
chart.Correlation(cor.pearson.Bin.Shell, histogram=TRUE, pch=19)
```



#### # Creation of the correlation table

```
table.corr.pearson.Bin.Shell <- rcorr(as.matrix(MixedData[,c(3,4,5,6,7,8,26,29,46,56,65)]), type="pearson")
table.corr.pearson.Bin.Shell.r <- table.corr.pearson.Bin.Shell$r # Pearson correlation coefficients
table.corr.pearson.Bin.Shell.p <- table.corr.pearson.Bin.Shell$P # P values of
```

### the correlations

table.cor.pearson.Bin.Shell.r

##	JulianDate	A.Temp	W.Temp	Time	Mass
## JulianDate	1.000000000	0.52662360	0.66868310	-0.051688682	0.07560208
## A.Temp	0.526623600	1.000000000	0.83887108	0.171967476	0.05252964
## W.Temp	0.668683098	0.83887108	1.000000000	0.166126059	0.13221505
## Time	-0.051688682	0.17196748	0.16612606	1.000000000	-0.05245171
## Mass	0.075602082	0.05252964	0.13221505	-0.052451709	1.000000000
## PL	0.089516177	0.09250283	0.14606475	0.009461487	0.93311455
## for.veg.1000	-0.159757526	-0.20286665	-0.35442189	0.048125471	-0.23967087
## wet.300	0.285505234	0.02957879	-0.02150728	-0.039628347	-0.08629485
## agri.1000	0.093070389	0.19452574	0.21528845	-0.078150322	0.15949820
## urban.1000	-0.003462429	0.04205111	0.14394243	-0.006102219	0.12738974
## water.900	0.140565598	0.09143754	0.25309673	0.016579901	0.13233284
##	PL	for.veg.1000	wet.300	agri.1000	urban.1000
0					
## JulianDate	0.089516177	-0.15975753	0.28550523	0.09307039	-0.00346242
9					
## A.Temp	0.092502829	-0.20286665	0.02957879	0.19452574	0.04205111
0					
## W.Temp	0.146064749	-0.35442189	-0.02150728	0.21528845	0.14394242
9					
## Time	0.009461487	0.04812547	-0.03962835	-0.07815032	-0.00610221
9					
## Mass	0.933114554	-0.23967087	-0.08629485	0.15949820	0.12738974
3					
## PL	1.000000000	-0.17433634	-0.05147288	0.16850474	0.06394514
9					
## for.veg.1000	-0.174336336	1.000000000	0.30140143	-0.47018472	-0.74379822
3					
## wet.300	-0.051472876	0.30140143	1.000000000	-0.27005276	-0.30777568
4					
## agri.1000	0.168504745	-0.47018472	-0.27005276	1.000000000	-0.03513526
8					
## urban.1000	0.063945149	-0.74379822	-0.30777568	-0.03513527	1.000000000
0					
## water.900	0.083709114	-0.26815368	-0.09294993	-0.02285347	-0.16249035
8					
##	water.900				
## JulianDate	0.14056560				
## A.Temp	0.09143754				
## W.Temp	0.25309673				
## Time	0.01657990				
## Mass	0.13233284				
## PL	0.08370911				
## for.veg.1000	-0.26815368				
## wet.300	-0.09294993				
## agri.1000	-0.02285347				



```
## urban.1000    -0.16249036
## water.900     1.00000000
```

```
table.cor.pearson.Bin.Shell.p
```

```
##          JulianDate      A.Temp      W.Temp      Time      M
ass
## JulianDate      NA 0.000000e+00 0.000000e+00 2.322103e-01 8.033738e
-02
## A.Temp      0.000000e+00      NA 0.000000e+00 6.283828e-05 2.246911e
-01
## W.Temp      0.000000e+00 0.000000e+00      NA 2.346306e-04 3.499660e
-03
## Time      2.322103e-01 6.283828e-05 2.346306e-04      NA 2.253805e
-01
## Mass      8.033738e-02 2.246911e-01 3.499660e-03 2.253805e-01
NA
## PL      3.846835e-02 3.242105e-02 1.256393e-03 8.271659e-01 0.000000e
+00
## for.veg.1000 2.040904e-04 2.189891e-06 8.881784e-16 2.660399e-01 1.932793e
-08
## wet.300      1.637268e-11 4.943861e-01 6.362342e-01 3.598346e-01 4.583253e
-02
## agri.1000      3.121016e-02 5.719872e-06 1.666280e-06 7.062947e-02 2.090744e
-04
## urban.1000      9.362578e-01 3.311946e-01 1.464223e-03 8.879110e-01 3.132312e
-03
## water.900      1.102607e-03 3.430866e-02 1.534812e-08 7.017326e-01 2.139968e
-03
##          PL for.veg.1000      wet.300      agri.1000      urban.1
000
## JulianDate      3.846835e-02 2.040904e-04 1.637268e-11 3.121016e-02 9.362578e
-01
## A.Temp      3.242105e-02 2.189891e-06 4.943861e-01 5.719872e-06 3.311946e
-01
## W.Temp      1.256393e-03 8.881784e-16 6.362342e-01 1.666280e-06 1.464223e
-03
## Time      8.271659e-01 2.660399e-01 3.598346e-01 7.062947e-02 8.879110e
-01
## Mass      0.000000e+00 1.932793e-08 4.583253e-02 2.090744e-04 3.132312e
-03
## PL      NA 5.033503e-05 2.346072e-01 8.986288e-05 1.396467e
-01
## for.veg.1000 5.033503e-05      NA 1.017852e-12 0.000000e+00 0.000000e
+00
## wet.300      2.346072e-01 1.017852e-12      NA 2.070131e-10 3.179679e
-13
## agri.1000      8.986288e-05 0.000000e+00 2.070131e-10      NA 4.169111e
-01
## urban.1000      1.396467e-01 0.000000e+00 3.179679e-13 4.169111e-01
```

```

NA
## water.900      5.298219e-02 2.797003e-10 3.143027e-02 5.975481e-01 1.579156e
-04
##
## JulianDate    1.102607e-03
## A.Temp        3.430866e-02
## W.Temp        1.534812e-08
## Time          7.017326e-01
## Mass          2.139968e-03
## PL            5.298219e-02
## for.veg.1000  2.797003e-10
## wet.300       3.143027e-02
## agri.1000     5.975481e-01
## urban.1000    1.579156e-04
## water.900     NA

```

### *Spearman correlation coefficients*

*# Creation of the correlation table*

```

table.corr.spearman.Bin.Shell <- rcorr(as.matrix(MixedData[,c(3,4,5,6,7,8,26,
29,46,56,65)]), type="spearman")
table.cor.spearman.Bin.Shell.r <- table.corr.spearman.Bin.Shell$r # Pearson c
orrelation coefficients
table.cor.spearman.Bin.Shell.p <- table.corr.spearman.Bin.Shell$P # P values
of the correlations
table.cor.spearman.Bin.Shell.r

```

##	JulianDate	A.Temp	W.Temp	Time	Mass
## JulianDate	1.00000000	0.458421690	0.56884913	-0.09892087	0.08038929
## A.Temp	0.45842169	1.000000000	0.83718368	0.13849890	0.06589666
## W.Temp	0.56884913	0.837183684	1.00000000	0.17365903	0.13270468
## Time	-0.09892087	0.138498905	0.17365903	1.00000000	-0.04025948
## Mass	0.08038929	0.065896659	0.13270468	-0.04025948	1.00000000
## PL	0.08898378	0.104425464	0.15193967	-0.01364455	0.94856246
## for.veg.1000	-0.13614271	-0.170018984	-0.28928407	0.11957044	-0.19217208
## wet.300	0.32499215	-0.007493503	0.05630636	0.12226733	-0.07340554
## agri.1000	0.14975219	0.307433887	0.33671601	-0.12473748	0.18211450
## urban.1000	0.05728527	0.131045369	0.21782810	-0.07145219	0.10868203
## water.900	-0.23064372	-0.101001782	-0.01834662	0.03769977	0.04847697
##	PL	for.veg.1000	wet.300	agri.1000	urban.1000
## JulianDate	0.08898378	-0.1361427	0.324992150	0.1497522	0.05728527
## A.Temp	0.10442546	-0.1700190	-0.007493503	0.3074339	0.13104537
## W.Temp	0.15193967	-0.2892841	0.056306361	0.3367160	0.21782810
## Time	-0.01364455	0.1195704	0.122267329	-0.1247375	-0.07145219
## Mass	0.94856246	-0.1921721	-0.073405540	0.1821145	0.10868203
## PL	1.00000000	-0.1623624	-0.088758890	0.1885881	0.07998962
## for.veg.1000	-0.16236237	1.0000000	0.291174883	-0.5429549	-0.82572013
## wet.300	-0.08875889	0.2911749	1.000000000	-0.3173058	-0.32221448
## agri.1000	0.18858810	-0.5429549	-0.317305773	1.0000000	0.33862008
## urban.1000	0.07998962	-0.8257201	-0.322214475	0.3386201	1.00000000
## water.900	0.03358886	-0.2295566	-0.387776581	-0.1408985	0.06336506

```
##          water.900
## JulianDate -0.23064372
## A.Temp     -0.10100178
## W.Temp     -0.01834662
## Time       0.03769977
## Mass       0.04847697
## PL         0.03358886
## for.veg.1000 -0.22955661
## wet.300    -0.38777658
## agri.1000  -0.14089853
## urban.1000 0.06336506
## water.900  1.00000000
```

table.cor.spearman.Bin.Shell.p

```
##          JulianDate      A.Temp      W.Temp      Time      M
ass
## JulianDate      NA 0.000000e+00 0.000000e+00 0.0219948308 6.291206e
-02
## A.Temp          0.000000e+00      NA 0.000000e+00 0.0013067179 1.275805e
-01
## W.Temp          0.000000e+00 0.000000e+00      NA 0.0001191832 3.379071e
-03
## Time           2.199483e-02 1.306718e-03 1.191832e-04      NA 3.522285e
-01
## Mass           6.291206e-02 1.275805e-01 3.379071e-03 0.3522284632
NA
## PL             3.964107e-02 1.567793e-02 7.879312e-04 0.7528559899 0.000000e
+00
## for.veg.1000 1.581516e-03 7.625616e-05 8.007350e-11 0.0055757019 7.445223e
-06
## wet.300        1.199041e-14 8.625850e-01 2.153146e-01 0.0045864816 8.954843e
-02
## agri.1000      5.041266e-04 3.388401e-13 2.398082e-14 0.0038225745 2.216001e
-05
## urban.1000     1.854229e-01 2.366128e-03 1.246624e-06 0.0984350178 1.181043e
-02
## water.900      6.652959e-08 1.934139e-02 6.866163e-01 0.3837078901 2.625597e
-01
##              PL for.veg.1000      wet.300      agri.1000      urban.1
000
## JulianDate     3.964107e-02 1.581516e-03 1.199041e-14 5.041266e-04 1.854229e
-01
## A.Temp         1.567793e-02 7.625616e-05 8.625850e-01 3.388401e-13 2.366128e
-03
## W.Temp         7.879312e-04 8.007350e-11 2.153146e-01 2.398082e-14 1.246624e
-06
## Time           7.528560e-01 5.575702e-03 4.586482e-03 3.822575e-03 9.843502e
-02
## Mass           0.000000e+00 7.445223e-06 8.954843e-02 2.216001e-05 1.181043e
```

```

-02
## PL NA 1.621239e-04 4.014546e-02 1.126664e-05 6.448653e
-02
## for.veg.1000 1.621239e-04 NA 6.200374e-12 0.000000e+00 0.000000e
+00
## wet.300 4.014546e-02 6.200374e-12 NA 5.284662e-14 2.042810e
-14
## agri.1000 1.126664e-05 0.000000e+00 5.284662e-14 NA 8.881784e
-16
## urban.1000 6.448653e-02 0.000000e+00 2.042810e-14 8.881784e-16
NA
## water.900 4.381540e-01 7.694788e-08 0.000000e+00 1.072620e-03 1.429059e
-01
## water.900
## JulianDate 6.652959e-08
## A.Temp 1.934139e-02
## W.Temp 6.866163e-01
## Time 3.837079e-01
## Mass 2.625597e-01
## PL 4.381540e-01
## for.veg.1000 7.694788e-08
## wet.300 0.000000e+00
## agri.1000 1.072620e-03
## urban.1000 1.429059e-01
## water.900 NA

```

The Pearson correlation coefficient of -0.74 between for.veg.1000 and urban.1000, 0.93 between Mass and PL, and 0.84 between W.Temp and A.Temp confirmed the deletion of for.veg.1000, Mass, and W.Temp with the calculation of the  $GVIF^{(1/(2*df))}$ .

### Verification of the assumptions with the initial model

```

mod.Bin.Shell.full <- glmer(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1|Site) + (1|Code), data = MixedData, family = binomial, na.action=na.exclude)

```

```

summary(mod.Bin.Shell.full)

```

```

## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
## PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled +
## water.900.Scaled + (1 | Site) + (1 | Code)
## Data: MixedData
##
## AIC      BIC    logLik deviance df.resid
## 637.6    688.3   -306.8    613.6     494
##

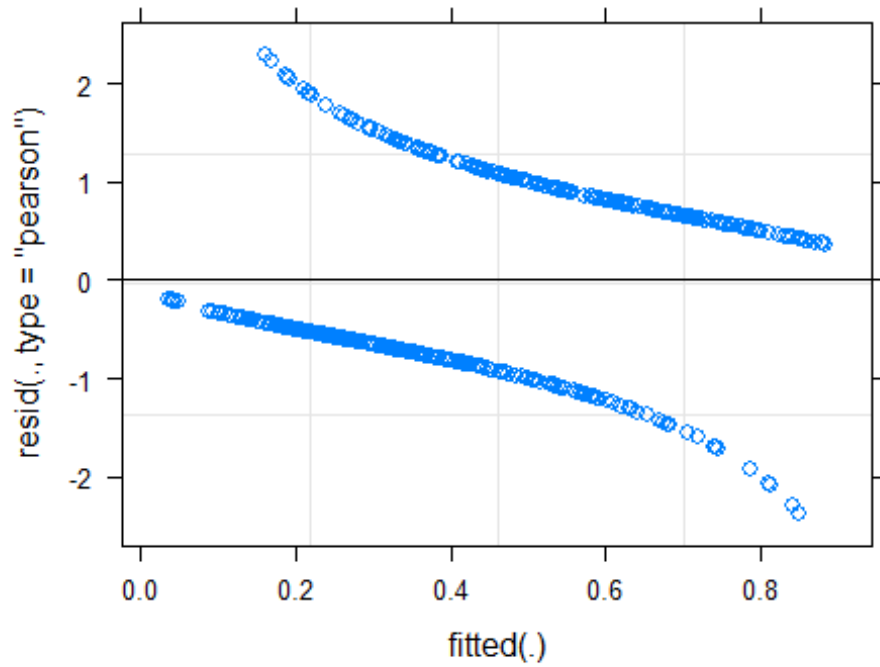
```

```

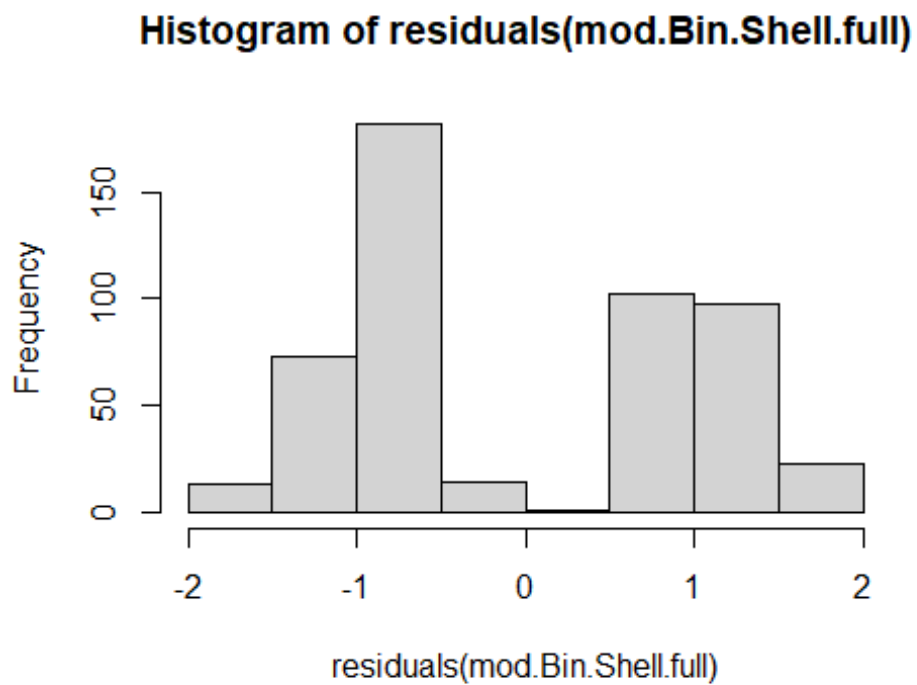
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3731 -0.6924 -0.4231  0.7804  2.2883
##
## Random effects:
##   Groups Name      Variance Std.Dev.
##   Code   (Intercept) 0.394958 0.62846
##   Site   (Intercept) 0.002675 0.05172
## Number of obs: 506, groups:  Code, 478; Site, 23
##
## Fixed effects:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   -0.273471   0.188311  -1.452 0.146437
## JulianDate.Scaled -0.519502   0.147058  -3.533 0.000411 ***
## A.Temp.Scaled    -0.444253   0.143668  -3.092 0.001987 **
## Time.Scaled      -0.308826   0.121947  -2.532 0.011326 *
## PL.Scaled        0.225610   0.138687   1.627 0.103787
## SexM             -0.074558   0.248956  -0.299 0.764571
## wet.300.Scaled   0.244893   0.125513   1.951 0.051041 .
## agri.1000.Scaled 0.001697   0.116766   0.015 0.988404
## urban.1000.Scaled -0.264439   0.121799  -2.171 0.029924 *
## water.900.Scaled -0.016795   0.112129  -0.150 0.880939
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S Tm.Scl PL.Scl SexM   w.300. a.1000 u.1000
## JulnDt.Scld -0.079
## A.Temp.Scld  0.177 -0.369
## Time.Scaled  0.166  0.159 -0.058
## PL.Scaled    -0.464 -0.038 -0.170 -0.099
## SexM         -0.779  0.119 -0.074 -0.090  0.476
## wt.300.Scld -0.084 -0.366 -0.016  0.010  0.094  0.028
## agr.1000.Sc -0.014 -0.035 -0.141  0.150 -0.109  0.003  0.341
## urbn.1000.S -0.027 -0.035  0.071  0.122 -0.062  0.086  0.330  0.184
## wtr.900.Scl  0.024 -0.151 -0.001 -0.016 -0.129 -0.044  0.221  0.105  0.232
## optimizer (Nelder_Mead) convergence code: 0 (OK)
## Model failed to converge with max|grad| = 0.025219 (tol = 0.002, component
1)

plot(mod.Bin.Shell.full)

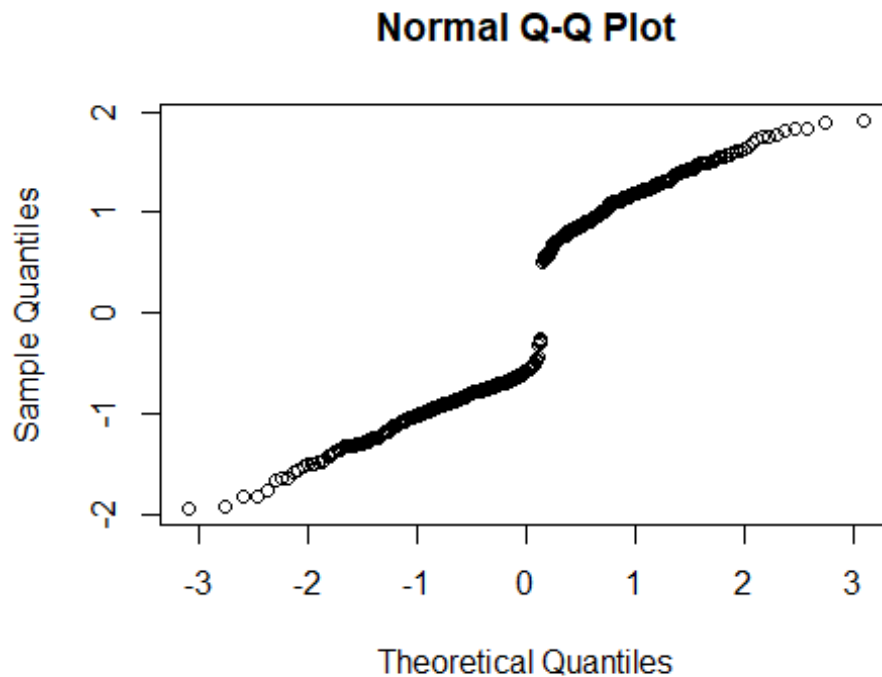
```



```
hist(residuals(mod.Bin.Shell.full))
```



```
qqnorm(resid(mod.Bin.Shell.full))
```



## Total time spent moving (Move)

### Multicollinearity: Generalized variance inflation factor ( $\text{GVIF}^{1/(2 \cdot \text{df})}$ )

$\text{GVIF}^{1/(2 \cdot \text{df})} > 2$  indicates the presence of multicollinearity, so I will remove variables with values over 2, starting with the highest value.

```
mod.Move.vif <- lm(Move ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled +
  Time.Scaled + Mass.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Sc
  aled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled, data = MixedD
  ata, na.action=na.exclude)
gvif(mod.Move.vif)
```

##		GVIF	df	$\text{GVIF}^{1/(2 \cdot \text{df})}$
##	JulianDate.Scaled	2.1511	1	1.4667
##	A.Temp.Scaled	4.0975	1	2.0242
##	W.Temp.Scaled	5.8063	1	2.4096
##	Time.Scaled	1.1339	1	1.0648
##	Mass.Scaled	9.0248	1	3.0041
##	PL.Scaled	7.9817	1	2.8252
##	Sex	1.4918	1	1.2214
##	for.veg.300.Scaled	3.7113	1	1.9265
##	wet.600.Scaled	1.9893	1	1.4104
##	agri.100.Scaled	1.5977	1	1.2640
##	urban.1000.Scaled	3.1887	1	1.7857
##	water.1000.Scaled	2.2055	1	1.4851

Turtle mass (Mass) has the highest  $GVIF^{1/(2*df)} > 2$ , so I will remove it and recalculate the factors.

```
mod.Move.vif <- lm(Move ~ JulianDate.Scaled + A.Temp.Scaled + W.Temp.Scaled +
Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.Move.vif)
```

##		GVIF	df	$GVIF^{1/(2*df)}$
##	JulianDate.Scaled	2.1479	1	1.4656
##	A.Temp.Scaled	4.0531	1	2.0132
##	W.Temp.Scaled	5.7873	1	2.4057
##	Time.Scaled	1.1162	1	1.0565
##	PL.Scaled	1.3618	1	1.1670
##	Sex	1.3759	1	1.1730
##	for.veg.300.Scaled	3.7113	1	1.9265
##	wet.600.Scaled	1.9893	1	1.4104
##	agri.100.Scaled	1.5920	1	1.2617
##	urban.1000.Scaled	3.1226	1	1.7671
##	water.1000.Scaled	2.1404	1	1.4630

Water temperature (W.Temp) has the highest  $GVIF^{1/(2*df)} > 2$ , so I will remove it and recalculate the factors.

```
mod.Move.vif <- lm(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + P
L.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urba
n.1000.Scaled + water.1000.Scaled, data = MixedData, na.action=na.exclude)
gvif(mod.Move.vif)
```

##		GVIF	df	$GVIF^{1/(2*df)}$
##	JulianDate.Scaled	1.6927	1	1.3010
##	A.Temp.Scaled	1.5492	1	1.2447
##	Time.Scaled	1.1070	1	1.0521
##	PL.Scaled	1.3734	1	1.1719
##	Sex	1.3747	1	1.1725
##	for.veg.300.Scaled	3.2982	1	1.8161
##	wet.600.Scaled	1.8972	1	1.3774
##	agri.100.Scaled	1.5658	1	1.2513
##	urban.1000.Scaled	2.5930	1	1.6103
##	water.1000.Scaled	1.8443	1	1.3581

All of the  $GVIF^{1/(2*df)} < 2$ , so I will not remove any more variables.

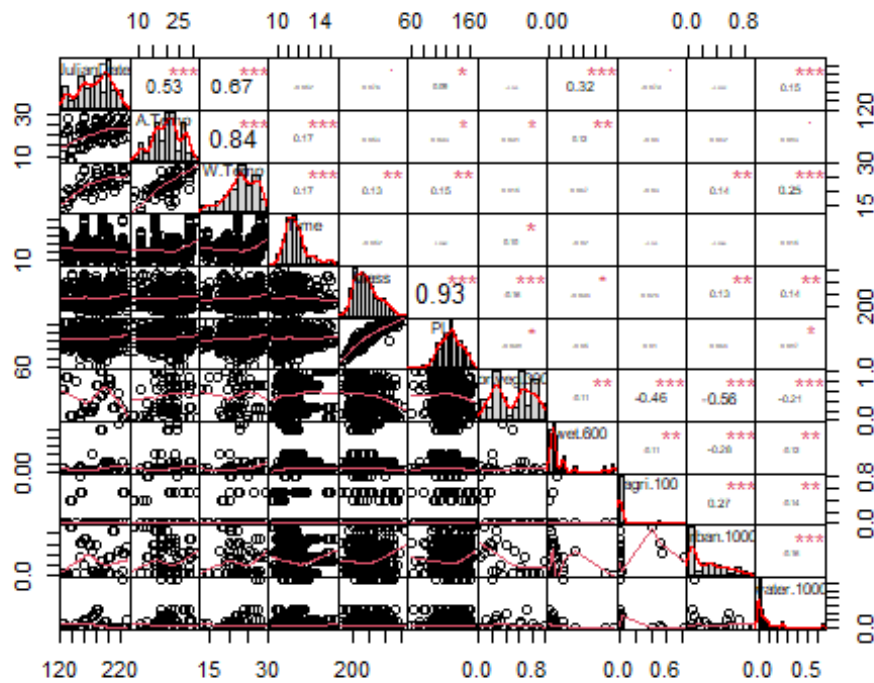
## Calculation of Pearson and Spearman correlation coefficients

### Pearson correlation coefficients

*# Visualization of the correlations*

```
cor.pearson.Move <- MixedData[, c(3,4,5,6,7,8,19,32,37,56,66)]
chart.Correlation(cor.pearson.Move, histogram=TRUE, pch=19)
```





*# Creation of the correlation table*

```
table.corr.pearson.Move <- rcorr(as.matrix(MixedData[,c(3,4,5,6,7,8,19,32,37,
56,66)]), type="pearson")
```

```
table.cor.pearson.Move.r <- table.corr.pearson.Move$r # Pearson correlation coefficients
```

```
table.cor.pearson.Move.p <- table.corr.pearson.Move$P # P values of the correlations
```

```
table.cor.pearson.Move.r
```

##	JulianDate	A.Temp	W.Temp	Time	Mass
## JulianDate	1.000000000	0.52662360	0.66868310	-0.051688682	0.07560208
## A.Temp	0.526623600	1.00000000	0.83887108	0.171967476	0.05252964
## W.Temp	0.668683098	0.83887108	1.00000000	0.166126059	0.13221505
## Time	-0.051688682	0.17196748	0.16612606	1.00000000	-0.05245171
## Mass	0.075602082	0.05252964	0.13221505	-0.052451709	1.00000000
## PL	0.089516177	0.09250283	0.14606475	0.009461487	0.93311455
## for.veg.300	-0.013496855	0.09108232	0.01541856	0.100593611	-0.15655831
## wet.600	0.317315461	0.12209698	0.06743185	-0.020258434	-0.09392134
## agri.100	-0.078714526	-0.03995312	-0.03022571	-0.013077109	0.02406704
## urban.1000	-0.003462429	0.04205111	0.14394243	-0.006102219	0.12738974
## water.1000	0.149646573	0.08311559	0.24535011	0.015131887	0.13576043
##	PL	for.veg.300	wet.600	agri.100	urban.1000
## JulianDate	0.089516177	-0.01349685	0.31731546	-0.07871453	-0.003462429
## A.Temp	0.092502829	0.09108232	0.12209698	-0.03995312	0.042051110
## W.Temp	0.146064749	0.01541856	0.06743185	-0.03022571	0.143942429
## Time	0.009461487	0.10059361	-0.02025843	-0.01307711	-0.006102219
## Mass	0.933114554	-0.15655831	-0.09392134	0.02406704	0.127389743

```
## PL      1.000000000 -0.09827262 -0.04979076  0.01037365  0.063945149
## for.veg.300 -0.098272621  1.000000000 -0.11488524 -0.46193445 -0.559879830
## wet.600    -0.049790756 -0.11488524  1.000000000 -0.11309548 -0.277413533
## agri.100   0.010373653 -0.46193445 -0.11309548  1.000000000  0.271994655
## urban.1000 0.063945149 -0.55987983 -0.27741353  0.27199465  1.000000000
## water.1000 0.086809165 -0.20733260 -0.11525621 -0.13673053 -0.163064570
##          water.1000
## JulianDate 0.14964657
## A.Temp     0.08311559
## W.Temp     0.24535011
## Time       0.01513189
## Mass       0.13576043
## PL         0.08680917
## for.veg.300 -0.20733260
## wet.600     -0.11525621
## agri.100    -0.13673053
## urban.1000  -0.16306457
## water.1000  1.000000000
```

table.cor.pearson.Move.p

```
##          JulianDate      A.Temp      W.Temp      Time      Ma
ss
## JulianDate      NA 0.000000e+00 0.000000e+00 2.322103e-01 0.08033738
37
## A.Temp          0.000000e+00      NA 0.000000e+00 6.283828e-05 0.22469108
77
## W.Temp          0.000000e+00 0.000000e+00      NA 2.346306e-04 0.00349965
95
## Time            2.322103e-01 6.283828e-05 2.346306e-04      NA 0.22538053
35
## Mass            8.033738e-02 2.246911e-01 3.499660e-03 2.253805e-01
NA
## PL              3.846835e-02 3.242105e-02 1.256393e-03 8.271659e-01 0.00000000
00
## for.veg.300     7.552233e-01 3.501640e-02 7.345696e-01 1.983870e-02 0.00027413
92
## wet.600         5.284662e-14 4.643928e-03 1.376963e-01 6.398062e-01 0.02969254
49
## agri.100        6.861526e-02 3.559077e-01 5.061950e-01 7.626040e-01 0.57822999
89
## urban.1000      9.362578e-01 3.311946e-01 1.464223e-03 8.879110e-01 0.00313231
18
## water.1000      5.088140e-04 5.446734e-02 4.280305e-08 7.266910e-01 0.00163080
42
##          PL    for.veg.300      wet.600      agri.100      urban.100
0
## JulianDate      0.038468345 7.552233e-01 5.284662e-14 6.861526e-02 9.362578e-0
1
## A.Temp          0.032421052 3.501640e-02 4.643928e-03 3.559077e-01 3.311946e-0
```

```

1
## W.Temp      0.001256393 7.345696e-01 1.376963e-01 5.061950e-01 1.464223e-0
3
## Time        0.827165890 1.983870e-02 6.398062e-01 7.626040e-01 8.879110e-0
1
## Mass        0.000000000 2.741392e-04 2.969254e-02 5.782300e-01 3.132312e-0
3
## PL           NA 2.301169e-02 2.502745e-01 8.108043e-01 1.396467e-0
1
## for.veg.300 0.023011686 NA 7.758640e-03 0.000000e+00 0.000000e+0
0
## wet.600     0.250274454 7.758640e-03 NA 8.776242e-03 6.305312e-1
1
## agri.100    0.810804345 0.000000e+00 8.776242e-03 NA 1.518086e-1
0
## urban.1000  0.139646656 0.000000e+00 6.305312e-11 1.518086e-10 NA
A
## water.1000  0.044750150 1.287993e-06 7.561412e-03 1.508389e-03 1.495518e-0
4
##             water.1000
## JulianDate  5.088140e-04
## A.Temp      5.446734e-02
## W.Temp      4.280305e-08
## Time        7.266910e-01
## Mass        1.630804e-03
## PL          4.475015e-02
## for.veg.300 1.287993e-06
## wet.600     7.561412e-03
## agri.100    1.508389e-03
## urban.1000  1.495518e-04
## water.1000  NA

```

### *Spearman correlation coefficients*

*# Creation of the correlation table*

```

table.corr.spearman.Move <- rcorr(as.matrix(MixedData[,c(3,4,5,6,7,8,19,32,37
,56,66)]), type="spearman")
table.cor.spearman.Move.r <- table.corr.spearman.Move$r # Pearson correlation
coefficients
table.cor.spearman.Move.p <- table.corr.spearman.Move$P # P values of the cor
relations
table.cor.spearman.Move.r

```

```

##             JulianDate      A.Temp      W.Temp      Time      Mass
## JulianDate  1.000000000 0.45842169 0.56884913 -0.09892087 0.08038929
## A.Temp      0.458421690 1.00000000 0.83718368 0.13849890 0.06589666
## W.Temp      0.568849127 0.83718368 1.00000000 0.17365903 0.13270468
## Time        -0.098920865 0.13849890 0.17365903 1.00000000 -0.04025948
## Mass        0.080389290 0.06589666 0.13270468 -0.04025948 1.00000000
## PL          0.088983783 0.10442546 0.15193967 -0.01364455 0.94856246
## for.veg.300 0.009875034 0.10037382 0.08247229 0.08653131 -0.11905658

```

```

## wet.600      0.408715978  0.18072693  0.24428729  0.16690425 -0.12345744
## agri.100     -0.089341274 -0.02430981 -0.06768594 -0.06305996  0.04983615
## urban.1000   0.057285275  0.13104537  0.21782810 -0.07145219  0.10868203
## water.1000  -0.228592159 -0.10346354 -0.01803073  0.03380113  0.05068902
##              PL    for.veg.300      wet.600      agri.100      urban.1000
## JulianDate   0.08898378  0.009875034  0.40871598 -0.08934127  0.05728527
## A.Temp       0.10442546  0.100373821  0.18072693 -0.02430981  0.13104537
## W.Temp       0.15193967  0.082472291  0.24428729 -0.06768594  0.21782810
## Time        -0.01364455  0.086531310  0.16690425 -0.06305996 -0.07145219
## Mass         0.94856246 -0.119056577 -0.12345744  0.04983615  0.10868203
## PL           1.00000000 -0.085385576 -0.11558586  0.05678992  0.07998962
## for.veg.300 -0.08538558  1.000000000  0.08511604 -0.49468619 -0.54813534
## wet.600      -0.11558586  0.085116043  1.00000000 -0.19656797 -0.28542464
## agri.100     0.05678992 -0.494686189 -0.19656797  1.00000000  0.27938984
## urban.1000   0.07998962 -0.548135336 -0.28542464  0.27938984  1.00000000
## water.1000   0.03595837 -0.233574219 -0.48557676 -0.07369328  0.06274805
##              water.1000
## JulianDate   -0.22859216
## A.Temp       -0.10346354
## W.Temp       -0.01803073
## Time         0.03380113
## Mass         0.05068902
## PL           0.03595837
## for.veg.300 -0.23357422
## wet.600      -0.48557676
## agri.100     -0.07369328
## urban.1000   0.06274805
## water.1000   1.00000000

```

table.cor.spearman.Move.p

```

##              JulianDate      A.Temp      W.Temp      Time      Mas
S
## JulianDate      NA 0.000000e+00 0.000000e+00 0.0219948308 0.06291206
1
## A.Temp          0.000000e+00      NA 0.000000e+00 0.0013067179 0.12758048
3
## W.Temp          0.000000e+00 0.000000e+00      NA 0.0001191832 0.00337907
1
## Time            2.199483e-02 1.306718e-03 1.191832e-04      NA 0.35222846
3
## Mass            6.291206e-02 1.275805e-01 3.379071e-03 0.3522284632      N
A
## PL              3.964107e-02 1.567793e-02 7.879312e-04 0.7528559899 0.00000000
0
## for.veg.300    8.195721e-01 2.011107e-02 6.928619e-02 0.0452390586 0.00578463
5
## wet.600        0.000000e+00 2.564232e-05 4.913921e-08 0.0001034490 0.00420271
2
## agri.100       3.866754e-02 5.744009e-01 1.362161e-01 0.1448442211 0.24939811

```

```

4
## urban.1000 1.854229e-01 2.366128e-03 1.246624e-06 0.0984350178 0.01181043
0
## water.1000 8.749580e-08 1.656624e-02 6.917336e-01 0.4348325411 0.24137815
2
##          PL  for.veg.300      wet.600      agri.100      urban.10
00
## JulianDate 0.0396410665 8.195721e-01 0.000000e+00 3.866754e-02 1.854229e-
01
## A.Temp      0.0156779333 2.011107e-02 2.564232e-05 5.744009e-01 2.366128e-
03
## W.Temp      0.0007879312 6.928619e-02 4.913921e-08 1.362161e-01 1.246624e-
06
## Time        0.7528559899 4.523906e-02 1.034490e-04 1.448442e-01 9.843502e-
02
## Mass        0.0000000000 5.784635e-03 4.202712e-03 2.493981e-01 1.181043e-
02
## PL          NA 4.838620e-02 7.445541e-03 1.896750e-01 6.448653e-
02
## for.veg.300 0.0483861978          NA 4.888954e-02 0.000000e+00 0.000000e+
00
## wet.600     0.0074455412 4.888954e-02          NA 4.538554e-06 1.659783e-
11
## agri.100    0.1896750465 0.000000e+00 4.538554e-06          NA 4.554446e-
11
## urban.1000 0.0644865292 0.000000e+00 1.659783e-11 4.554446e-11
NA
## water.1000 0.4065146753 4.478494e-08 0.000000e+00 8.829560e-02 1.468465e-
01
##          water.1000
## JulianDate 8.749580e-08
## A.Temp      1.656624e-02
## W.Temp      6.917336e-01
## Time        4.348325e-01
## Mass        2.413782e-01
## PL          4.065147e-01
## for.veg.300 4.478494e-08
## wet.600     0.000000e+00
## agri.100    8.829560e-02
## urban.1000 1.468465e-01
## water.1000          NA

```

The Pearson correlation coefficient of 0.93 between Mass and PL and 0.84 between W.Temp and A.Temp confirmed the deletion of Mass and W.Temp with the calculation of the  $GVIF^{(1/(2*df))}$ .

#### Verification of the assumptions with the initial model

```

mod.Move.full <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled
+ PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + u
rban.1000.Scaled + water.1000.Scaled + (1|Code) + (1|Site), data = MixedData,

```

```
REML = TRUE, na.action=na.exclude)
```

```
summary(mod.Move.full)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled +
```

```
## Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled +
## urban.1000.Scaled + water.1000.Scaled + (1 | Code) + (1 | Site)
## Data: MixedData
```

```
##
## REML criterion at convergence: 4604.6
##
```

```
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.94964 -0.52458  0.06559  0.55080  1.67014
##
```

```
## Random effects:
## Groups Name Variance Std.Dev.
## Code (Intercept) 756.5 27.50
## Site (Intercept) 231.9 15.23
## Residual 894.8 29.91
## Number of obs: 453, groups: Code, 429; Site, 23
##
```

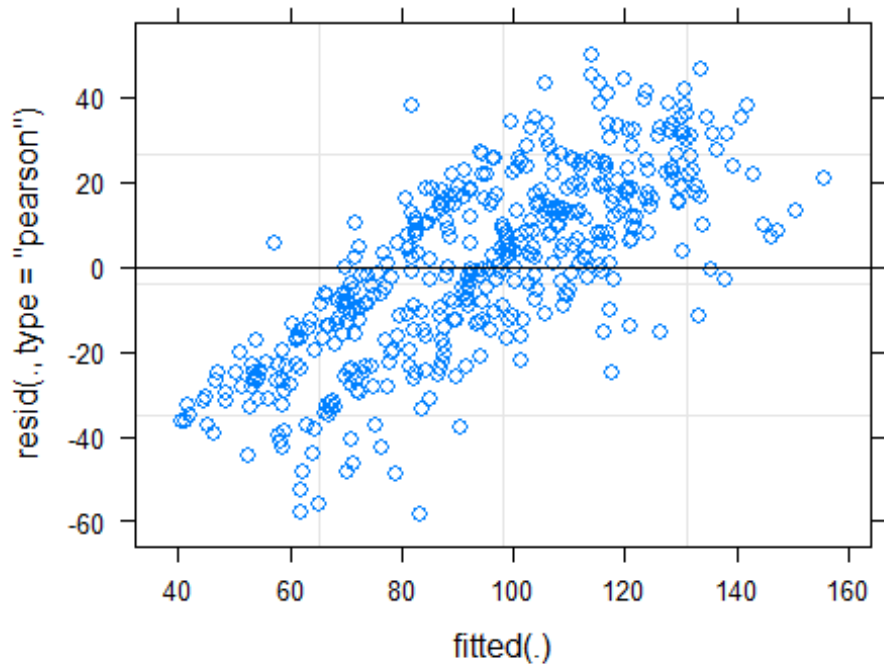
```
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    89.7725    4.7506  29.2985  18.897 <2e-16 ***
## JulianDate.Scaled  2.8778    4.5793  19.6989   0.628  0.5369
## A.Temp.Scaled   -1.6130    3.4232 105.3638  -0.471  0.6385
## Time.Scaled     1.2102    2.1178 432.5464   0.571  0.5680
## PL.Scaled       -0.6353    2.5282 429.7145  -0.251  0.8017
## SexM            4.9045    4.5844 422.3354   1.070  0.2853
## for.veg.300.Scaled 0.3362    6.6030  15.7704   0.051  0.9600
## wet.600.Scaled   -0.8010    4.8360  16.8054  -0.166  0.8704
## agri.100.Scaled   7.6336    4.2909  19.6989   1.779  0.0907 .
## urban.1000.Scaled 5.9955    5.8880  16.6737   1.018  0.3231
## water.1000.Scaled -3.2380    4.6730  15.5003  -0.693  0.4986
```

```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S Tm.Sc1 PL.Sc1 SexM   f..300 w.600. a.100.
## JulnDt.Sc1d -0.056
## A.Temp.Sc1d  0.090 -0.404
## Time.Sc1d   0.045  0.101 -0.227
## PL.Sc1d     -0.280 -0.022 -0.050 -0.038
## SexM        -0.557  0.050 -0.108 -0.053  0.467
## fr.vg.300.S 0.110 -0.022 -0.127 -0.024  0.039  0.011
## wt.600.Sc1d -0.005 -0.329  0.005 -0.017  0.064  0.012  0.498
```

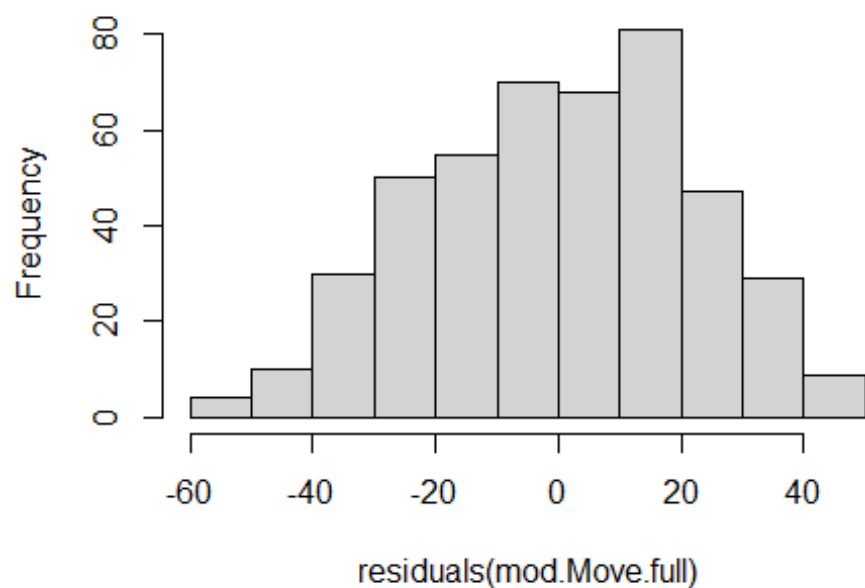
```
## agr.100.Scl 0.032 0.008 -0.057 0.011 0.012 -0.034 0.592 0.343
## urbn.1000.S 0.003 -0.194 -0.007 -0.004 0.034 0.043 0.686 0.591 0.308
## wtr.1000.Sc 0.010 -0.260 -0.023 -0.029 -0.028 -0.015 0.553 0.489 0.413
##
## u.1000
## JulnDt.Scld
## A.Temp.Scld
## Time.Scaled
## PL.Scaled
## SexM
## fr.vg.300.S
## wt.600.Scld
## agr.100.Scl
## urbn.1000.S
## wtr.1000.Sc 0.558

plot(mod.Move.full)
```



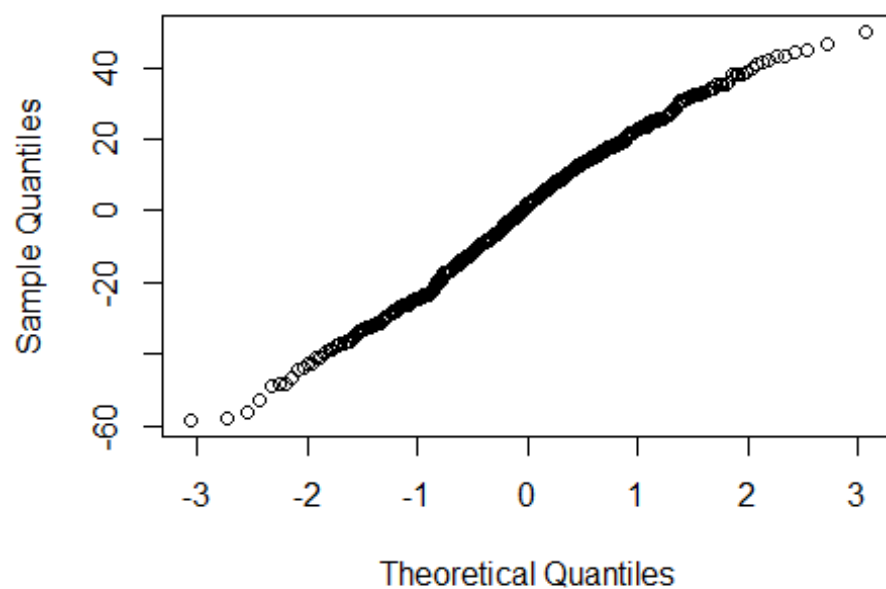
```
hist(residuals(mod.Move.full))
```

**Histogram of residuals(mod.Move.full)**



```
qqnorm(resid(mod.Move.full))
```

**Normal Q-Q Plot**





## Model selection

### Active defensive behaviours

#### Random variable

I am testing the significance of turtle ID and site identity by using likelihood ratio tests to see if the addition of these random variables has a significant effect on the initial model. I am using a dummy variable (the same value for all the observations) to create a null mixed model to compared with the different combinations of mixed models.

#### Creation of the different mixed models

##### **## Null mixed model**

```
mod.Aggression.null <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scaled +  
Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000  
.Scaled + water.400.Scaled + (1|Dummy), data = MixedData, na.action=na.exclud  
e, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

##### **## only site identity as random variable**

```
mod.Aggression.dummy.site <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Sca  
led + Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urba  
n.1000.Scaled + water.400.Scaled + (1|Dummy) + (1|Site), data = MixedData, na  
.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FA  
LSE)
```

##### **## only turtle ID as random variable**

```
mod.Aggression.dummy.code <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Sca  
led + Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urba  
n.1000.Scaled + water.400.Scaled + (1|Dummy) + (1|Code), data = MixedData, na  
.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FA  
LSE)
```

## Likelihood ratio tests between the mixed models

#### **#anova with null model and dummy + site model**

```
anova(mod.Aggression.null, mod.Aggression.dummy.site)
```

```
## Data: MixedData
```

```
## Models:
```

```
## mod.Aggression.null: Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time  
.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000.Sca  
led + water.400.Scaled + (1 | Dummy)
```

```
## mod.Aggression.dummy.site: Aggression ~ JulianDate.Scaled + A.Temp.Scaled  
+ Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.10  
00.Scaled + water.400.Scaled + (1 | Dummy) + (1 | Site)
```

```
##
```

	npar	AIC	BIC	logLik	deviance	Chisq	Df
--	------	-----	-----	--------	----------	-------	----

## mod.Aggression.null	12	1519.2	1570.3	-747.61	1495.2		
------------------------	----	--------	--------	---------	--------	--	--

## mod.Aggression.dummy.site	13	1521.2	1576.5	-747.59	1495.2	0.0522	1
------------------------------	----	--------	--------	---------	--------	--------	---

```
## Pr(>Chisq)
```

```

## mod.Aggression.null
## mod.Aggression.dummy.site      0.8192

#anova with null model and dummy + code model
anova(mod.Aggression.null, mod.Aggression.dummy.code)

## Data: MixedData
## Models:
## mod.Aggression.null: Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time
.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000.Sca
led + water.400.Scaled + (1 | Dummy)
## mod.Aggression.dummy.code: Aggression ~ JulianDate.Scaled + A.Temp.Scaled
+ Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.10
00.Scaled + water.400.Scaled + (1 | Dummy) + (1 | Code)
##
##      npar      AIC      BIC    logLik deviance  Chisq Df
## mod.Aggression.null      12 1519.2 1570.3 -747.61    1495.2
## mod.Aggression.dummy.code  13 1486.8 1542.2 -730.41    1460.8 34.397  1
##
##      Pr(>Chisq)
## mod.Aggression.null
## mod.Aggression.dummy.code  4.494e-09 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

If a test has a significant p-value (less than 0.05) then the random effect is significant. Turtle ID (Code) is significant by itself but site identity (Site) is not. I will see if Code and Site together are more significant than Code by itself.

```

## turtle ID without the dummy variable
mod.Aggression.code <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scaled +
Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000
.Scaled + water.400.Scaled + (1|Code), data = MixedData, na.action=na.exclude
, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)

## turtle ID and site identity without the dummy variable
mod.Aggression.code.site <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scal
ed + Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban
.1000.Scaled + water.400.Scaled + (1|Code) + (1|Site), data = MixedData, na.a
ction=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALS
E)

#anova with code model and code + site model
anova(mod.Aggression.code, mod.Aggression.code.site)

## Data: MixedData
## Models:
## mod.Aggression.code: Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time
.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000.Sca
led + water.400.Scaled + (1 | Code)
## mod.Aggression.code.site: Aggression ~ JulianDate.Scaled + A.Temp.Scaled +
Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000
.Scaled + water.400.Scaled + (1 | Code) + (1 | Site)

```

```
##               npar    AIC    BIC  logLik deviance Chisq Df
## mod.Aggression.code      12 1484.8 1535.9 -730.41   1460.8
## mod.Aggression.code.site  13 1486.8 1542.2 -730.41   1460.8 8e-04  1
##               Pr(>Chisq)
## mod.Aggression.code
## mod.Aggression.code.site    0.9772
```

Code and Site together are not more significant than Code by itself ( $p > 0.05$ ), so I will only keep Code.

## Predictor variables

I am selecting the final model with a backward selection procedure. At each step, I deleted the fixed effect with the highest p value. I confirmed the deletion of each fixed effect with a likelihood ratio test. I created a new dataset at each step to use only the rows with complete observations for all the fixed effects, so that the likelihood ratio tests do not run between two models with a different number of observations.

```
mod.Aggression.full <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scaled +
Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000
.Scaled + water.400.Scaled + (1|Code), data = MixedData, na.action=na.exclude
, REML = FALSE)
summary(mod.Aggression.full)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
##      PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled
##      +
##      water.400.Scaled + (1 | Code)
##      Data: MixedData
##
##      AIC      BIC    logLik deviance df.resid
##    1484.8    1535.9   -730.4   1460.8      510
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.60879 -0.32199 -0.05001  0.31016  1.57965
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
##  Code     (Intercept)  0.8978     0.9475
##  Residual                  0.1433     0.3785
## Number of obs: 522, groups:  Code, 492
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    0.99134    0.07176 416.06835  13.815 < 2e-16 ***
## JulianDate.Scaled -0.06446    0.05923 513.61544  -1.088  0.27698
```

```
## A.Temp.Scaled      0.14055      0.05411 490.26081    2.597  0.00968 **
## Time.Scaled        -0.03584      0.04230 328.28211   -0.847  0.39746
## PL.Scaled          -0.03876      0.05547 492.74772   -0.699  0.48508
## SexM               0.05669      0.09405 251.64085    0.603  0.54720
## wet.200.Scaled     -0.04635      0.05135 495.02947   -0.903  0.36713
## agri.100.Scaled    -0.06135      0.04946 493.06408   -1.240  0.21538
## urban.1000.Scaled  0.09016      0.04996 492.73077    1.805  0.07172 .
## water.400.Scaled   0.13219      0.04929 492.59868    2.682  0.00757 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) JlnD.S A.Tm.S Tm.Scl PL.Scl SexM   w.200. a.100. u.1000
## JulnDt.Scl -0.073
## A.Temp.Scl  0.069 -0.523
## Time.Scaled 0.030  0.162 -0.242
## PL.Scaled   -0.378 -0.051 -0.068 -0.014
## SexM        -0.767  0.068 -0.059 -0.051  0.461
## wt.200.Scl -0.042 -0.312  0.109  0.011  0.097  0.025
## agr.100.Scl 0.084  0.090 -0.015  0.033 -0.088 -0.104  0.013
## urbn.1000.S -0.074 -0.045 -0.023  0.017  0.015  0.077  0.319 -0.223
## wtr.400.Scl 0.048  0.154 -0.074  0.005 -0.138 -0.074  0.173  0.198  0.117
```

I deleted turtle sex (Sex).

```
mod.Aggression.1 <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time
e.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled +
water.400.Scaled + (1|Code), data = MixedData, na.action=na.exclude, REML = F
ALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Aggression, MixedData$
JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$
PL.Scaled, MixedData$Sex, MixedData$wet.200.Scaled, MixedData$agri.100.Scaled
, MixedData$urban.1000.Scaled, MixedData$water.400.Scaled),]
```

```
mod.full.adjust <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time
.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000.Sca
led + water.400.Scaled + (1|Code), data = MixedData.adjust, na.action=na.excl
ude, REML = FALSE)
```

```
mod.Aggression.1.adjust <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scale
d + Time.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.S
caled + water.400.Scaled + (1|Code), data = MixedData.adjust, na.action=na.ex
clude, REML = FALSE)
```

```
anova(mod.Aggression.1.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Aggression.1.adjust: Aggression ~ JulianDate.Scaled + A.Temp.Scaled +
```

```

Time.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled + (1 | Code)
## mod.full.adjust: Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled + (1 | Code)
##
##          npar      AIC      BIC  logLik deviance  Chisq Df
## mod.Aggression.1.adjust    11 1483.2 1530.0 -730.59   1461.2
## mod.full.adjust           12 1484.8 1535.9 -730.41   1460.8 0.3555  1
##
##          Pr(>Chisq)
## mod.Aggression.1.adjust
## mod.full.adjust           0.551

summary(mod.Aggression.1)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
##          PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled +
##          water.400.Scaled + (1 | Code)
## Data: MixedData
##
##          AIC      BIC  logLik deviance df.resid
## 1528.6 1575.7 -753.3 1506.6      524
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.93844 -0.39021 -0.05932  0.37266  2.46919
##
## Random effects:
## Groups   Name      Variance Std.Dev.
## Code     (Intercept) 0.8257   0.9087
## Residual              0.2124   0.4608
## Number of obs: 535, groups: Code, 504
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    1.02630    0.04530 500.68223  22.653 < 2e-16 ***
## JulianDate.Scaled -0.05129    0.05874 518.60051  -0.873  0.38300
## A.Temp.Scaled    0.13273    0.05427 531.82399   2.446  0.01478 *
## Time.Scaled     -0.01832    0.04320 427.49445  -0.424  0.67171
## PL.Scaled       -0.06187    0.04531 530.22999  -1.365  0.17271
## wet.200.Scaled  -0.05092    0.05084 508.59182  -1.002  0.31700
## agri.100.Scaled -0.04561    0.04831 501.97540  -0.944  0.34559
## urban.1000.Scaled 0.08383    0.04924 504.46854   1.702  0.08929 .
## water.400.Scaled 0.14028    0.04786 503.15370   2.931  0.00353 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S Tm.Scl PL.Scl w.200. a.100. u.1000

```

```
## JulnDt.Scl d -0.033
## A.Temp.Scl d 0.021 -0.529
## Time.Scaled 0.001 0.168 -0.235
## PL.Scaled 0.021 -0.096 -0.012 -0.022
## wt.200.Scl d -0.021 -0.317 0.123 0.002 0.065
## agr.100.Scl -0.002 0.101 -0.025 0.026 -0.017 0.016
## urbn.1000.S -0.021 -0.042 -0.019 0.012 -0.035 0.313 -0.225
## wtr.400.Scl -0.031 0.154 -0.091 0.010 -0.089 0.181 0.187 0.133
```

I deleted time of testing (Time).

```
mod.Aggression.2 <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scaled + PL.
Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Sca
led + (1|Code), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Aggression, MixedData$
JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$
PL.Scaled, MixedData$wet.200.Scaled, MixedData$agri.100.Scaled, MixedData$urb
an.1000.Scaled, MixedData$water.400.Scaled),]
```

```
mod.full.adjust <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time
.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled +
water.400.Scaled + (1|Code), data = MixedData.adjust, na.action=na.exclude, R
EML = FALSE)
```

```
mod.Aggression.2.adjust <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scale
d + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.
400.Scaled + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML =
FALSE)
```

```
anova(mod.Aggression.2.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Aggression.2.adjust: Aggression ~ JulianDate.Scaled + A.Temp.Scaled +
PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.
Scaled + (1 | Code)
```

```
## mod.full.adjust: Aggression ~ JulianDate.Scaled + A.Temp.Scaled + Time.Sca
led + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + wate
r.400.Scaled + (1 | Code)
```

```
##          npar      AIC      BIC logLik deviance Chisq Df
## mod.Aggression.2.adjust    10 1526.8 1569.6 -753.38   1506.8
## mod.full.adjust           11 1528.6 1575.7 -753.29   1506.6 0.1798  1
##          Pr(>Chisq)
```

```
## mod.Aggression.2.adjust
```

```
## mod.full.adjust          0.6715
```

```
summary(mod.Aggression.2)
```

```
## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
```

```

## Formula: Aggression ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled +
##      wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled +
##      (1 | Code)
##      Data: MixedData
##
##      AIC      BIC    logLik deviance df.resid
##    1526.8    1569.6   -753.4   1506.8     525
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.90627 -0.39311 -0.06746  0.37682  2.42856
##
## Random effects:
##   Groups   Name                Variance Std.Dev.
##   Code     (Intercept)  0.8259     0.9088
##   Residual                  0.2125     0.4610
## Number of obs: 535, groups: Code, 504
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    1.02632    0.04531 500.63443   22.650 < 2e-16 ***
## JulianDate.Scaled -0.04711    0.05792 519.45485    -0.813  0.41638
## A.Temp.Scaled    0.12731    0.05276 529.72570     2.413  0.01616 *
## PL.Scaled        -0.06229    0.04531 530.44429    -1.375  0.16976
## wet.200.Scaled   -0.05088    0.05085 508.54422    -1.001  0.31744
## agri.100.Scaled  -0.04508    0.04830 501.83505    -0.933  0.35111
## urban.1000.Scaled 0.08408    0.04924 504.42215     1.707  0.08837 .
## water.400.Scaled 0.14048    0.04786 503.19256     2.935  0.00349 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S PL.Scl w.200. a.100. u.1000
## JulnDt.Scld -0.033
## A.Temp.Scld  0.022 -0.511
## PL.Scaled    0.021 -0.094 -0.017
## wt.200.Scld -0.021 -0.321  0.127  0.065
## agr.100.Scl -0.002  0.098 -0.020 -0.016  0.016
## urbn.1000.S -0.021 -0.045 -0.016 -0.035  0.313 -0.226
## wtr.400.Scl -0.031  0.154 -0.092 -0.089  0.181  0.187  0.133

```

I deleted Julian date of testing (JulianDate).

```

mod.Aggression.3 <- lmer(Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled +
agri.100.Scaled + urban.1000.Scaled + water.400.Scaled + (1|Code), data
= MixedData, na.action=na.exclude, REML = FALSE)

```

```

MixedData.adjust <- MixedData[complete.cases(MixedData$Aggression, MixedData$
JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$PL.Scaled, MixedData$we

```

```
t.200.Scaled, MixedData$agri.100.Scaled, MixedData$urban.1000.Scaled, MixedData$water.400.Scaled),]
```

```
mod.full.adjust <- lmer(Aggression ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.Aggression.3.adjust <- lmer(Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.Aggression.3.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Aggression.3.adjust: Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled + (1 | Code)
```

```
## mod.full.adjust: Aggression ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled + (1 | Code)
```

```
##
```

	npars	AIC	BIC	logLik	deviance	Chisq	Df
## mod.Aggression.3.adjust	9	1525.4	1564.0	-753.71	1507.4		
## mod.full.adjust	10	1526.8	1569.6	-753.38	1506.8	0.6609	1

```
##
```

```
## Pr(>Chisq)
```

```
## mod.Aggression.3.adjust
```

```
## mod.full.adjust
```

```
summary(mod.Aggression.3)
```

```
## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
```

```
## method [lmerModLmerTest]
```

```
## Formula:
```

```
## Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled +
```

```
## urban.1000.Scaled + water.400.Scaled + (1 | Code)
```

```
## Data: MixedData
```

```
##
```

```
##
```

	AIC	BIC	logLik	deviance	df.resid
##	1525.4	1564.0	-753.7	1507.4	526

```
##
```

```
## Scaled residuals:
```

```
##
```

	Min	1Q	Median	3Q	Max
##	-1.89243	-0.39528	-0.07591	0.38030	2.43447

```
##
```

```
##
```

```
## Random effects:
```

```
##
```

	Groups	Name	Variance	Std.Dev.
##	Code	(Intercept)	0.8258	0.9088
##	Residual		0.2137	0.4622

```
##
```

```
##
```

```
## Number of obs: 535, groups: Code, 504
```

```
##
```



```
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    1.02509    0.04531 500.81975   22.625 < 2e-16 ***
## A.Temp.Scaled    0.10532    0.04536 534.99437    2.322  0.02063 *
## PL.Scaled       -0.06569    0.04513 530.40418   -1.456  0.14611
## wet.200.Scaled  -0.06418    0.04817 504.98435   -1.332  0.18340
## agri.100.Scaled -0.04122    0.04809 501.88280   -0.857  0.39178
## urban.1000.Scaled 0.08227    0.04922 503.91281    1.672  0.09522 .
## water.400.Scaled 0.14648    0.04731 503.27379    3.096  0.00207 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) A.Tm.S PL.Scl w.200. a.100. u.1000
## A.Temp.Scl d  0.006
## PL.Scaled     0.018 -0.076
## wt.200.Scl d -0.033 -0.046  0.037
## agr.100.Scl  0.001  0.036 -0.007  0.050
## urbn.1000.S -0.023 -0.046 -0.039  0.316 -0.223
## wtr.400.Scl -0.026 -0.015 -0.076  0.246  0.175  0.142
```

I deleted proportion of agricultural area at 100m (agri.100).

```
mod.Aggression.4 <- lmer(Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled + urban.1000.Scaled + water.400.Scaled + (1|Code), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Aggression, MixedData$A.Temp.Scaled, MixedData$PL.Scaled, MixedData$wet.200.Scaled, MixedData$agri.100.Scaled, MixedData$urban.1000.Scaled, MixedData$water.400.Scaled),]
```

```
mod.full.adjust <- lmer(Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.Aggression.4.adjust <- lmer(Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled + urban.1000.Scaled + water.400.Scaled + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.Aggression.4.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Aggression.4.adjust: Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled + urban.1000.Scaled + water.400.Scaled + (1 | Code)
```

```
## mod.full.adjust: Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled + agri.100.Scaled + urban.1000.Scaled + water.400.Scaled + (1 | Code)
```

```
##              npar      AIC      BIC  logLik deviance Chisq Df Pr(>ChiSq)
```

```
## mod.Aggression.4.adjust      8 1524.2 1558.4 -754.07   1508.2
```

```

## mod.full.adjust          9 1525.4 1564.0 -753.71   1507.4 0.734 1    0
.3916

summary(mod.Aggression.4)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula:
## Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled + urban.1000.Scaled +
## water.400.Scaled + (1 | Code)
## Data: MixedData
##
##      AIC      BIC    logLik deviance df.resid
##  1524.1   1558.4   -754.1   1508.1     527
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.89448 -0.38885 -0.07051  0.38111  2.43647
##
## Random effects:
## Groups Name Variance Std.Dev.
## Code (Intercept) 0.8279  0.9099
## Residual 0.2133  0.4618
## Number of obs: 535, groups: Code, 504
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    1.02512    0.04534 500.93942   22.608 < 2e-16 ***
## A.Temp.Scaled    0.10675    0.04537 534.99140    2.353  0.01898 *
## PL.Scaled      -0.06600    0.04517 530.47415   -1.461  0.14453
## wet.200.Scaled -0.06210    0.04815 505.23814   -1.290  0.19776
## urban.1000.Scaled 0.07288    0.04802 503.27997    1.518  0.12973
## water.400.Scaled 0.15356    0.04662 503.63761    3.294  0.00106 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) A.Tm.S PL.Scl w.200. u.1000
## A.Temp.Scld  0.006
## PL.Scaled    0.018 -0.076
## wt.200.Scld -0.033 -0.048  0.038
## urbn.1000.S -0.023 -0.039 -0.042  0.336
## wtr.400.Scl -0.027 -0.022 -0.076  0.241  0.189

```

I deleted proportion of wetland area at 200m (wet.200).

```

mod.Aggression.5 <- lmer(Aggression ~ A.Temp.Scaled + PL.Scaled + urban.1000.
Scaled + water.400.Scaled + (1|Code), data = MixedData, na.action=na.exclude,
REML = FALSE)

```

```

MixedData.adjust <- MixedData[complete.cases(MixedData$Aggression, MixedData$
A.Temp.Scaled, MixedData$PL.Scaled, MixedData$wet.200.Scaled, MixedData$urban
.1000.Scaled, MixedData$water.400.Scaled),]

mod.full.adjust <- lmer(Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scal
ed + urban.1000.Scaled + water.400.Scaled + (1|Code), data = MixedData.adjust
, na.action=na.exclude, REML = FALSE)

mod.Agression.5.adjust <- lmer(Aggression ~ A.Temp.Scaled + PL.Scaled + urba
n.1000.Scaled + water.400.Scaled + (1|Code), data = MixedData.adjust, na.acti
on=na.exclude, REML = FALSE)

anova(mod.Agression.5.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.Agression.5.adjust: Aggression ~ A.Temp.Scaled + PL.Scaled + urban.10
00.Scaled + water.400.Scaled + (1 | Code)
## mod.full.adjust: Aggression ~ A.Temp.Scaled + PL.Scaled + wet.200.Scaled +
urban.1000.Scaled + water.400.Scaled + (1 | Code)
##
##          npar      AIC      BIC logLik deviance Chisq Df
## mod.Agression.5.adjust      7 1523.8 1553.8 -754.90    1509.8
## mod.full.adjust            8 1524.2 1558.4 -754.07    1508.2 1.6606  1
##
##          Pr(>Chisq)
## mod.Agression.5.adjust
## mod.full.adjust          0.1975

summary(mod.Agression.5)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Aggression ~ A.Temp.Scaled + PL.Scaled + urban.1000.Scaled +
##          water.400.Scaled + (1 | Code)
## Data: MixedData
##
##          AIC      BIC logLik deviance df.resid
## 1523.8    1553.8  -754.9   1509.8      528
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.88555 -0.38883 -0.06362  0.39251  2.44042
##
## Random effects:
## Groups   Name                Variance Std.Dev.
## Code     (Intercept) 0.8306    0.9114
## Residual                0.2137    0.4623
## Number of obs: 535, groups: Code, 504
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)

```

```
## (Intercept)          1.02318    0.04539 501.20871  22.542 < 2e-16 ***
## A.Temp.Scaled        0.10395    0.04538 534.99225   2.290 0.022383 *
## PL.Scaled            -0.06381    0.04520 530.40562  -1.412 0.158673
## urban.1000.Scaled    0.09366    0.04530 504.79137   2.067 0.039204 *
## water.400.Scaled     0.16806    0.04532 504.82354   3.708 0.000232 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) A.Tm.S PL.Scl u.1000
## A.Temp.Scl d  0.004
## PL.Scaled     0.019 -0.074
## urbn.1000.S  -0.013 -0.024 -0.058
## wtr.400.Scl  -0.019 -0.011 -0.088  0.118
```

I deleted plastron length (PL).

```
mod.Aggression.6 <- lmer(Aggression ~ A.Temp.Scaled + urban.1000.Scaled + wat
er.400.Scaled + (1|Code), data = MixedData, na.action=na.exclude, REML = FALS
E)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Aggression, MixedData$
A.Temp.Scaled, MixedData$PL.Scaled, MixedData$urban.1000.Scaled, MixedData$wa
ter.400.Scaled),]
```

```
mod.full.adjust <- lmer(Aggression ~ A.Temp.Scaled + PL.Scaled + urban.1000.S
caled + water.400.Scaled + (1|Code), data = MixedData.adjust, na.action=na.ex
clude, REML = FALSE)
```

```
mod.Aggression.6.adjust <- lmer(Aggression ~ A.Temp.Scaled + urban.1000.Scale
d + water.400.Scaled + (1|Code), data = MixedData.adjust, na.action=na.exclud
e, REML = FALSE)
```

```
anova(mod.Aggression.6.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Aggression.6.adjust: Aggression ~ A.Temp.Scaled + urban.1000.Scaled +
water.400.Scaled + (1 | Code)
```

```
## mod.full.adjust: Aggression ~ A.Temp.Scaled + PL.Scaled + urban.1000.Scale
d + water.400.Scaled + (1 | Code)
```

```
##              npar      AIC      BIC logLik deviance Chisq Df Pr(>Chi
sq)
```

```
## mod.Aggression.6.adjust      6 1523.8 1549.5 -755.9   1511.8
```

```
## mod.full.adjust              7 1523.8 1553.8 -754.9   1509.8 1.983  1    0.
1591
```

```
summary(mod.Aggression.6)
```

```
## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
```

```

## Formula: Aggression ~ A.Temp.Scaled + urban.1000.Scaled + water.400.Scaled
+
##      (1 | Code)
##      Data: MixedData
##
##      AIC      BIC    logLik deviance df.resid
##  1525.7    1551.4   -756.8   1513.7     530
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.84490 -0.38534 -0.05837  0.39662  2.43163
##
## Random effects:
##  Groups   Name            Variance Std.Dev.
##  Code     (Intercept) 0.8264    0.9091
##  Residual                0.2187    0.4677
## Number of obs: 536, groups: Code, 505
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    1.02412    0.04535 502.37553   22.583 < 2e-16 ***
## A.Temp.Scaled    0.09838    0.04513 535.87535    2.180 0.029684 *
## urban.1000.Scaled 0.08987    0.04524 505.62984    1.987 0.047510 *
## water.400.Scaled 0.16271    0.04514 506.00437    3.605 0.000344 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) A.Tm.S u.1000
## A.Temp.Scld  0.002
## urbn.1000.S -0.012 -0.030
## wtr.400.Scl -0.016 -0.015  0.114

```

All of the fixed effects are statistically significant, so I will stop the backwards selection process.

## Final model

### Summary statistics

I changed the REML to TRUE to calculate the summary statistics of the final model.

```

mod.Aggression.final <- lmer(Aggression ~ A.Temp.Scaled + urban.1000.Scaled +
water.400.Scaled + (1|Code), data = MixedData, na.action=na.exclude, REML = T
RUE)
summary(mod.Aggression.final)

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]

```

```

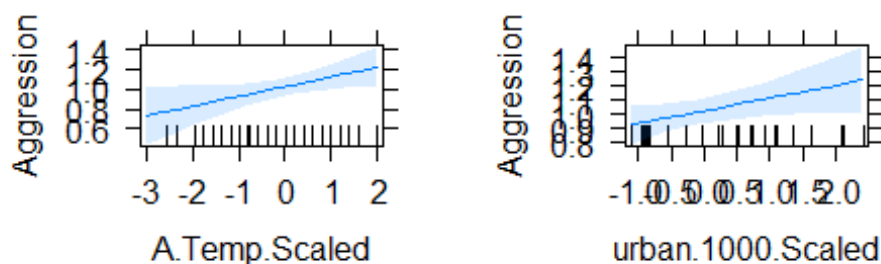
## Formula: Aggression ~ A.Temp.Scaled + urban.1000.Scaled + water.400.Scaled
+
##      (1 | Code)
##      Data: MixedData
##
## REML criterion at convergence: 1531.1
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.84545 -0.38266 -0.05814  0.39317  2.42771
##
## Random effects:
##   Groups   Name                Variance Std.Dev.
##   Code     (Intercept) 0.8343     0.9134
##   Residual                0.2191     0.4680
## Number of obs: 536, groups: Code, 505
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    1.02412    0.04553 498.54384  22.495 < 2e-16 ***
## A.Temp.Scaled    0.09844    0.04530 531.88334   2.173 0.030203 *
## urban.1000.Scaled 0.08987    0.04542 501.74100   1.979 0.048389 *
## water.400.Scaled 0.16269    0.04532 502.10846   3.590 0.000363 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) A.Tm.S u.1000
## A.Temp.Scld  0.002
## urbn.1000.S -0.012 -0.030
## wtr.400.Scl -0.016 -0.015  0.114

```

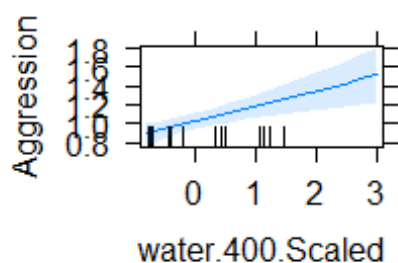
### Visualization of the predictor effects

```
plot(allEffects(mod.Aggression.final))
```

## A.Temp.Scaled effect ploturban.1000.Scaled effect plot



## water.400.Scaled effect plot



## Calculation of the marginal and conditional variance explained by the final model

```
r.squaredGLMM(mod.Aggression.final)
```

```
##           R2m           R2c
## [1,] 0.03730533 0.7997945
```

Marginal R2: fixed effects R2. Conditional R2: fixed and random effects R2.

## Calculation of the 95% confidence intervals

```
confint(mod.Aggression.final, level = 0.95, method = "Wald")
```

```
##                2.5 %    97.5 %
## .sig01             NA         NA
## .sigma             NA         NA
## (Intercept)    0.9348892047 1.1133529
## A.Temp.Scaled    0.0096607255 0.1872179
## urban.1000.Scaled 0.0008539708 0.1788790
## water.400.Scaled 0.0738757498 0.2515142
```

## Creation of the prediction figure for urban area at 1000m

```
pred.con.model.Aggression.urban.1000 <- ggpredict(mod.Aggression.final, terms
= "urban.1000.Scaled")
pred.con.model.Aggression.urban.1000
```

```
## # Predicted values of Aggression
##
## urban.1000.Scaled | Predicted |          95% CI
## -----|-----|-----
##          -1.20 |          0.92 | [0.78, 1.06]
##          -0.80 |          0.95 | [0.84, 1.07]
##          -0.20 |          1.01 | [0.91, 1.10]
##           0.20 |          1.04 | [0.95, 1.13]
##           0.60 |          1.08 | [0.97, 1.18]
##           1.00 |          1.11 | [0.99, 1.24]
##           1.40 |          1.15 | [1.00, 1.30]
##           2.40 |          1.24 | [1.01, 1.47]
##
## Adjusted for:
## *      A.Temp.Scaled = 0.00
## * water.400.Scaled = 0.00
## *              Code = 0 (population-level)
# New dataset to only have complete observations for all the variables
MixedData.adjust.final.model.Aggression <- MixedData[complete.cases(MixedData
$A.Temp.Scaled, MixedData$urban.1000.Scaled, MixedData$water.400.Scaled),]
```

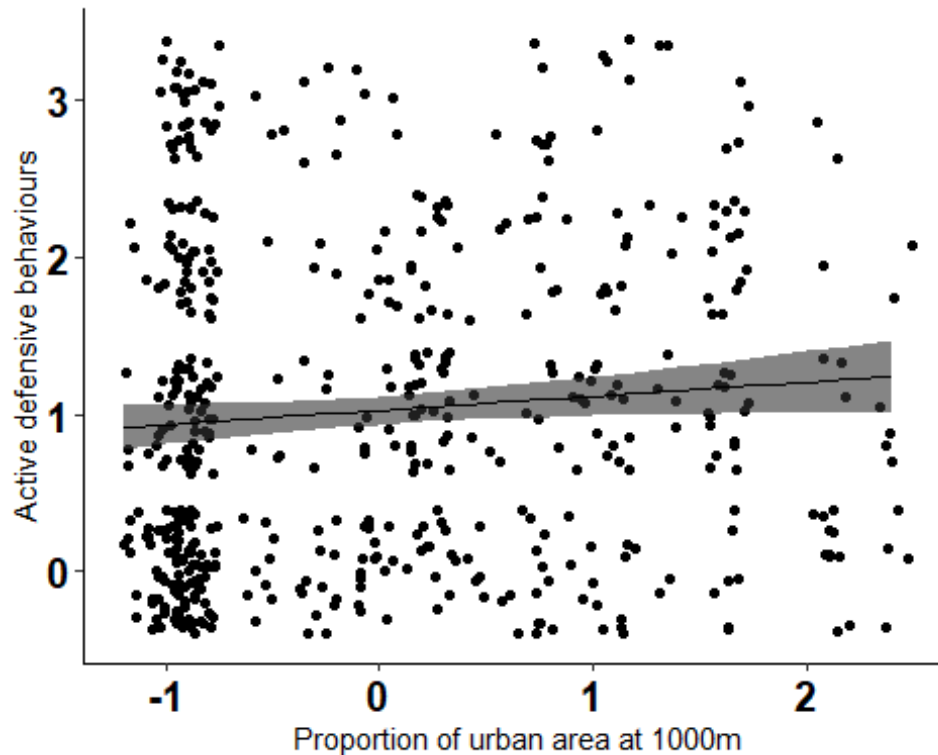
## Figure for proportion of urban area at 1000m

Data was jittered.

```
graph.con.Aggression.urban.1000 <- ggplot(data=pred.con.model.Aggression.urba
n.1000, aes(x, predicted)) + geom_point(data=MixedData, aes(urban.1000.Scaled
, Aggression), position = position_jitter(width = 0.1)) +
  geom_ribbon(data=pred.con.model.Aggression.urban.1000, aes(ymin=conf.low, y
max= conf.high), alpha=0.6) +
  geom_line(data=pred.con.model.Aggression.urban.1000, color="black")+
  theme_bw() + theme(panel.border = element_blank(),
                    panel.grid.major = element_blank(),
                    panel.grid.minor = element_blank(),
                    axis.line = element_line(colour = "black")) +
  theme(axis.text=element_text(size=15, colour="black",face = "bold")) +
  ylab("Active defensive behaviours") +
  xlab("Proportion of urban area at 1000m")

graph.con.Aggression.urban.1000
```





## Model selection

### Time of shell emergence (binary)

#### Random variable

I am testing the significance of turtle ID and site identity by using likelihood ratio tests to see if the addition of these random variables make a significant effect on the initial model. I am using a dummy variable (same value for all the observations) to create a null mixed model to compared with the different combinations of mixed models.

#### Creation of the different mixed models

##### ## Null mixed model

```
mod.Bin.Shell.null <- glmer(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1|Dummy), data = MixedData, family = binomial, control=glmerControl(check.nlev.gtr.1="ignore"), na.action=na.exclude)
```

##### ## only site identity as random variable

```
mod.Bin.Shell.dummy.site <- glmer(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1|Dummy) + (1|Site), data = MixedData, family = binomial, control=glmerControl(check.nlev.gtr.1="ignore"), na.action=na.exclude)
```

*## only turtle ID as random variable*

```
mod.Bin.Shell.dummy.code <- glmer(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1|Dummy) + (1|Code), data = MixedData, family = binomial, control=glmerControl(check.nlev.gtr.1="ignore"), na.action=na.exclude)
```

## Likelihood ratio tests between the mixed models

*#anova with null model and dummy + site model*

```
anova(mod.Bin.Shell.null, mod.Bin.Shell.dummy.site)
```

## Data: MixedData

## Models:

## mod.Bin.Shell.null: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1 | Dummy)

## mod.Bin.Shell.dummy.site: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1 | Dummy) + (1 | Site)

	npars	AIC	BIC	logLik	deviance	Chisq	Df
## mod.Bin.Shell.null	11	636.60	683.09	-307.30	614.60		
## mod.Bin.Shell.dummy.site	12	638.52	689.24	-307.26	614.52	0.0737	1

## Pr(>Chisq)

## mod.Bin.Shell.null

## mod.Bin.Shell.dummy.site 0.786

*#anova with null model and dummy + code model*

```
anova(mod.Bin.Shell.null, mod.Bin.Shell.dummy.code)
```

## Data: MixedData

## Models:

## mod.Bin.Shell.null: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1 | Dummy)

## mod.Bin.Shell.dummy.code: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1 | Dummy) + (1 | Code)

	npars	AIC	BIC	logLik	deviance	Chisq	Df
## mod.Bin.Shell.null	11	636.60	683.09	-307.30	614.60		
## mod.Bin.Shell.dummy.code	12	637.56	688.28	-306.78	613.56	1.0357	1

## Pr(>Chisq)

## mod.Bin.Shell.null

## mod.Bin.Shell.dummy.code 0.3088

If a test has a significant p-value (less than 0.05) then the random effect is significant. Turtle ID (Code) and site identity (Site) are not significant by themselves so I will see if they are more significant together.

### *## Turtle ID without the dummy variable*

```
mod.Bin.Shell.code <- glmer(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1|Code), data = MixedData, family = binomial, control=glmerControl(check.nlev.gtr.1="ignore"), na.action=na.exclude)
```

### *## Site identity without the dummy variable*

```
mod.Bin.Shell.site <- glmer(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1|Site), data = MixedData, family = binomial, control=glmerControl(check.nlev.gtr.1="ignore"), na.action=na.exclude)
```

### *## Turtle and site identity without the dummy variable*

```
mod.Bin.Shell.code.site <- glmer(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1|Code) + (1|Site), data = MixedData, family = binomial, control=glmerControl(check.nlev.gtr.1="ignore"), na.action=na.exclude)
```

### *#anova with code model and code + site model*

```
anova(mod.Bin.Shell.code, mod.Bin.Shell.code.site)
```

```
## Data: MixedData
```

```
## Models:
```

```
## mod.Bin.Shell.code: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1 | Code)
```

```
## mod.Bin.Shell.code.site: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1 | Code) + (1 | Site)
```

```
##          npar    AIC    BIC  logLik deviance Chisq Df Pr(>Chisq)
```

```
## mod.Bin.Shell.code          11 635.56 682.05 -306.78    613.56
```

```
## mod.Bin.Shell.code.site    12 637.56 688.28 -306.78    613.56 9e-04  1      0.9758
```

### *#anova with site model and code + site model*

```
anova(mod.Bin.Shell.site, mod.Bin.Shell.code.site)
```

```
## Data: MixedData
```

```
## Models:
```

```
## mod.Bin.Shell.site: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1 | Site)
```

```
## mod.Bin.Shell.code.site: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled + (1 | Code) + (1 | Site)
```

```
##          npar    AIC    BIC  logLik deviance  Chisq Df
```

```
## mod.Bin.Shell.site          11 636.52 683.01 -307.26    614.52
```

```
## mod.Bin.Shell.code.site    12 637.56 688.28 -306.78    613.56 0.9623  1
```

```
##          Pr(>Chisq)
```

```
## mod.Bin.Shell.site
## mod.Bin.Shell.code.site      0.3266
```

Site and Code are not more significant together ( $p > 0.05$ ), so I will not keep any of the random effects.

## Predictor variables

I am selecting the final model with a backward selection procedure. At each step, I deleted the fixed effect with the highest p value. I confirmed the deletion of each fixed effect with a likelihood ratio test. I created a new dataset at each step to use only the rows with complete observations for all the fixed effects, so that the likelihood ratio tests do not run between two models with a different number of observations.

```
mod.Bin.Shell.full <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled, data = MixedData, family = binomial, na.action=na.exclude)
```

```
summary(mod.Bin.Shell.full)
```

```
##
## Call:
## glm(formula = Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled +
##      Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled +
##      urban.1000.Scaled + water.900.Scaled, family = binomial,
##      data = MixedData, na.action = na.exclude)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.0259  -0.9577  -0.6259   1.0474   2.0174
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   -0.25033    0.16361  -1.530  0.126007
## JulianDate.Scaled -0.48214    0.12832  -3.757  0.000172 ***
## A.Temp.Scaled   -0.40595    0.12447  -3.261  0.001108 **
## Time.Scaled     -0.28796    0.11140  -2.585  0.009743 **
## PL.Scaled        0.20347    0.12329   1.650  0.098851 .
## SexM            -0.06964    0.22757  -0.306  0.759583
## wet.300.Scaled   0.22514    0.11105   2.027  0.042618 *
## agri.1000.Scaled -0.00587    0.10625  -0.055  0.955941
## urban.1000.Scaled -0.24275    0.10823  -2.243  0.024906 *
## water.900.Scaled -0.01244    0.10186  -0.122  0.902753
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
```

```
## Null deviance: 694.8 on 505 degrees of freedom
## Residual deviance: 614.6 on 496 degrees of freedom
## (30 observations deleted due to missingness)
## AIC: 634.6
##
## Number of Fisher Scoring iterations: 4
```

I deleted proportion of agricultural area at 1000m (agri.1000).

```
mod.Bin.Shell.1 <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled + water.900.Scaled, data = MixedData, family = binomial, na.action=na.exclude)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Bin.Shell, MixedData$JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Scaled, MixedData$Sex, MixedData$wet.300.Scaled, MixedData$agri.1000.Scaled, MixedData$urban.1000.Scaled, MixedData$water.900.Scaled),]
```

```
mod.full.adjust <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled + water.900.Scaled, data = MixedData.adjust, family = binomial, na.action=na.exclude)
```

```
mod.Bin.Shell.1.adjust <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled + water.900.Scaled, data = MixedData.adjust, family = binomial, na.action=na.exclude)
```

```
anova(mod.Bin.Shell.1.adjust, mod.full.adjust, test="Chisq")
```

```
## Analysis of Deviance Table
```

```
##
```

```
## Model 1: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled + water.900.Scaled
```

```
## Model 2: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + agri.1000.Scaled + urban.1000.Scaled +
```

```
## water.900.Scaled
```

```
## Resid. Df Resid. Dev Df Deviance Pr(>Chi)
```

```
## 1 497 614.6
```

```
## 2 496 614.6 1 0.0030514 0.9559
```

```
summary(mod.Bin.Shell.1)
```

```
##
```

```
## Call:
```

```
## glm(formula = Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled + water.900.Scaled, family = binomial, data = MixedData, na.action = na.exclude)
```

```
##
```

```
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.0238  -0.9571  -0.6261   1.0475   2.0150
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   -0.25013    0.16358  -1.529 0.126249
## JulianDate.Scaled -0.48226    0.12829  -3.759 0.000171 ***
## A.Temp.Scaled   -0.40671    0.12371  -3.288 0.001010 **
## Time.Scaled     -0.28691    0.10974  -2.615 0.008935 **
## PL.Scaled        0.20250    0.12203   1.659 0.097019 .
## SexM            -0.06972    0.22760  -0.306 0.759356
## wet.300.Scaled   0.22714    0.10497   2.164 0.030468 *
## urban.1000.Scaled -0.24161    0.10623  -2.274 0.022949 *
## water.900.Scaled -0.01189    0.10138  -0.117 0.906634
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 694.8  on 505  degrees of freedom
## Residual deviance: 614.6  on 497  degrees of freedom
## (30 observations deleted due to missingness)
## AIC: 632.6
##
## Number of Fisher Scoring iterations: 4
```

I deleted proportion of open water at 900m (water.900).

```
mod.Bin.Shell.2 <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled, data = MixedData, family = binomial, na.action=na.exclude)

MixedData.adjust <- MixedData[complete.cases(MixedData$Bin.Shell, MixedData$JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Scaled, MixedData$Sex, MixedData$wet.300.Scaled, MixedData$urban.1000.Scaled, MixedData$water.900.Scaled),]

mod.full.adjust <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled + water.900.Scaled, data = MixedData.adjust, family = binomial, na.action=na.exclude)

mod.Bin.Shell.2.adjust <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled, data = MixedData.adjust, family = binomial, na.action=na.exclude)

anova(mod.Bin.Shell.2.adjust, mod.full.adjust, test="Chisq")

## Analysis of Deviance Table
##
```

```
## Model 1: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
##   PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled
## Model 2: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
##   PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled + water.900.Scaled
##   Resid. Df Resid. Dev Df Deviance Pr(>Chi)
## 1      498      614.61
## 2      497      614.60  1 0.013799  0.9065

summary(mod.Bin.Shell.2)

##
## Call:
## glm(formula = Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled +
##   Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled,
##   family = binomial, data = MixedData, na.action = na.exclude)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.0239  -0.9561  -0.6238   1.0490   2.0177
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   -0.24913    0.16337  -1.525 0.127276
## JulianDate.Scaled -0.48497    0.12614  -3.845 0.000121 ***
## A.Temp.Scaled   -0.40646    0.12365  -3.287 0.001012 **
## Time.Scaled     -0.28742    0.10970  -2.620 0.008792 **
## PL.Scaled        0.20064    0.12100   1.658 0.097284 .
## SexM            -0.07128    0.22721  -0.314 0.753732
## wet.300.Scaled   0.22964    0.10279   2.234 0.025486 *
## urban.1000.Scaled -0.23916    0.10415  -2.296 0.021655 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 694.80  on 505  degrees of freedom
## Residual deviance: 614.61  on 498  degrees of freedom
##   (30 observations deleted due to missingness)
## AIC: 630.61
##
## Number of Fisher Scoring iterations: 4
```

I deleted turtle sex (Sex).

```
mod.Bin.Shell.3 <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
  PL.Scaled + wet.300.Scaled + urban.1000.Scaled, data = MixedData, family = binomial, na.action=na.exclude)

MixedData.adjust <- MixedData[complete.cases(MixedData$Bin.Shell, MixedData$JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Scaled, MixedData$wet.300.Scaled, MixedData$urban.1000.Scaled), ]
```

```
L.Scaled, MixedData$Sex, MixedData$wet.300.Scaled, MixedData$urban.1000.Scaled),]
```

```
mod.full.adjust <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled, data = MixedData.adjust, family = binomial, na.action=na.exclude)
```

```
mod.Bin.Shell.3.adjust <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + wet.300.Scaled + urban.1000.Scaled, data = MixedData.adjust, family = binomial, na.action=na.exclude)
```

```
anova(mod.Bin.Shell.3.adjust, mod.full.adjust, test="Chisq")
```

```
## Analysis of Deviance Table
```

```
##
```

```
## Model 1: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +  
##      PL.Scaled + wet.300.Scaled + urban.1000.Scaled
```

```
## Model 2: Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +  
##      PL.Scaled + Sex + wet.300.Scaled + urban.1000.Scaled
```

```
##      Resid. Df Resid. Dev Df Deviance Pr(>Chi)
```

```
## 1          499        614.71
```

```
## 2          498        614.61  1 0.098397  0.7538
```

```
summary(mod.Bin.Shell.3)
```

```
##
```

```
## Call:
```

```
## glm(formula = Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled +  
##      Time.Scaled + PL.Scaled + wet.300.Scaled + urban.1000.Scaled,  
##      family = binomial, data = MixedData, na.action = na.exclude)
```

```
##
```

```
## Deviance Residuals:
```

```
##      Min       1Q   Median       3Q      Max  
## -2.0034  -0.9421  -0.6419   1.0446   2.0357
```

```
##
```

```
## Coefficients:
```

```
##              Estimate Std. Error z value Pr(>|z|)  
## (Intercept)   -0.30912    0.09807  -3.152 0.001621 **  
## JulianDate.Scaled -0.47153    0.12449  -3.788 0.000152 ***  
## A.Temp.Scaled  -0.40945    0.12254  -3.341 0.000834 ***  
## Time.Scaled    -0.28586    0.10878  -2.628 0.008590 **  
## PL.Scaled       0.24691    0.10480   2.356 0.018475 *  
## wet.300.Scaled  0.23790    0.10254   2.320 0.020334 *  
## urban.1000.Scaled -0.22537    0.10310  -2.186 0.028823 *
```

```
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
##
```

```
## (Dispersion parameter for binomial family taken to be 1)
```

```
##
```

```
##      Null deviance: 700.61  on 510  degrees of freedom
```



```
## Residual deviance: 620.61 on 504 degrees of freedom
## (25 observations deleted due to missingness)
## AIC: 634.61
##
## Number of Fisher Scoring iterations: 4
```

All of the fixed effects are statistically significant, so I will stop the backwards selection process.

## Final model

### Summary statistics

Data was jittered.

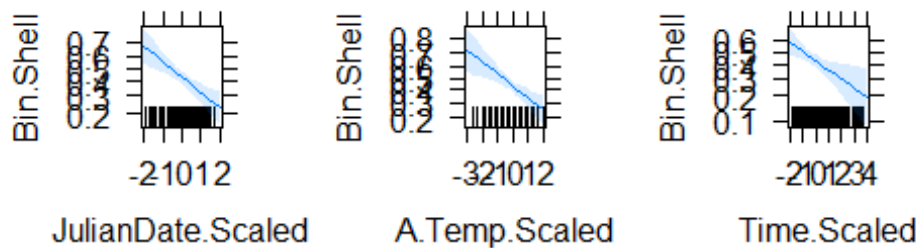
```
mod.Bin.Shell.final <- glm(Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + wet.300.Scaled + urban.1000.Scaled, data = MixedData.adjust, family = binomial, na.action=na.exclude)
summary(mod.Bin.Shell.final)

##
## Call:
## glm(formula = Bin.Shell ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + wet.300.Scaled + urban.1000.Scaled, family = binomial, data = MixedData.adjust, na.action = na.exclude)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.0117  -0.9527  -0.6273   1.0516   2.0170
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -0.2900     0.0986  -2.942  0.003263 **
## JulianDate.Scaled -0.4807     0.1253  -3.837  0.000124 ***
## A.Temp.Scaled   -0.4093     0.1233  -3.320  0.000901 ***
## Time.Scaled     -0.2906     0.1092  -2.662  0.007777 **
## PL.Scaled        0.2189     0.1063   2.060  0.039428 *
## wet.300.Scaled   0.2307     0.1027   2.245  0.024744 *
## urban.1000.Scaled -0.2364     0.1037  -2.279  0.022651 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 694.80 on 505 degrees of freedom
## Residual deviance: 614.71 on 499 degrees of freedom
## AIC: 628.71
##
## Number of Fisher Scoring iterations: 4
```

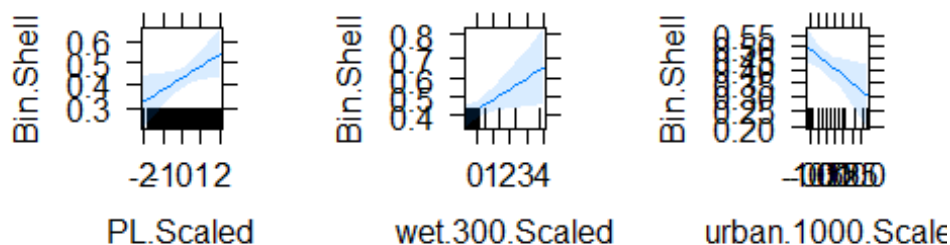
## Visualization of the predictor effects

```
plot(allEffects(mod.Bin.Shell.final,))
```

**Date.Scaled effect** **Temp.Scaled effect** **Time.Scaled effect**



**L.Scaled effect** **wet.300.Scaled effect** **urban.1000.Scaled effect**



## Calculation of the marginal and conditional variance explained by the final model

```
r.squaredGLMM(mod.Bin.Shell.final)
```

```
##               R2m      R2c
## theoretical 0.1955055 0.1955055
## delta      0.1647505 0.1647505
```

Marginal R2: fixed effects R2. Conditional R2: fixed and random effects R2. The delta method can be used with all distributions and link functions.

## Calculation of the 95% confidence intervals

```
confint(mod.Bin.Shell.final, level = 0.95, method = "Wald")
```

```
##               2.5 %      97.5 %
## (Intercept)  -0.48486594 -0.09791228
## JulianDate.Scaled -0.72953960 -0.23753990
## A.Temp.Scaled   -0.65442480 -0.17004497
## Time.Scaled     -0.51113185 -0.08181381
## PL.Scaled       0.01175867  0.42895858
## wet.300.Scaled  0.03024778  0.43466269
## urban.1000.Scaled -0.44148747 -0.03422159
```

## Creation of the prediction figure for urban area at 1000m

```
pred.con.model.Bin.Shell.final <- ggpredict(mod.Bin.Shell.final, terms = "urban.1000.Scaled")
pred.con.model.Bin.Shell.final

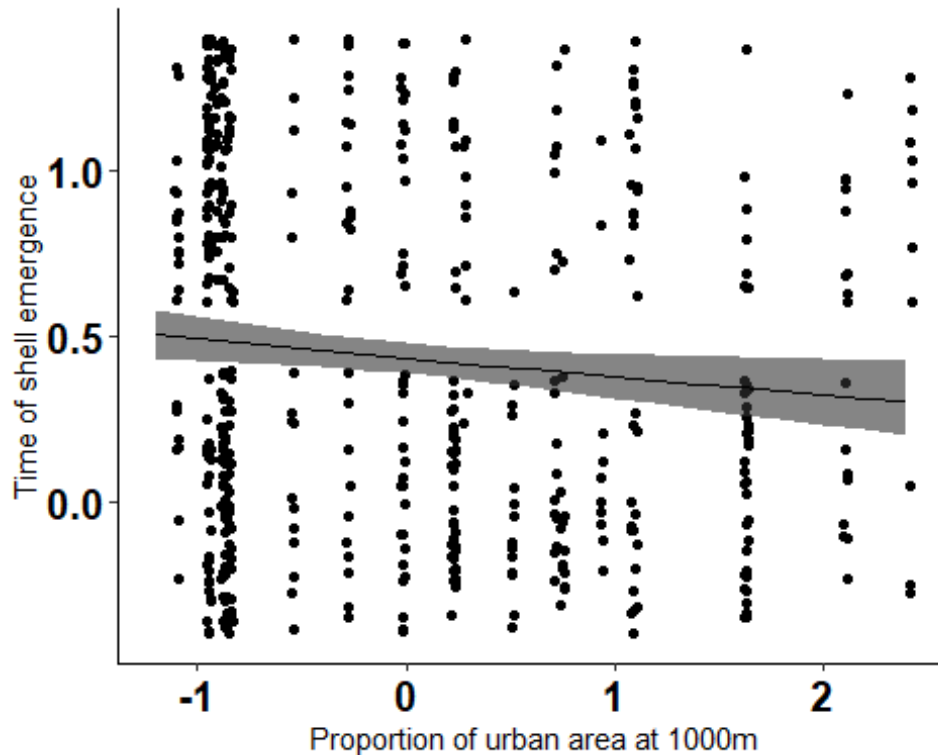
## # Predicted probabilities of Bin.Shell
##
## urban.1000.Scaled | Predicted |          95% CI
## -----|-----|-----
##          -1.20 |          0.50 | [0.43, 0.58]
##          -0.80 |          0.48 | [0.42, 0.54]
##          -0.20 |          0.44 | [0.40, 0.49]
##           0.20 |          0.42 | [0.37, 0.47]
##           0.60 |          0.40 | [0.34, 0.46]
##           1.00 |          0.38 | [0.31, 0.45]
##           1.40 |          0.35 | [0.28, 0.44]
##           2.40 |          0.30 | [0.20, 0.43]
##
## Adjusted for:
## * JulianDate.Scaled = 0.02
## *       A.Temp.Scaled = -0.02
## *       Time.Scaled = -0.02
## *       PL.Scaled = 0.06
## *       wet.300.Scaled = 0.01
```

## Figure for proportion of urban area at 1000m

Data was jittered.

```
graph.con.Bin.Shell.urban.1000 <- ggplot(data=pred.con.model.Bin.Shell.final,
aes(x, predicted)) + geom_point(data=MixedData, aes(urban.1000.Scaled, Bin.Shell),
position = position_jitter(width = 0.01)) +
  geom_ribbon(data=pred.con.model.Bin.Shell.final, aes(ymin=conf.low, ymax=conf.high),
alpha=0.6) +
  geom_line(data=pred.con.model.Bin.Shell.final, color="black")+
  theme_bw() + theme(panel.border = element_blank(),
    panel.grid.major = element_blank(),
    panel.grid.minor = element_blank(),
    axis.line = element_line(colour = "black")) +
  theme(axis.text=element_text(size=15, colour="black",face = "bold")) +
  ylab("Time of shell emergence") +
  xlab("Proportion of urban area at 1000m")

graph.con.Bin.Shell.urban.1000
```



## Model selection

### Time of initial movement

#### Random variable

I am testing the significance of turtle ID and site identity by using likelihood ratio tests to see if the addition of these random variables make a significant effect on the initial model. I am using a dummy variable (same value for all the observations) to create a null mixed model to compared with the different combinations of mixed models.

#### Creation of the different mixed models

##### ## Null mixed model

```
mod.Start.null <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Dummy), data = MixedData, na.action = na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

##### ## only site identity as random variable

```
mod.Start.dummy.site <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Dummy) + (1|Site), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

*## only turtle ID as random variable*

```
mod.Start.dummy.code <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time
.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Sc
aled + urban.1000.Scaled + water.500.Scaled + (1|Dummy) + (1|Code), data = Mi
xedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore")
, REML = FALSE)
```

## Likelihood ratio tests between the mixed models

*#anova with null model and dummy + site model*

```
anova(mod.Start.null, mod.Start.dummy.site)
```

## Data: MixedData

## Models:

## mod.Start.null: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +  
PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urb  
an.1000.Scaled + water.500.Scaled + (1 | Dummy)

## mod.Start.dummy.site: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Sca  
led + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled  
+ urban.1000.Scaled + water.500.Scaled + (1 | Dummy) + (1 | Site)

##

	npar	AIC	BIC	logLik	deviance	Chisq	Df	Pr(>Chi sq)
--	------	-----	-----	--------	----------	-------	----	----------------

## mod.Start.null	13	5633.0	5686.6	-2803.5	5607.0			
-------------------	----	--------	--------	---------	--------	--	--	--

## mod.Start.dummy.site	14	5622.5	5680.2	-2797.2	5594.5	12.539	1	0.0003 985
-------------------------	----	--------	--------	---------	--------	--------	---	---------------

##

## mod.Start.null

## mod.Start.dummy.site \*\*\*

## ---

## Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

*#anova with null model and dummy + code model*

```
anova(mod.Start.null, mod.Start.dummy.code)
```

## Data: MixedData

## Models:

## mod.Start.null: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +  
PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urb  
an.1000.Scaled + water.500.Scaled + (1 | Dummy)

## mod.Start.dummy.code: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Sca  
led + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled  
+ urban.1000.Scaled + water.500.Scaled + (1 | Dummy) + (1 | Code)

##

	npar	AIC	BIC	logLik	deviance	Chisq	Df	Pr(>Chi sq)
--	------	-----	-----	--------	----------	-------	----	----------------

## mod.Start.null	13	5633.0	5686.6	-2803.5	5607.0			
-------------------	----	--------	--------	---------	--------	--	--	--

## mod.Start.dummy.code	14	5633.2	5690.9	-2802.6	5605.2	1.8135	1	0.1 781
-------------------------	----	--------	--------	---------	--------	--------	---	------------

If a test has a significant p-value (less than 0.05) then the random effect is significant. Site identity (Site) is significant by itself but turtle identity (Code) is not. I will see if Site and Code together are more significant than Site by itself.

### *## Site identity without the dummy variable*

```
mod.Start.site <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

### *## Turtle and site identity without the dummy variable*

```
mod.Start.code.site <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Code) + (1|Site), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

### *#anova with site model and code + site model*

```
anova(mod.Start.site, mod.Start.code.site)
```

```
## Data: MixedData
```

```
## Models:
```

```
## mod.Start.site: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urban.1000.Scaled + water.500.Scaled + (1 | Site)
```

```
## mod.Start.code.site: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urban.1000.Scaled + water.500.Scaled + (1 | Code) + (1 | Site)
```

```
##          npar    AIC    BIC logLik deviance Chisq Df Pr(>Chisq)
```

```
## mod.Start.site      13 5620.5 5674.0 -2797.2    5594.5
```

```
## mod.Start.code.site  14 5621.6 5679.2 -2796.8    5593.6 0.9154  1      0.33
```

```
87
```

Site and Code together are not more significant than Site by itself ( $p > 0.05$ ), so I will only keep Site.

## Predictor variables

I am selecting the final model with a backward selection procedure. At each step, I deleted the fixed effect with the highest p value. I confirmed the deletion of each fixed effect with a likelihood ratio test. I created a new dataset at each step to use only the rows with complete observations for all the fixed effects, so that the likelihood ratio tests do not run between two models with a different number of observations.

```
mod.Start.full <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```

summary(mod.Start.full)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled +
## Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled +
## urban.1000.Scaled + water.500.Scaled + (1 | Site)
## Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
## 5620.5  5674.0 -2797.3  5594.5      441
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.7795 -0.6498 -0.2979  0.4847  4.2002
##
## Random effects:
## Groups Name Variance Std.Dev.
## Site (Intercept) 1300 36.05
## Residual 12480 111.71
## Number of obs: 454, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    121.563    12.188  42.712   9.974 1.01e-12 ***
## JulianDate.Scaled -18.702    11.118  24.906  -1.682  0.10504
## A.Temp.Scaled    -3.933     9.352 115.551  -0.421  0.67485
## Time.Scaled     -5.669     5.832 447.861  -0.972  0.33151
## PL.Scaled       18.916     6.836 451.192   2.767  0.00589 **
## SexM             3.427    12.723 449.098   0.269  0.78779
## for.veg.200.Scaled -15.890    16.774  19.514  -0.947  0.35506
## wet.400.Scaled   17.139    15.218  18.677   1.126  0.27435
## agri.600.Scaled   1.413    10.337  22.144   0.137  0.89250
## urban.1000.Scaled -16.261    15.873  18.101  -1.024  0.31912
## water.500.Scaled -12.932    12.231  18.034  -1.057  0.30432
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S Tm.Sc1 PL.Sc1 SexM f..200 w.400. a.600.
## JulnDt.Sc1d -0.066
## A.Temp.Sc1d  0.110 -0.399
## Time.Sc1d   0.053  0.095 -0.233
## PL.Sc1d    -0.307 -0.020 -0.041 -0.035
## SexM       -0.604  0.049 -0.113 -0.059  0.472
## fr.vg.200.S -0.013 -0.109 -0.257  0.048 -0.016  0.059
## wt.400.Sc1d -0.063 -0.315 -0.116  0.036  0.011  0.046  0.704
## agr.600.Sc1 -0.092 -0.060 -0.172  0.067 -0.060  0.009  0.579  0.541

```

```
## urbn.1000.S -0.077 -0.218 -0.120 0.055 -0.018 0.067 0.750 0.725 0.495
## wtr.500.Scl -0.042 -0.179 -0.146 0.019 -0.072 0.011 0.581 0.595 0.418
##          u.1000
## JulnDt.Scl
## A.Temp.Scl
## Time.Scaled
## PL.Scaled
## SexM
## fr.vg.200.S
## wt.400.Scl
## agr.600.Scl
## urbn.1000.S
## wtr.500.Scl 0.619
```

I deleted proportion of agricultural area at 600m (agri.600).

```
mod.Start.1 <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + w
ater.500.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FA
LSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$Julia
nDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Sc
aled, MixedData$Sex, MixedData$for.veg.200.Scaled, MixedData$wet.400.Scaled,
MixedData$agri.600.Scaled, MixedData$urban.1000.Scaled, MixedData$water.500.S
caled),]
```

```
mod.full.adjust <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scal
ed + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled
+ urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData.adjust, n
a.action=na.exclude, REML = FALSE)
```

```
mod.Start.1.adjust <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.S
caled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Sc
aled + water.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exc
lude, REML = FALSE)
```

```
anova(mod.Start.1.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Start.1.adjust: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scale
d + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled
+ water.500.Scaled + (1 | Site)
```

```
## mod.full.adjust: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + agri.600.Scaled + urb
an.1000.Scaled + water.500.Scaled + (1 | Site)
```

```
##          npar    AIC    BIC  logLik deviance  Chisq Df Pr(>Chisq
)
```

```
## mod.Start.1.adjust    12 5618.5 5667.9 -2797.3   5594.5
```



```

## mod.full.adjust      13 5620.5 5674.0 -2797.2   5594.5 0.0186 1      0.891
4

summary(mod.Start.1)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scal
ed +
## Sex + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled +
## water.500.Scaled + (1 | Site)
## Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
##    5618.5    5667.9  -2797.3   5594.5     442
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.7825 -0.6512 -0.2920  0.4800  4.2021
##
## Random effects:
## Groups Name Variance Std.Dev.
## Site (Intercept) 1305 36.12
## Residual 12479 111.71
## Number of obs: 454, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    121.725     12.146   43.855   10.022 6.47e-13 ***
## JulianDate.Scaled -18.619     11.109   25.189    -1.676  0.10612
## A.Temp.Scaled    -3.713      9.216  105.768    -0.403  0.68786
## Time.Scaled     -5.716      5.819  446.144    -0.982  0.32644
## PL.Scaled       18.963      6.823  452.210     2.779  0.00568 **
## SexM             3.413     12.722  449.091     0.268  0.78864
## for.veg.200.Scaled -17.212     13.691   22.126    -1.257  0.22178
## wet.400.Scaled    16.014     12.810   20.789     1.250  0.22514
## urban.1000.Scaled -17.332     13.808   20.578    -1.255  0.22345
## water.500.Scaled  -13.628     11.122   18.941    -1.225  0.23548
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnDt.S A.Tm.S Tm.Scl PL.Scl SexM f..200 w.400. u.1000
## JulnDt.Scl -0.072
## A.Temp.Scl 0.096 -0.416
## Time.Scaled 0.060 0.099 -0.226
## PL.Scaled -0.314 -0.024 -0.052 -0.031
## SexM -0.606 0.050 -0.113 -0.060 0.473
## fr.vg.200.S 0.050 -0.092 -0.195 0.011 0.023 0.066
## wt.400.Scl -0.016 -0.336 -0.027 -0.001 0.052 0.050 0.570

```

```
## urbn.1000.S -0.037 -0.217 -0.041 0.026 0.013 0.072 0.654 0.626
## wtr.500.Scl -0.004 -0.170 -0.083 -0.010 -0.051 0.008 0.458 0.483 0.522
```

I deleted turtle sex (Sex).

```
mod.Start.2 <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Scaled, MixedData$Sex, MixedData$for.veg.200.Scaled, MixedData$wet.400.Scaled, MixedData$urban.1000.Scaled, MixedData$water.500.Scaled),]
```

```
mod.full.adjust <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.Start.2.adjust <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.Start.2.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Start.2.adjust: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1 | Site)
```

```
## mod.full.adjust: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1 | Site)
```

```
##          npar    AIC    BIC  logLik deviance  Chisq Df Pr(>Chisq)
```

```
## mod.Start.2.adjust    11 5616.6 5661.9 -2797.3    5594.6
```

```
## mod.full.adjust      12 5618.5 5667.9 -2797.3    5594.5 0.0719  1      0.7885
```

```
summary(mod.Start.2)
```

```
## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
```

```
## Formula: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled +
```

```
##      for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled +
```

```
##      water.500.Scaled + (1 | Site)
```

```
## Data: MixedData
```

```
##
```

```
##      AIC      BIC  logLik deviance df.resid
```

```
##    5673.8    5719.2   -2825.9    5651.8        448
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.7853 -0.6524 -0.3028  0.4765  4.2409
##
## Random effects:
##   Groups      Name            Variance Std.Dev.
##   Site      (Intercept)    1326      36.41
##   Residual                12354     111.15
## Number of obs: 459, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    123.672      9.686   18.364  12.768 1.42e-10 ***
## JulianDate.Scaled -18.545     11.105   25.227  -1.670  0.10729
## A.Temp.Scaled    -3.860      9.103  111.426  -0.424  0.67239
## Time.Scaled      -5.444      5.773  449.706  -0.943  0.34621
## PL.Scaled        18.144      5.860  458.871   3.096  0.00208 **
## for.veg.200.Scaled -16.984     13.674   22.029  -1.242  0.22729
## wet.400.Scaled    15.834     12.834   20.853   1.234  0.23101
## urban.1000.Scaled -17.532     13.806   20.520  -1.270  0.21836
## water.500.Scaled  -13.818     11.085   18.582  -1.247  0.22806
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S Tm.Scl PL.Scl f..200 w.400. u.1000
## JulnDt.Scl -0.053
## A.Temp.Scl  0.036 -0.411
## Time.Scaled 0.032  0.102 -0.236
## PL.Scaled   -0.029 -0.050 -0.003 -0.012
## fr.vg.200.S 0.112 -0.098 -0.186  0.016  0.000
## wt.400.Scl  0.017 -0.340 -0.022  0.003  0.035  0.569
## urbn.1000.S 0.008 -0.222 -0.033  0.031 -0.020  0.652  0.625
## wtr.500.Scl -0.004 -0.176 -0.081 -0.007 -0.041  0.458  0.485  0.525
```

I deleted air temperature (A.Temp).

```
mod.Start.3 <- lmer(Start ~ JulianDate.Scaled + Time.Scaled + PL.Scaled + for
.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1|
Site), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$Julia
nDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Sc
aled, MixedData$for.veg.200.Scaled, MixedData$wet.400.Scaled, MixedData$urban
.1000.Scaled, MixedData$water.500.Scaled),]
```

```
mod.full.adjust <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scal
ed + PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + wa
```

```

ter.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

mod.Start.3.adjust <- lmer(Start ~ JulianDate.Scaled + Time.Scaled + PL.Scaled +
d + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled +
d + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

anova(mod.Start.3.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.Start.3.adjust: Start ~ JulianDate.Scaled + Time.Scaled + PL.Scaled +
for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled +
(1 | Site)
## mod.full.adjust: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.5
00.Scaled + (1 | Site)
##
##          npar      AIC      BIC  logLik deviance  Chisq Df Pr(>Chisq
)
## mod.Start.3.adjust    10 5672.0 5713.3 -2826.0    5652.0
## mod.full.adjust      11 5673.8 5719.2 -2825.9    5651.8 0.1797  1      0.671
6

summary(mod.Start.3)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula:
## Start ~ JulianDate.Scaled + Time.Scaled + PL.Scaled + for.veg.200.Scaled +
##      wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1 |      Site
)
##      Data: MixedData
##
##           AIC          BIC    logLik deviance df.resid
##    5672.0    5713.3   -2826.0    5652.0      449
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.7806 -0.6478 -0.2925  0.4666  4.2266
##
## Random effects:
##      Groups      Name              Variance Std.Dev.
##      Site      (Intercept)    1331      36.49
##      Residual                12357     111.16
## Number of obs: 459, groups: Site, 23
##
## Fixed effects:
##
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    123.830      9.692   18.514  12.776 1.26e-10 ***
## JulianDate.Scaled -20.486     10.139   20.873  -2.021  0.0563 .

```

```
## Time.Scaled          -6.015          5.611 452.475  -1.072    0.2843
## PL.Scaled            18.129          5.862 458.881   3.093    0.0021 **
## for.veg.200.Scaled  -18.054         13.454  21.081  -1.342    0.1939
## wet.400.Scaled       15.719         12.847  20.991   1.224    0.2347
## urban.1000.Scaled   -17.721         13.817  20.525  -1.283    0.2139
## water.500.Scaled    -14.197         11.064  18.379  -1.283    0.2154
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) JlnDt.S Tm.Scl PL.Scl f..200 w.400. u.1000
## JulnDt.Scl -0.042
## Time.Scaled  0.041  0.006
## PL.Scaled   -0.029 -0.056 -0.013
## fr.vg.200.S  0.121 -0.194 -0.029  0.000
## wt.400.Scl   0.018 -0.383 -0.003  0.035  0.575
## urbn.1000.S  0.009 -0.258  0.024 -0.020  0.658  0.625
## wtr.500.Scl -0.001 -0.231 -0.027 -0.041  0.452  0.484  0.524
```

I deleted time of testing (Time).

```
mod.Start.4 <- lmer(Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$JulianDate.Scaled, MixedData$Time.Scaled, MixedData$PL.Scaled, MixedData$for.veg.200.Scaled, MixedData$wet.400.Scaled, MixedData$urban.1000.Scaled, MixedData$water.500.Scaled),]
```

```
mod.full.adjust <- lmer(Start ~ JulianDate.Scaled + Time.Scaled + PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.Start.4.adjust <- lmer(Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.Start.4.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Start.4.adjust: Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1 | Site)
```

```
## mod.full.adjust: Start ~ JulianDate.Scaled + Time.Scaled + PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1 | Site)
```

```
##          npar    AIC    BIC  logLik deviance  Chisq Df Pr(>Chisq)
```

```
## mod.Start.4.adjust    9 5671.1 5708.3 -2826.6   5653.1
```

```

## mod.full.adjust      10 5672.0 5713.3 -2826.0   5652.0 1.1215  1    0.289
6

summary(mod.Start.4)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled +
##      wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1 |      Site
## )
## Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
##  5671.1   5708.3  -2826.5   5653.1     450
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.8000 -0.6361 -0.2864  0.4597  4.2948
##
## Random effects:
## Groups   Name                Variance Std.Dev.
## Site     (Intercept)      1439      37.94
## Residual                    12356     111.16
## Number of obs: 459, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    124.468     9.945  19.395  12.516 9.68e-11 ***
## JulianDate.Scaled -20.603    10.396  21.741  -1.982  0.06028 .
## PL.Scaled        17.880     5.867 458.649   3.047  0.00244 **
## for.veg.200.Scaled -18.337    13.788  21.868  -1.330  0.19725
## wet.400.Scaled    15.734    13.173  21.805   1.194  0.24513
## urban.1000.Scaled -17.283    14.166  21.472  -1.220  0.23568
## water.500.Scaled  -14.425    11.359  19.172  -1.270  0.21930
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S PL.Scl f..200 w.400. u.1000
## JulnDt.Scld -0.042
## PL.Scaled   -0.028 -0.054
## fr.vg.200.S 0.126 -0.191 -0.001
## wt.400.Scld 0.019 -0.382  0.035  0.574
## urbn.1000.S 0.009 -0.258 -0.019  0.658  0.625
## wtr.500.Scl 0.001 -0.232 -0.041  0.450  0.485  0.525

```

I removed proportion of wetland area at 400m (wet.400).

```

mod.Start.5 <- lmer(Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)

```

```

MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$JulianDate.Scaled, MixedData$PL.Scaled, MixedData$for.veg.200.Scaled, MixedData$wet.400.Scaled, MixedData$urban.1000.Scaled, MixedData$water.500.Scaled),]

mod.full.adjust <- lmer(Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

mod.Start.5.adjust <- lmer(Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

anova(mod.Start.5.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.Start.5.adjust: Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled + water.500.Scaled + (1 | Site)
## mod.full.adjust: Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1 | Site)
##
##          npar    AIC    BIC  logLik deviance  Chisq Df Pr(>Chisq)
## mod.Start.5.adjust      8 5670.5 5703.5 -2827.2   5654.5
## mod.full.adjust        9 5671.1 5708.3 -2826.6   5653.1 1.3923  1      0.238
##
summary(mod.Start.5)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled +
##          urban.1000.Scaled + water.500.Scaled + (1 | Site)
## Data: MixedData
##
##          AIC          BIC   logLik deviance df.resid
## 5670.5    5703.5  -2827.2   5654.5      451
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.7520 -0.6381 -0.2847  0.4564  4.3022
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
##  Site      (Intercept)    1566      39.58
##  Residual                    12359    111.17
## Number of obs: 459, groups: Site, 23
##
## Fixed effects:
##
##              Estimate Std. Error      df t value Pr(>|t|)

```

```
## (Intercept)          124.448      10.241  19.456  12.152 1.55e-10 ***
## JulianDate.Scaled    -16.049       9.885  20.932  -1.624  0.11942
## PL.Scaled            17.468       5.872 458.468   2.975  0.00309 **
## for.veg.200.Scaled   -27.653      11.626  20.165  -2.378  0.02738 *
## urban.1000.Scaled    -27.792      11.379  21.202  -2.442  0.02343 *
## water.500.Scaled     -20.940      10.236  19.145  -2.046  0.05478 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) JlnDt.S PL.Scl f..200 u.1000
## JulnDt.Scl d -0.036
## PL.Scaled    -0.029 -0.043
## fr.vg.200.S  0.143  0.039 -0.025
## urbn.1000.S  -0.004 -0.027 -0.051  0.469
## wtr.500.Scl  -0.009 -0.060 -0.064  0.238  0.326
```

I removed Julian date of testing (JulianDate).

```
mod.Start.6 <- lmer(Start ~ PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$JulianDate.Scaled, MixedData$PL.Scaled, MixedData$for.veg.200.Scaled, MixedData$urban.1000.Scaled, MixedData$water.500.Scaled),]
```

```
mod.full.adjust <- lmer(Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.Start.6.adjust <- lmer(Start ~ PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.Start.6.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Start.6.adjust: Start ~ PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled + water.500.Scaled + (1 | Site)
```

```
## mod.full.adjust: Start ~ JulianDate.Scaled + PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled + water.500.Scaled + (1 | Site)
```

```
##          npar    AIC    BIC  logLik deviance  Chisq Df Pr(>Chisq)
```

```
## mod.Start.6.adjust      7 5671.1 5700.0 -2828.5   5657.1
```

```
## mod.full.adjust        8 5670.5 5703.5 -2827.2   5654.5 2.5754  1      0.1085
```

```
summary(mod.Start.6)
```



```

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Start ~ PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled +
## water.500.Scaled + (1 | Site)
## Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
## 5671.1  5700.0  -2828.5   5657.1     452
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.7710 -0.6399 -0.3070  0.4709  4.2815
##
## Random effects:
## Groups   Name                Variance Std.Dev.
## Site     (Intercept)         1750     41.83
## Residual                    12383    111.28
## Number of obs: 459, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    124.112    10.650  18.392  11.654 6.26e-10 ***
## PL.Scaled       16.875     5.882 458.345   2.869  0.00431 **
## for.veg.200.Scaled -26.692    12.080  19.150  -2.210  0.03951 *
## urban.1000.Scaled -28.209    11.818  19.997  -2.387  0.02699 *
## water.500.Scaled -21.911    10.635  18.236  -2.060  0.05392 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) PL.Scl f..200 u.1000
## PL.Scaled    -0.030
## fr.vg.200.S  0.149 -0.023
## urbn.1000.S -0.005 -0.050  0.470
## wtr.500.Scl -0.010 -0.065  0.239  0.325

```

I removed proportion of open water area at 500m (water.500).

```

mod.Start.7 <- lmer(Start ~ PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)

```

```

MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$PL.Scaled, MixedData$for.veg.200.Scaled, MixedData$urban.1000.Scaled, MixedData$water.500.Scaled),]

```

```

mod.full.adjust <- lmer(Start ~ PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

```

```

mod.Start.7.adjust <- lmer(Start ~ PL.Scaled + for.veg.200.Scaled + urban.100

```

```

0.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = F
ALSE)

anova(mod.Start.7.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.Start.7.adjust: Start ~ PL.Scaled + for.veg.200.Scaled + urban.1000.Sc
aled + (1 | Site)
## mod.full.adjust: Start ~ PL.Scaled + for.veg.200.Scaled + urban.1000.Scale
d + water.500.Scaled + (1 | Site)
##
##          npar    AIC    BIC  logLik deviance  Chisq Df Pr(>Chisq
)
## mod.Start.7.adjust      6 5672.9 5697.6 -2830.4   5660.9
## mod.full.adjust       7 5671.1 5700.0 -2828.5   5657.1 3.8075  1    0.0510
2 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

summary(mod.Start.7)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Start ~ PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled +
## (1 | Site)
## Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
##  5672.9   5697.6  -2830.4   5660.9      453
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.7411 -0.6467 -0.2877  0.4708  4.2924
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
##  Site      (Intercept)    2302      47.98
##  Residual                    12362    111.19
## Number of obs: 459, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    124.575     11.799  19.883  10.558 1.34e-09 ***
## PL.Scaled       15.786      5.889 457.518   2.681  0.00761 **
## for.veg.200.Scaled -20.325     12.985  20.168  -1.565  0.13308
## urban.1000.Scaled -20.139     12.330  21.432  -1.633  0.11700
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:

```

```
##          (Intr) PL.Scl f..200
## PL.Scaled -0.029
## fr.vg.200.S 0.166 -0.008
## urbn.1000.S -0.002 -0.028 0.428
```

Since the LRT P-value is almost significant (0.051) and the removal of water.500 changed urban.1000 from non-significant to significant, I decided to calculate the AICc values for mod.Start.6 and mod.Start.7 in order to compare their parsimony.

## AIC test

I had to recreate the initial model for initial time of movement for the AIC test.

```
mod.Start.0 <- lmer(Start ~ agri.600.Scaled + JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.200.Scaled + wet.400.Scaled + urban.1000.Scaled + water.500.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)

#define list of models
Start.models <- list(mod.Start.0, mod.Start.1, mod.Start.2, mod.Start.3, mod.Start.4, mod.Start.5, mod.Start.6, mod.Start.7)

#specify model names
Start.mod.names <- c('mod.Start.0', 'mod.Start.1', 'mod.Start.2', 'mod.Start.3', 'mod.Start.4', 'mod.Start.5', 'mod.Start.6', 'mod.Start.7')

#calculate AICc of each model
aictab(cand.set = Start.models, modnames = Start.mod.names)

##
## Model selection based on AICc:
##
##          K      AICc Delta_AICc AICcWt Cum.Wt      LL
## mod.Start.1 12 5619.23      0.00  0.74  0.74 -2797.26
## mod.Start.0 13 5621.33      2.10  0.26  1.00 -2797.25
## mod.Start.5  8 5670.80     51.57  0.00  1.00 -2827.24
## mod.Start.6  7 5671.31     52.08  0.00  1.00 -2828.53
## mod.Start.4  9 5671.49     52.26  0.00  1.00 -2826.55
## mod.Start.3 10 5672.46     53.23  0.00  1.00 -2825.99
## mod.Start.7  6 5673.05     53.82  0.00  1.00 -2830.43
## mod.Start.2 11 5674.38     55.15  0.00  1.00 -2825.90
```

mod.Start.6 and mod.Start.7 had an AICc value within 2 of each other which suggests that they are equally parsimonious, so I decided to average them.

## Averaging mod.Start.6 and mod.Start.7

```
# Average the two models using model.avg
averaged.model.Start <- model.avg(mod.Start.6, mod.Start.7)
```

```
summary(averaged.model.Start)

##
## Call:
## model.avg(object = mod.Start.6, mod.Start.7)
##
## Component model call:
## lmer(formula = <2 unique values>, data = MixedData, REML = FALSE,
##       na.action = na.exclude)
##
## Component models:
##      df  logLik   AICc delta weight
## 1234  7 -2828.53 5671.31  0.00  0.71
## 123   6 -2830.43 5673.05  1.74  0.29
##
## Term codes:
## for.veg.200.Scaled      PL.Scaled  urban.1000.Scaled  water.500.Scale
d
##              1              2              3
4
##
## Model-averaged coefficients:
## (full average)
##              Estimate Std. Error Adjusted SE z value Pr(>|z|)
## (Intercept)    124.249    11.003     11.032  11.262 < 2e-16 ***
## PL.Scaled       16.554     5.905      5.920   2.796  0.00517 **
## for.veg.200.Scaled -24.815    12.690     12.723   1.951  0.05112 .
## urban.1000.Scaled -25.830    12.524     12.555   2.057  0.03965 *
## water.500.Scaled -15.453    13.400     13.416   1.152  0.24938
##
## (conditional average)
##              Estimate Std. Error Adjusted SE z value Pr(>|z|)
## (Intercept)    124.249    11.003     11.032  11.262 < 2e-16 ***
## PL.Scaled       16.554     5.905      5.920   2.796  0.00517 **
## for.veg.200.Scaled -24.815    12.690     12.723   1.951  0.05112 .
## urban.1000.Scaled -25.830    12.524     12.555   2.057  0.03965 *
## water.500.Scaled -21.911    10.635     10.663   2.055  0.03989 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

To create the final model I will extract the conditional average coefficients, as they are only averaged over the models where the fixed effects appear.

```
#Extracting the conditional average coefficients from the averaged model
con.avg.coefs.Start <- averaged.model.Start$coefs[,1]
```

## Final model

### Summary statistics

I changed the REML to TRUE to calculate the summary statistics of the final model.

```
#Creating model with just the conditional average coefficients
con.averaged.model.Start <- lmer(Start ~ PL.Scaled + for.veg.200.Scaled + urb
an.1000.Scaled + water.500.Scaled + (1|Site),
                                data = MixedData,
                                REML = TRUE,
                                na.action = na.exclude,
                                weights = con.avg.coefs.Start)
summary(con.averaged.model.Start)

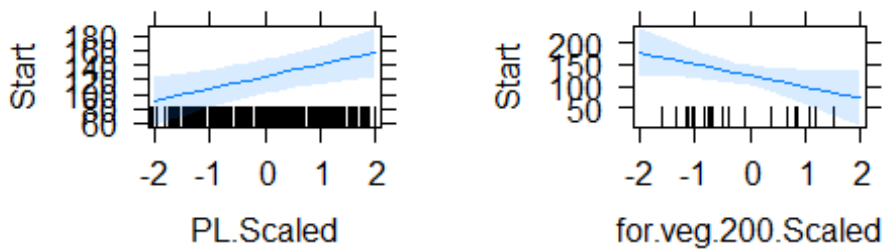
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: Start ~ PL.Scaled + for.veg.200.Scaled + urban.1000.Scaled +
##      water.500.Scaled + (1 | Site)
##      Data: MixedData
## Weights: con.avg.coefs.Start
##
## REML criterion at convergence: 5624.9
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.8024 -0.6303 -0.2929  0.4734  4.2838
##
## Random effects:
##      Groups   Name                Variance Std.Dev.
##      Site     (Intercept)    2380      48.79
##      Residual                    12393     111.32
## Number of obs: 459, groups:  Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    124.820    11.956  15.948  10.440 1.55e-08 ***
## PL.Scaled       16.389     5.909 453.319   2.774 0.00577 **
## for.veg.200.Scaled -26.082    13.534  16.464  -1.927 0.07139 .
## urban.1000.Scaled -28.002    13.211  17.100  -2.120 0.04896 *
## water.500.Scaled -21.845    11.946  15.733  -1.829 0.08648 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) PL.Scl f..200 u.1000
## PL.Scaled    -0.028
## fr.vg.200.S  0.160 -0.022
## urbn.1000.S  -0.005 -0.045  0.469
## wtr.500.Scl -0.008 -0.059  0.234  0.326
```

Proportion of urban area at 1000m (urban.1000) has a statistically significant effect on time of initial movement.

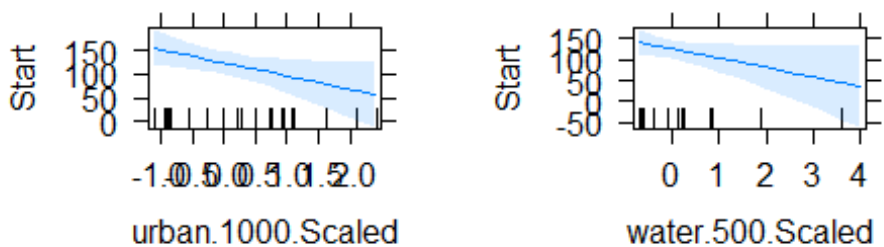
## Visualization of the predictor effects

```
plot(allEffects(con.averaged.model.Start))
```

### PL.Scaled effect plot for.veg.200.Scaled effect plot



### urban.1000.Scaled effect plot water.500.Scaled effect plot



## Calculation of the marginal and conditional variance explained by the final model

```
r.squaredGLMM(con.averaged.model.Start)
```

```
##           R2m           R2c
## [1,] 0.0650692 0.215701
```

Marginal R2: fixed effects R2. Conditional R2: fixed and random effects R2

## Calculation of the 95% confidence intervals

```
confint(con.averaged.model.Start, level = 0.95, method = "Wald")
```

```
##           2.5 %           97.5 %
## .sig01           NA           NA
## .sigma           NA           NA
## (Intercept)    101.387414 148.2528846
## PL.Scaled       4.807912  27.9706575
## for.veg.200.Scaled -52.607602  0.4437976
```

```
## urban.1000.Scaled -53.894410 -2.1098848
## water.500.Scaled -45.259851 1.5689630
```

## Creation of the prediction figure for urban area at 1000m

```
pred.con.averaged.model.Start.urban.1000 <- ggpredict(con.averaged.model.Start, terms = "urban.1000.Scaled")
pred.con.averaged.model.Start.urban.1000
```

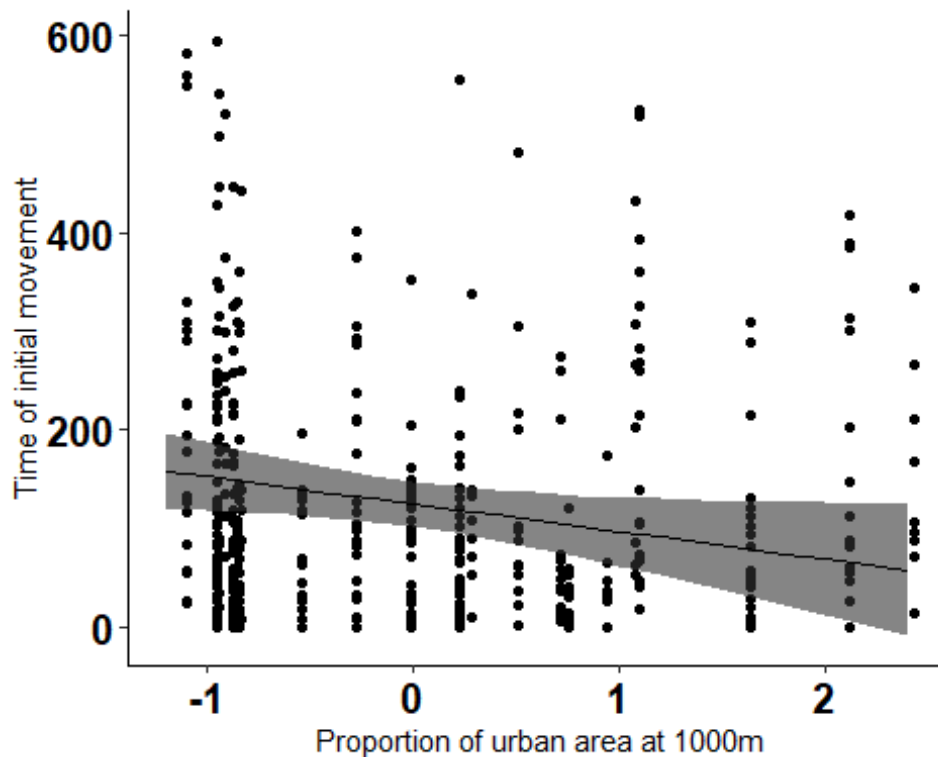
```
## # Predicted values of Start
##
## urban.1000.Scaled | Predicted |          95% CI
## -----|-----|-----
##          -1.20 |    158.07 | [119.26, 196.88]
##          -0.80 |    146.87 | [115.67, 178.07]
##          -0.20 |    130.07 | [106.05, 154.09]
##           0.20 |    118.87 | [ 94.78, 142.96]
##           0.60 |    107.67 | [ 79.41, 135.93]
##           1.00 |     96.47 | [ 61.38, 131.56]
##           1.40 |     85.27 | [ 41.93, 128.61]
##           2.40 |     57.26 | [-9.33, 123.86]
##
## Adjusted for:
## *          PL.Scaled = 0.02
## * for.veg.200.Scaled = 0.01
## *   water.500.Scaled = 0.02
## *          Site = 0 (population-level)
```

```
# New dataset to only have complete observations for all the variables
MixedData.adjust.final.model.Start <- MixedData[complete.cases(MixedData$PL.Scaled, MixedData$for.veg.200.Scaled, MixedData$urban.1000.Scaled, MixedData$water.500.Scaled),]
```

## Figure for proportion of urban area at 1000m

```
graph.con.Start.urban.1000 <- ggplot(data=pred.con.averaged.model.Start.urban.1000, aes(x, predicted)) + geom_point(data=MixedData, aes(urban.1000.Scaled, Start)) +
  geom_ribbon(data=pred.con.averaged.model.Start.urban.1000, aes(ymin=conf.low, ymax= conf.high), alpha=0.6) +
  geom_line(data=pred.con.averaged.model.Start.urban.1000, color="black")+
  theme_bw() + theme(panel.border = element_blank(),
                     panel.grid.major = element_blank(),
                     panel.grid.minor = element_blank(),
                     axis.line = element_line(colour = "black")) +
  theme(axis.text=element_text(size=15, colour="black",face = "bold")) +
  ylab("Time of initial movement") +
  xlab("Proportion of urban area at 1000m")

graph.con.Start.urban.1000
```



## Model selection

### Log(x+1) transformed version of time of initial movement

#### Random variable

I am testing the significance of turtle ID and site identity by using likelihood ratio tests to see if the addition of these random variables make a significant effect on the initial model. I am using a dummy variable (same value for all the observations) to create a null mixed model to compared with the different combinations of mixed models.

#### Creation of the different mixed models

##### ## Null mixed model

```
mod.log.Start.null <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1|Dummy), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

##### ## only site identity as random variable

```
mod.log.Start.dummy.site <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1|Dummy) + (1|Site), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```



```
## only turtle ID as random variable
```

```
mod.log.Start.dummy.code <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1|Dummy) + (1|Code), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

## Likelihood ratio tests between the mixed models

```
#anova with null model and dummy + site model
```

```
anova(mod.log.Start.null, mod.log.Start.dummy.site)
```

```
## Data: MixedData
```

```
## Models:
```

```
## mod.log.Start.null: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1 | Dummy)
```

```
## mod.log.Start.dummy.site: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1 | Dummy) + (1 | Site)
```

```
##               npar      AIC      BIC  logLik deviance  Chisq Df
## mod.log.Start.null          12 1692.3 1741.7 -834.16   1668.3
## mod.log.Start.dummy.site    13 1686.3 1739.8 -830.15   1660.3 8.0127  1
##               Pr(>Chisq)
```

```
## mod.log.Start.null
```

```
## mod.log.Start.dummy.site    0.004645 **
```

```
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
#anova with null model and dummy + code model
```

```
anova(mod.log.Start.null, mod.log.Start.dummy.code)
```

```
## Data: MixedData
```

```
## Models:
```

```
## mod.log.Start.null: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1 | Dummy)
```

```
## mod.log.Start.dummy.code: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1 | Dummy) + (1 | Code)
```

```
##               npar      AIC      BIC  logLik deviance  Chisq Df
## mod.log.Start.null          12 1692.3 1741.7 -834.16   1668.3
## mod.log.Start.dummy.code    13 1693.4 1746.9 -833.69   1667.4 0.936  1
##               Pr(>Chisq)
```

```
## mod.log.Start.null
```

```
## mod.log.Start.dummy.code    0.3333
```

If a test has a significant p-value (less than 0.05) then the random effect is significant. Site identity (Site) is significant by itself but turtle identity (Code) is not. I will see if Site and Code together are more significant than Site by itself.

### *## Site identity without the dummy variable*

```
mod.log.Start.site <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1|Site), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

### *## Turtle and site identity without the dummy variable*

```
mod.log.Start.code.site <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1|Code) + (1|Site), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

### *#anova with site model and code + site model*

```
anova(mod.log.Start.site, mod.log.Start.code.site)
```

```
## Data: MixedData
```

```
## Models:
```

```
## mod.log.Start.site: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1 | Site)
```

```
## mod.log.Start.code.site: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1 | Code) + (1 | Site)
```

```
##               npar      AIC      BIC  logLik deviance  Chisq Df
## mod.log.Start.site      12 1684.3 1733.7 -830.15   1660.3
## mod.log.Start.code.site  13 1685.5 1739.0 -829.73   1659.5 0.8478  1
##               Pr(>Chisq)
```

```
## mod.log.Start.site
```

```
## mod.log.Start.code.site      0.3572
```

Site and Code together are not more significant than Site by itself ( $p > 0.05$ ), so I will only keep Site.

## Predictor variables

I am selecting the final model with a backward selection procedure. At each step, I deleted the fixed effect with the highest p value. I confirmed the deletion of each fixed effect with a likelihood ratio test. I created a new dataset at each step to use only the rows with complete observations for all the fixed effects, so that the likelihood ratio tests do not run between two models with a different number of observations.

```
mod.log.Start.full <- lmer(Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
summary(mod.log.Start.full)
```

```

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled +
## Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled +
## water.100.Scaled + (1 | Site)
## Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
## 5619.1   5668.6  -2797.6   5595.1     442
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.7757 -0.6533 -0.3065  0.4705  4.1817
##
## Random effects:
## Groups   Name      Variance Std.Dev.
## Site     (Intercept) 1352     36.77
## Residual                12481    111.72
## Number of obs: 454, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    121.087    12.282  41.750   9.859 1.83e-12 ***
## JulianDate.Scaled -22.644    11.241  27.449  -2.014  0.05385 .
## A.Temp.Scaled    -7.177     9.167  97.611  -0.783  0.43557
## Time.Scaled     -4.954     5.852 451.163  -0.847  0.39772
## PL.Scaled       18.912     6.828 451.897   2.770  0.00584 **
## SexM             4.566    12.721 448.796   0.359  0.71980
## wet.400.Scaled  26.169    10.552  18.449   2.480  0.02299 *
## agri.400.Scaled  5.182     8.308  25.656   0.624  0.53828
## urban.1000.Scaled -3.624     9.976  19.570  -0.363  0.72028
## water.100.Scaled -10.805    11.088  18.579  -0.974  0.34233
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnDt.S A.Tm.S Tm.Scl PL.Scl SexM  w.400. a.400. u.1000
## JulnDt.Scl -0.069
## A.Temp.Scl  0.113 -0.412
## Time.Scl    0.050  0.085 -0.237
## PL.Scl      -0.312 -0.044 -0.054 -0.030
## SexM        -0.602  0.041 -0.107 -0.058  0.474
## wt.400.Scl -0.051 -0.256  0.130 -0.015  0.048  0.005
## agr.400.Scl -0.092  0.065 -0.007  0.046 -0.050 -0.038  0.161
## urbn.1000.S -0.084 -0.170  0.122  0.030  0.022  0.050  0.327 -0.019
## wtr.100.Scl  0.035  0.173  0.132 -0.083 -0.059 -0.052  0.234  0.081  0.003

```

I deleted proportion of urban area at 1000m (urban.1000.Scaled).

```

mod.log.Start.1 <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.
Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)

MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Scaled, MixedData$Sex, MixedData$wet.400.Scaled, MixedData$agri.400.Scaled, MixedData$urban.1000.Scaled, MixedData$water.100.Scaled),]

mod.full.adjust <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

mod.log.Start.1.adjust <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

anova(mod.log.Start.1.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.log.Start.1.adjust: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled + (1 | Site)
## mod.full.adjust: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + urban.1000.Scaled + water.100.Scaled + (1 | Site)
##
##               npar      AIC      BIC  logLik deviance  Chisq Df Pr(>Chi
##               sq)
## mod.log.Start.1.adjust    11 1683.3 1728.7 -830.68   1661.3
## mod.full.adjust          12 1684.3 1733.7 -830.15   1660.3 1.0476  1    0
##               .3061

summary(mod.log.Start.1)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
##          PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled
##          +
##          (1 | Site)
## Data: MixedData
##
##          AIC      BIC  logLik deviance df.resid
##    1683.4    1728.7  -830.7   1661.4      443
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max

```

```
## -3.2853 -0.4431 0.1587 0.6841 2.3076
##
## Random effects:
## Groups Name Variance Std.Dev.
## Site (Intercept) 0.1972 0.444
## Residual 2.1656 1.472
## Number of obs: 454, groups: Site, 23
##
## Fixed effects:
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) 4.17950 0.15545 46.17081 26.887 <2e-16 ***
## JulianDate.Scaled -0.20921 0.13942 29.10646 -1.501 0.1442
## A.Temp.Scaled -0.21045 0.11700 90.93424 -1.799 0.0754 .
## Time.Scaled 0.06512 0.07675 447.58010 0.848 0.3966
## PL.Scaled 0.21406 0.08973 453.01909 2.386 0.0175 *
## SexM -0.25630 0.16706 451.59760 -1.534 0.1257
## wet.400.Scaled 0.26264 0.12429 19.58936 2.113 0.0476 *
## agri.400.Scaled 0.15050 0.10438 27.45810 1.442 0.1607
## water.100.Scaled -0.27459 0.13814 19.21437 -1.988 0.0613 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
## (Intr) JlnD.S A.Tm.S Tm.Scl PL.Scl SexM w.400. a.400.
## JulnDt.Scl -0.087
## A.Temp.Scl 0.125 -0.410
## Time.Scaled 0.054 0.093 -0.242
## PL.Scaled -0.322 -0.042 -0.061 -0.030
## SexM -0.622 0.051 -0.113 -0.062 0.475
## wt.400.Scl -0.023 -0.216 0.099 -0.026 0.044 -0.011
## agr.400.Scl -0.084 0.060 -0.007 0.050 -0.053 -0.038 0.177
## wtr.100.Scl 0.037 0.184 0.134 -0.087 -0.063 -0.054 0.242 0.080
```

I deleted time of testing (Time).

```
mod.log.Start.2 <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled + (1|Site),
data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Scaled, MixedData$Sex, MixedData$wet.400.Scaled, MixedData$agri.400.Scaled, MixedData$water.100.Scaled),]
```

```
mod.full.adjust <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.log.Start.2.adjust <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled + (1|
```

```

Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

anova(mod.log.Start.2.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.log.Start.2.adjust: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL
.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled + (1 | Si
te)
## mod.full.adjust: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scal
ed + PL.Scaled + Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled +
(1 | Site)
##
##          npar    AIC    BIC  logLik deviance  Chisq Df Pr(>C
hisq)
## mod.log.Start.2.adjust    10 1682.1 1723.2 -831.03    1662.1
## mod.full.adjust          11 1683.3 1728.7 -830.68    1661.3 0.7111  1    0
.3991

summary(mod.log.Start.2)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + Sex +
##      wet.400.Scaled + agri.400.Scaled + water.100.Scaled + (1 |      Site)
##      Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
## 1682.1    1723.2   -831.0    1662.1      444
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.2866 -0.4403  0.1559  0.7085  2.2337
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
##  Site      (Intercept)  0.1884     0.434
##  Residual                    2.1721     1.474
## Number of obs: 454, groups:  Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    4.16989    0.15395 46.48752  27.085  <2e-16 ***
## JulianDate.Scaled -0.21898    0.13733 28.94543  -1.594    0.1217
## A.Temp.Scaled   -0.18946    0.11291 87.59331  -1.678    0.0969 .
## PL.Scaled        0.21769    0.08978 453.13083   2.425    0.0157 *
## SexM             -0.24670    0.16689 452.86811  -1.478    0.1400
## wet.400.Scaled   0.26570    0.12260 19.56986   2.167    0.0427 *
## agri.400.Scaled  0.14678    0.10310 27.38158   1.424    0.1658
## water.100.Scaled -0.26598    0.13575 18.91447  -1.959    0.0650 .
## ---

```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) JlnD.S A.Tm.S PL.Scl SexM   w.400. a.400.
## JulnDt.Scld -0.093
## A.Temp.Scld  0.143 -0.404
## PL.Scaled   -0.324 -0.039 -0.072
## SexM        -0.626  0.058 -0.132  0.475
## wt.400.Scld -0.022 -0.215  0.096  0.044 -0.013
## agr.400.Scl -0.084  0.055  0.006 -0.052 -0.036  0.178
## wtr.100.Scl  0.042  0.196  0.118 -0.067 -0.061  0.239  0.085
```

I deleted proportion of agricultural area at 400m (agri.400).

```
mod.log.Start.3 <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Sc
aled + Sex + wet.400.Scaled + water.100.Scaled + (1|Site), data = MixedData,
na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$Julia
nDate.Scaled, MixedData$A.Temp.Scaled, MixedData$PL.Scaled, MixedData$Sex, Mi
xedData$wet.400.Scaled, MixedData$agri.400.Scaled, MixedData$water.100.Scaled
),]
```

```
mod.full.adjust <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Sc
aled + Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled + (1|Site),
data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.log.Start.3.adjust <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled
+ PL.Scaled + Sex + wet.400.Scaled + water.100.Scaled + (1|Site), data = Mixe
dData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.log.Start.3.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.log.Start.3.adjust: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL
.Scaled + Sex + wet.400.Scaled + water.100.Scaled + (1 | Site)
```

```
## mod.full.adjust: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled
+ Sex + wet.400.Scaled + agri.400.Scaled + water.100.Scaled + (1 | Site)
```

```
##               npar      AIC      BIC    logLik deviance  Chisq Df Pr(>C
hisq)
```

```
## mod.log.Start.3.adjust      9 1682.0 1719.0 -831.98   1664.0
```

```
## mod.full.adjust           10 1682.1 1723.2 -831.03   1662.1 1.8953   1      0
.1686
```

```
summary(mod.log.Start.3)
```

```
## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
```

```
## Formula: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + Sex +
##          wet.400.Scaled + water.100.Scaled + (1 | Site)
```

```
## Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
##    1682    1719    -832    1664     445
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.2463 -0.4611  0.1448  0.7046  2.2515
##
## Random effects:
## Groups Name Variance Std.Dev.
## Site (Intercept) 0.225 0.4744
## Residual 2.169 1.4728
## Number of obs: 454, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    4.1988    0.1589  48.8803  26.424 <2e-16 ***
## JulianDate.Scaled -0.2345    0.1435  30.5100  -1.634  0.1124
## A.Temp.Scaled    -0.1793    0.1157 105.6708  -1.550  0.1242
## PL.Scaled        0.2187    0.0898 452.9378   2.436  0.0152 *
## SexM             -0.2419    0.1670 451.9804  -1.448  0.1482
## wet.400.Scaled    0.2337    0.1275  20.8536   1.833  0.0811 .
## water.100.Scaled -0.2766    0.1431  20.3342  -1.933  0.0673 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnDt.S A.Tm.S PL.Scl SexM   w.400.
## JulnDt.Scl -0.086
## A.Temp.Scl 0.143 -0.395
## PL.Scaled -0.320 -0.036 -0.067
## SexM      -0.611 0.058 -0.133 0.473
## wt.400.Scl -0.006 -0.228 0.094 0.052 -0.007
## wtr.100.Scl 0.049 0.183 0.116 -0.059 -0.055 0.233
```

I deleted turtle sex (Sex).

```
mod.log.Start.4 <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + wet.400.Scaled + water.100.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$PL.Scaled, MixedData$Sex, MixedData$wet.400.Scaled, MixedData$water.100.Scaled),]
```

```
mod.full.adjust <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + Sex + wet.400.Scaled + water.100.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```



```

mod.log.Start.4.adjust <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled
+ PL.Scaled + wet.400.Scaled + water.100.Scaled + (1|Site), data = MixedData.
adjust, na.action=na.exclude, REML = FALSE)

anova(mod.log.Start.4.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.log.Start.4.adjust: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL
.Scaled + wet.400.Scaled + water.100.Scaled + (1 | Site)
## mod.full.adjust: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled
+ Sex + wet.400.Scaled + water.100.Scaled + (1 | Site)
##
##          npar  AIC   BIC  logLik deviance  Chisq Df Pr(>Chisq
)
## mod.log.Start.4.adjust      8 1682 1715 -833.02      1666
## mod.full.adjust            9 1682 1719 -831.98      1664 2.0844  1      0.148
8

summary(mod.log.Start.4)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula:
## log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + wet.400.Scaled
+
##   water.100.Scaled + (1 | Site)
##   Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
## 1698.4   1731.4  -841.2   1682.4      451
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.2243 -0.4728  0.1780  0.7005  2.2212
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
##  Site      (Intercept)  0.2194     0.4684
##  Residual                    2.1721     1.4738
## Number of obs: 459, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    4.05638    0.12454  20.75064  32.571 < 2e-16 ***
## JulianDate.Scaled -0.21411    0.14210  30.81161  -1.507  0.142046
## A.Temp.Scaled   -0.22010    0.11385 106.82315  -1.933  0.055859 .
## PL.Scaled        0.27503    0.07759 458.99096   3.544  0.000434 ***
## wet.400.Scaled   0.22597    0.12636  21.16093   1.788  0.088055 .
## water.100.Scaled -0.30304    0.14128  20.29591  -2.145  0.044235 *
## ---

```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S PL.Scl w.400.
## JulnDt.Scld -0.066
## A.Temp.Scld  0.079 -0.393
## PL.Scaled   -0.034 -0.067 -0.010
## wt.400.Scld -0.012 -0.228  0.092  0.058
## wtr.100.Scl  0.017  0.190  0.106 -0.038  0.231
```

I deleted Julian date of testing (JulianDate).

```
mod.log.Start.5 <- lmer(log.Start ~ A.Temp.Scaled + PL.Scaled + wet.400.Scaled + water.100.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$PL.Scaled, MixedData$wet.400.Scaled, MixedData$water.100.Scaled),]
```

```
mod.full.adjust <- lmer(log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + wet.400.Scaled + water.100.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.log.Start.5.adjust <- lmer(log.Start ~ A.Temp.Scaled + PL.Scaled + wet.400.Scaled + water.100.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.log.Start.5.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.log.Start.5.adjust: log.Start ~ A.Temp.Scaled + PL.Scaled + wet.400.Scaled + water.100.Scaled + (1 | Site)
```

```
## mod.full.adjust: log.Start ~ JulianDate.Scaled + A.Temp.Scaled + PL.Scaled + wet.400.Scaled + water.100.Scaled + (1 | Site)
```

```
##              npar      AIC      BIC  logLik deviance  Chisq Df Pr(>ChiSq)
```

```
## mod.log.Start.5.adjust      7 1698.6 1727.5 -842.31   1684.6
```

```
## mod.full.adjust      8 1698.4 1731.4 -841.19   1682.4 2.2357  1      0.1349
```

```
summary(mod.log.Start.5)
```

```
## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's method [lmerModLmerTest]
```

```
## Formula:
```

```
## log.Start ~ A.Temp.Scaled + PL.Scaled + wet.400.Scaled + water.100.Scaled +
```

```
##      (1 | Site)
```

```
##      Data: MixedData
```

```
##
##      AIC      BIC   logLik deviance df.resid
##  1698.6   1727.5   -842.3   1684.6     452
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.2385 -0.4510  0.1859  0.6909  2.1771
##
## Random effects:
##   Groups      Name      Variance Std.Dev.
##   Site      (Intercept) 0.2395   0.4894
##   Residual                2.1768   1.4754
## Number of obs: 459, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    4.04785    0.12794  19.43850   31.639 < 2e-16 ***
## A.Temp.Scaled  -0.28209    0.10626  77.82417   -2.655 0.009623 **
## PL.Scaled       0.26514    0.07761 458.99814    3.416 0.000692 ***
## wet.400.Scaled  0.18150    0.12664  19.69936    1.433 0.167499
## water.100.Scaled -0.26040    0.14297  17.99773   -1.821 0.085224 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) A.Tm.S PL.Scl w.400.
## A.Temp.Scld  0.059
## PL.Scaled   -0.038 -0.036
## wt.400.Scld -0.028  0.002  0.043
## wtr.100.Scl  0.031  0.196 -0.025  0.288
```

I deleted proportion of wetland area at 400m (wet.400).

```
mod.log.Start.6 <- lmer(log.Start ~ A.Temp.Scaled + PL.Scaled + water.100.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Start, MixedData$A.Temp.Scaled, MixedData$PL.Scaled, MixedData$wet.400.Scaled, MixedData$water.100.Scaled),]
```

```
mod.full.adjust <- lmer(log.Start ~ A.Temp.Scaled + PL.Scaled + wet.400.Scaled + water.100.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.log.Start.6.adjust <- lmer(log.Start ~ A.Temp.Scaled + PL.Scaled + water.100.Scaled + (1|Site), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.log.Start.6.adjust, mod.full.adjust)
```

```

## Data: MixedData.adjust
## Models:
## mod.log.Start.6.adjust: log.Start ~ A.Temp.Scaled + PL.Scaled + water.100.
Scaled + (1 | Site)
## mod.full.adjust: log.Start ~ A.Temp.Scaled + PL.Scaled + wet.400.Scaled +
water.100.Scaled + (1 | Site)
##               npar      AIC      BIC  logLik deviance  Chisq Df Pr(>C
hisq)
## mod.log.Start.6.adjust      6 1698.5 1723.3 -843.26   1686.5
## mod.full.adjust           7 1698.6 1727.5 -842.31   1684.6 1.9138  1    0
.1665

summary(mod.log.Start.6)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: log.Start ~ A.Temp.Scaled + PL.Scaled + water.100.Scaled + (1 |
## Site)
## Data: MixedData
##
##      AIC      BIC  logLik deviance df.resid
##  1698.5   1723.3  -843.3   1686.5      453
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.2482 -0.4496  0.1993  0.7078  2.2029
##
## Random effects:
## Groups   Name      Variance Std.Dev.
## Site     (Intercept) 0.2836   0.5325
## Residual                2.1736   1.4743
## Number of obs: 459, groups: Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    4.06011    0.13542  20.72958  29.981 < 2e-16 ***
## A.Temp.Scaled  -0.27120    0.10916  91.85078  -2.484  0.01479 *
## PL.Scaled       0.25683    0.07769  458.85146   3.306  0.00102 **
## water.100.Scaled -0.31479    0.14519  19.78048  -2.168  0.04253 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) A.Tm.S PL.Scl
## A.Temp.Scl d  0.060
## PL.Scaled     -0.035 -0.029
## wtr.100.Scl   0.042  0.195 -0.037

```

All of the fixed effects are statistically significant, so I will stop the backwards selection process.

## Final model

### Summary statistics

I changed the REML to TRUE to calculate the summary statistics of the final model.

```
mod.log.Start.final <- lmer(log.Start ~ A.Temp.Scaled + PL.Scaled + water.100
.Scaled + (1|Site), data = MixedData, na.action=na.exclude, REML = TRUE)

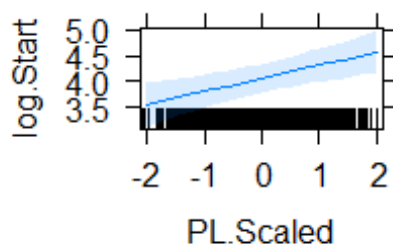
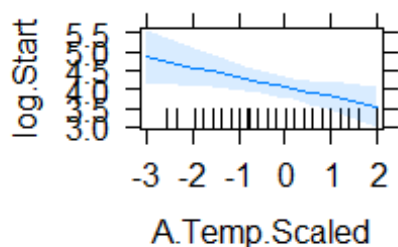
summary(mod.log.Start.final)

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: log.Start ~ A.Temp.Scaled + PL.Scaled + water.100.Scaled + (1 |
##      Site)
##      Data: MixedData
##
## REML criterion at convergence: 1696.5
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.2455 -0.4547  0.2023  0.7024  2.2234
##
## Random effects:
##      Groups      Name      Variance Std.Dev.
##      Site      (Intercept) 0.3389   0.5822
##      Residual                2.1801   1.4765
## Number of obs: 459, groups:  Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    4.06719    0.14435  19.04324  28.176 < 2e-16 ***
## A.Temp.Scaled  -0.25963    0.11241  95.45498  -2.310  0.02306 *
## PL.Scaled       0.25354    0.07802  454.88303   3.250  0.00124 **
## water.100.Scaled -0.31011    0.15498  18.31420  -2.001  0.06044 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) A.Tm.S PL.Scl
## A.Temp.Scld  0.061
## PL.Scaled   -0.034 -0.023
## wtr.100.Scl  0.043  0.186 -0.036
```

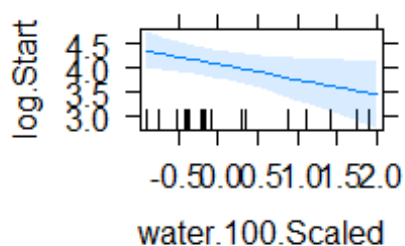
### Visualization of the predictor effects

```
plot(allEffects(mod.log.Start.final))
```

## A.Temp.Scaled effect plot    PL.Scaled effect plot



## water.100.Scaled effect plot



## Calculation of the marginal and conditional variance explained by the final model

```
r.squaredGLMM(mod.log.Start.final)
```

```
##           R2m           R2c
## [1,] 0.05926647 0.1858313
```

Marginal R2: fixed effects R2. Conditional R2: fixed and random effects R2.

## Calculation of the 95% confidence intervals

```
confint(mod.log.Start.final, level = 0.95, method = "Wald")
```

```
##           2.5 %           97.5 %
## .sig01           NA           NA
## .sigma           NA           NA
## (Intercept)    3.7842661    4.350113266
## A.Temp.Scaled  -0.4799473   -0.039307505
## PL.Scaled       0.1006257    0.406453293
## water.100.Scaled -0.6138571  -0.006360052
```

## Model selection

### Total time spent moving

#### Random variable

I am testing the significance of turtle ID and site identity by using likelihood ratio tests to see if the addition of these random variables make a significant effect on the initial model. I am using a dummy variable (same value for all the observations) to create a null mixed model to compared with the different combinations of mixed models.

#### Creation of the different mixed models

##### *#Null Mixed Model*

```
mod.Move.null <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Dummy), data = MixedData, na.action = na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

##### *## only site identity as random variable*

```
mod.Move.dummy.site <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Dummy) + (1|Site), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

##### *## only turtle ID as random variable*

```
mod.Move.dummy.code <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Dummy) + (1|Code), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)
```

## Likelihood ratio tests between the mixed models

##### *#anova with null model and dummy + site model*

```
anova(mod.Move.null, mod.Move.dummy.site)
```

```
## Data: MixedData
```

```
## Models:
```

```
## mod.Move.null: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Dummy)
```

```
## mod.Move.dummy.site: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Dummy) + (1 | Site)
```

```
##               npar      AIC      BIC  logLik deviance  Chisq Df Pr(>Chisq)
```

```
## mod.Move.null      13 4693.0 4746.6 -2333.5   4667.0
```

```
## mod.Move.dummy.site 14 4683.4 4741.0 -2327.7   4655.4 11.665  1 0.00063
```

```

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##
## mod.Move.null
## mod.Move.dummy.site ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

#anova with null model and dummy + code model
anova(mod.Move.null, mod.Move.dummy.code)

## Data: MixedData
## Models:
## mod.Move.null: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL
.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban
.1000.Scaled + water.1000.Scaled + (1 | Dummy)
## mod.Move.dummy.code: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Dummy) + (1 | Code)
##
      npar    AIC    BIC logLik deviance  Chisq Df Pr(>Chisq)
## mod.Move.null      13 4693.0 4746.6 -2333.5   4667.0
## mod.Move.dummy.code 14 4690.6 4748.2 -2331.3   4662.6 4.4534  1    0.034
83 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

If a test has a significant p-value (less than 0.05) then the random effect is significant. Site identity (Site) and turtle ID (Code) are significant by themselves. I will see if Site and Code together are more significant than Site and Code by themselves.

```

## Turtle ID without the dummy variable
mod.Move.code <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Code), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)

## Site identity without the dummy variable
mod.Move.site <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)

## Turtle and site identity without the dummy variable
mod.Move.code.site <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Code) + (1|Site), data = MixedData, na.action=na.exclude, control=lmerControl(check.nlev.gtr.1="ignore"), REML = FALSE)

#anova with site model and code + site model
anova(mod.Move.site, mod.Move.code.site)

```



```
## Data: MixedData
## Models:
## mod.Move.site: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL
.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban
.1000.Scaled + water.1000.Scaled + (1 | Site)
## mod.Move.code.site: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled
+ PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + u
rban.1000.Scaled + water.1000.Scaled + (1 | Code) + (1 | Site)
##
      npar    AIC    BIC  logLik deviance  Chisq Df Pr(>Chisq
)
## mod.Move.site          13 4681.4 4734.9 -2327.7   4655.4
## mod.Move.code.site     14 4679.5 4737.1 -2325.7   4651.5 3.9129  1    0.0479
2 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

#anova with code model and code + site model
anova(mod.Move.code, mod.Move.code.site)

## Data: MixedData
## Models:
## mod.Move.code: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL
.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban
.1000.Scaled + water.1000.Scaled + (1 | Code)
## mod.Move.code.site: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled
+ PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + u
rban.1000.Scaled + water.1000.Scaled + (1 | Code) + (1 | Site)
##
      npar    AIC    BIC  logLik deviance  Chisq Df Pr(>Chisq
)
## mod.Move.code          13 4688.6 4742.1 -2331.3   4662.6
## mod.Move.code.site     14 4679.5 4737.1 -2325.7   4651.5 11.125  1 0.000851
8 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Site and Code together are more significant than Site and Code by themselves ( $p < 0.05$ ), so I will keep both.

## Predictor variables

I am selecting the final model with a backward selection procedure. At each step, I deleted the fixed effect with the highest p value. I confirmed the deletion of each fixed effect with a likelihood ratio test. I created a new dataset at each step to use only the rows with complete observations for all the fixed effects, so that the likelihood ratio tests do not run between two models with a different number of observations.

```
mod.Move.full <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled
+ PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + u
rban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData,
```

```

na.action=na.exclude, REML = FALSE)
summary(mod.Move.full)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled +
## Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled +
## urban.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 | Code)
## Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
##  4679.5   4737.1  -2325.7   4651.5     439
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.94945 -0.53190  0.05023  0.55901  1.59557
##
## Random effects:
## Groups Name Variance Std.Dev.
## Code (Intercept) 746.7 27.33
## Site (Intercept) 123.4 11.11
## Residual 890.7 29.84
## Number of obs: 453, groups: Code, 429; Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)  90.03990   4.13573   51.89653  21.771 <2e-16 ***
## JulianDate.Scaled  2.53842   3.83232   27.68781   0.662  0.5132
## A.Temp.Scaled    -1.55475   3.13922   89.37380  -0.495  0.6216
## Time.Scaled       1.20480   2.08020  440.09897   0.579  0.5628
## PL.Scaled        -0.83400   2.49972  440.21016  -0.334  0.7388
## SexM              4.80496   4.54104  432.01412   1.058  0.2906
## for.veg.300.Scaled  0.03984   5.41914   22.81472   0.007  0.9942
## wet.600.Scaled    -0.96779   4.00018   25.31734  -0.242  0.8108
## agri.100.Scaled    7.28995   3.61269   32.42149   2.018  0.0519 .
## urban.1000.Scaled  5.87915   4.85745   23.95407   1.210  0.2380
## water.1000.Scaled -3.19655   3.83167   22.83416  -0.834  0.4128
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnDt.S A.Tm.S Tm.Scl PL.Scl SexM  f..300 w.600. a.100.
## JlnDt.Scl -0.077
## A.Temp.Scl 0.102 -0.439
## Time.Scl 0.050 0.115 -0.222
## PL.Scl -0.317 -0.021 -0.068 -0.037
## SexM -0.633 0.060 -0.105 -0.064 0.470
## fr.vg.300.S 0.086 -0.047 -0.138 -0.041 0.050 0.009
## wt.600.Scl -0.005 -0.332 -0.006 -0.023 0.074 0.013 0.517

```

```
## agr.100.Scl  0.044 -0.020 -0.062  0.008  0.012 -0.044  0.586  0.352
## urbn.1000.S -0.008 -0.195 -0.029 -0.015  0.039  0.048  0.695  0.597  0.312
## wtr.1000.Sc  0.017 -0.259 -0.033 -0.041 -0.034 -0.021  0.567  0.494  0.416
##              u.1000
## JulnDt.Scl d
## A.Temp.Scl d
## Time.Scaled
## PL.Scaled
## SexM
## fr.vg.300.S
## wt.600.Scl d
## agr.100.Scl
## urbn.1000.S
## wtr.1000.Sc  0.563
```

I deleted proportion of forest and vegetation area at 300m (for.veg.300).

```
mod.Move.1 <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + P
L.Scaled + Sex + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water
.1000.Scaled + (1|Site) + (1|Code), data = MixedData, na.action=na.exclude, R
EML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Move, MixedData$Julian
Date.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Sca
led, MixedData$Sex, MixedData$for.veg.300.Scaled, MixedData$wet.600.Scaled,
MixedData$agri.100.Scaled, MixedData$urban.1000.Scaled, MixedData$water.1000.
Scaled),]
```

```
mod.full.adjust <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.Move.1.adjust <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.Move.1.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Move.1.adjust: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 | Code)
```

```
## mod.full.adjust: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + for.veg.300.Scaled + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 | Code)
```

```
##              npar      AIC      BIC  logLik deviance Chisq Df Pr(>Chisq)
```

```

## mod.Move.1.adjust    13 4677.5 4731.0 -2325.7    4651.5
## mod.full.adjust     14 4679.5 4737.1 -2325.7    4651.5 1e-04  1      0.9941

summary(mod.Move.1)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled +
## Sex + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled +
## water.1000.Scaled + (1 | Site) + (1 | Code)
## Data: MixedData
##
##      AIC      BIC    logLik deviance df.resid
##  4677.5   4731.0   -2325.7   4651.5     440
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.94939 -0.53174  0.05024  0.55910  1.59533
##
## Random effects:
## Groups Name Variance Std.Dev.
## Code (Intercept) 746.7 27.33
## Site (Intercept) 123.4 11.11
## Residual 890.8 29.85
## Number of obs: 453, groups: Code, 429; Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    90.037     4.120  53.830   21.853 <2e-16 ***
## JulianDate.Scaled  2.540     3.828  27.309    0.663  0.5126
## A.Temp.Scaled   -1.552     3.109  87.320   -0.499  0.6190
## Time.Scaled     1.205     2.079 437.021    0.580  0.5622
## PL.Scaled       -0.835     2.497 440.696   -0.334  0.7382
## SexM            4.805     4.541 432.046    1.058  0.2906
## wet.600.Scaled  -0.983     3.425  21.684   -0.287  0.7768
## agri.100.Scaled  7.274     2.928  35.408    2.484  0.0179 *
## urban.1000.Scaled 5.854     3.493  20.559    1.676  0.1088
## water.1000.Scaled -3.212     3.157  20.092   -1.018  0.3210
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnDt.S A.Tm.S Tm.Scl PL.Scl SexM  w.600. a.100. u.1000
## JulnDt.Scl -0.073
## A.Temp.Scl  0.115 -0.451
## Time.Scl    0.054  0.114 -0.230
## PL.Scl      -0.323 -0.019 -0.061 -0.035
## SexM        -0.637  0.060 -0.105 -0.063  0.470
## wt.600.Scl -0.058 -0.360  0.077 -0.003  0.057  0.009

```

```
## agr.100.Scl -0.007  0.009  0.023  0.040 -0.021 -0.060  0.072
## urbn.1000.S -0.095 -0.226  0.094  0.019  0.007  0.058  0.386 -0.163
## wtr.1000.Sc -0.039 -0.283  0.056 -0.021 -0.075 -0.032  0.286  0.126  0.286
```

I deleted proportion of wetland area at 600m (wet.600).

```
mod.Move.2 <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + P
L.Scaled + Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1
|Site) + (1|Code), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Move, MixedData$Julian
Date.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Sca
led, MixedData$Sex, MixedData$wet.600.Scaled, MixedData$agri.100.Scaled, Mix
edData$urban.1000.Scaled, MixedData$water.1000.Scaled),]
```

```
mod.full.adjust <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.Move.2.adjust <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.Move.2.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Move.2.adjust: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 | Code)
```

```
## mod.full.adjust: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + wet.600.Scaled + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 | Code)
```

```
##          npar      AIC      BIC  logLik deviance  Chisq Df Pr(>Chisq)
## mod.Move.2.adjust    12 4675.6 4724.9 -2325.8   4651.6
## mod.full.adjust     13 4677.5 4731.0 -2325.7   4651.5 0.0821  1      0.7745
```

```
summary(mod.Move.2)
```

```
## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
```

```
## Formula: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled +
```

```
## Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled +
## (1 | Site) + (1 | Code)
```

```
## Data: MixedData
```

```
##
```

```
##      AIC      BIC  logLik deviance df.resid
## 4675.6  4724.9 -2325.8   4651.6      441
```

```
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.94827 -0.53101  0.05176  0.56139  1.59715
##
## Random effects:
##   Groups   Name      Variance Std.Dev.
##   Code     (Intercept) 746.2    27.32
##   Site     (Intercept) 124.6    11.16
##   Residual                891.0    29.85
## Number of obs: 453, groups:  Code, 429; Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    89.9635     4.1200   54.5571   21.836   <2e-16 ***
## JulianDate.Scaled  2.1491     3.5788   28.6360    0.601   0.5529
## A.Temp.Scaled    -1.4847     3.1037   89.0762   -0.478   0.6336
## Time.Scaled      1.2043     2.0789  437.4471    0.579   0.5627
## PL.Scaled        -0.7914     2.4928  440.8079   -0.317   0.7510
## SexM             4.8188     4.5409  432.0419    1.061   0.2892
## agri.100.Scaled   7.3381     2.9269   35.3741    2.507   0.0169 *
## urban.1000.Scaled 6.2406     3.2300   21.9675    1.932   0.0663 .
## water.1000.Scaled -2.9556     3.0334   20.9853   -0.974   0.3410
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S Tm.Scl PL.Scl SexM   a.100. u.1000
## JulnDt.SclD -0.101
## A.Temp.SclD  0.120 -0.454
## Time.Scaled  0.054  0.120 -0.231
## PL.Scaled    -0.321  0.002 -0.066 -0.035
## SexM         -0.636  0.068 -0.106 -0.063  0.470
## agr.100.Scl -0.004  0.038  0.018  0.040 -0.025 -0.061
## urbn.1000.S -0.079 -0.102  0.070  0.022 -0.016  0.059 -0.207
## wtr.1000.Sc -0.024 -0.201  0.035 -0.022 -0.095 -0.036  0.110  0.199
```

I deleted turtle plastron length (PL).

```
mod.Move.3 <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Move, MixedData$JulianDate.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$PL.Scaled, MixedData$Sex, MixedData$agri.100.Scaled, MixedData$urban.1000.Scaled, MixedData$water.1000.Scaled),]
```

```
mod.full.adjust <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + PL.Scaled + Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled
```

```

+ (1|Site) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML =
FALSE)

mod.Move.3.adjust <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Sca
led + Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site
) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

anova(mod.Move.3.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.Move.3.adjust: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled
+ Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Site)
+ (1 | Code)
## mod.full.adjust: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
PL.Scaled + Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (
1 | Site) + (1 | Code)
##
##          npar    AIC    BIC logLik deviance Chisq Df Pr(>Chisq)
## mod.Move.3.adjust    11 4673.7 4718.9 -2325.8   4651.7
## mod.full.adjust     12 4675.6 4724.9 -2325.8   4651.6 0.1004  1      0.7514

summary(mod.Move.3)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + Sex +
##          agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled +
##          (1 | Site) + (1 | Code)
## Data: MixedData
##
##          AIC          BIC    logLik deviance df.resid
##    4683.9    4729.2   -2331.0   4661.9      443
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.93328 -0.53864  0.05394  0.55345  1.59614
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
##  Code     (Intercept)    746.7      27.33
##  Site     (Intercept)    128.6      11.34
##  Residual                    889.6      29.83
## Number of obs: 454, groups: Code, 430; Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    89.4599     3.9264  44.4263  22.784  <2e-16 ***
## JulianDate.Scaled  2.0268     3.6044  28.5476   0.562   0.5783
## A.Temp.Scaled    -1.6561     3.1085  88.8618  -0.533   0.5955
## Time.Scaled       0.8925     2.0586  438.5359   0.434   0.6648

```

```
## SexM          5.4202      4.0072 443.6209    1.353    0.1769
## agri.100.Scaled  7.0283      2.9319  33.9761    2.397    0.0222 *
## urban.1000.Scaled 6.2252      3.2590  21.9516    1.910    0.0693 .
## water.1000.Scaled -2.9995      3.0476  20.5500   -0.984    0.3364
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S Tm.Scl SexM   a.100. u.1000
## JulnDt.Scl d -0.106
## A.Temp.Scl d  0.104 -0.455
## Time.Scaled  0.043  0.116 -0.240
## SexM          -0.578  0.075 -0.087 -0.055
## agr.100.Scl  -0.016  0.035  0.013  0.025 -0.058
## urbn.1000.S -0.089 -0.103  0.071  0.022  0.075 -0.209
## wtr.1000.Sc -0.058 -0.202  0.030 -0.023  0.011  0.111  0.199
```

I deleted time of testing (Time).

```
mod.Move.4 <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Sex + agri.100.
Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data =
MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Move, MixedData$Julian
Date.Scaled, MixedData$A.Temp.Scaled, MixedData$Time.Scaled, MixedData$Sex, M
ixedData$agri.100.Scaled, MixedData$urban.1000.Scaled, MixedData$water.1000.S
caled),]
```

```
mod.full.adjust <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scale
d + Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site)
+ (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.Move.4.adjust <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Sex + ag
ri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code),
data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.Move.4.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Move.4.adjust: Move ~ JulianDate.Scaled + A.Temp.Scaled + Sex + agri.1
00.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 | Code)
```

```
## mod.full.adjust: Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled +
Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Site) +
(1 | Code)
```

```
##              npar    AIC    BIC logLik deviance Chisq Df Pr(>Chisq)
## mod.Move.4.adjust   10 4682.1 4723.3 -2331.1   4662.1
## mod.full.adjust     11 4683.9 4729.2 -2331.0   4661.9 0.1877  1      0.6648
```

```
summary(mod.Move.4)
```



```

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Move ~ JulianDate.Scaled + A.Temp.Scaled + Sex + agri.100.Scaled
+
##      urban.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 |      Code)
##      Data: MixedData
##
##      AIC      BIC    logLik deviance df.resid
##    4682.1    4723.3   -2331.1    4662.1      444
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.95059 -0.53848  0.05562  0.55242  1.60079
##
## Random effects:
##  Groups      Name              Variance Std.Dev.
##  Code      (Intercept)  750.9      27.40
##  Site      (Intercept)  128.3      11.33
##  Residual                    886.4      29.77
## Number of obs: 454, groups:  Code, 430; Site, 23
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)      89.388      3.921   44.111   22.796 <2e-16 ***
## JulianDate.Scaled    1.846      3.578   27.817    0.516  0.6100
## A.Temp.Scaled      -1.333      3.017   88.263   -0.442  0.6597
## SexM                5.516      4.002  444.720    1.378  0.1688
## agri.100.Scaled     6.995      2.930   33.803    2.388  0.0227 *
## urban.1000.Scaled   6.195      3.256   21.849    1.903  0.0703 .
## water.1000.Scaled  -2.969      3.045   20.445   -0.975  0.3409
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S A.Tm.S SexM    a.100. u.1000
## JulnDt.Scld -0.112
## A.Temp.Scld  0.118 -0.443
## SexM         -0.577  0.082 -0.103
## agr.100.Scld -0.017  0.032  0.020 -0.057
## urbn.1000.S -0.090 -0.106  0.078  0.076 -0.210
## wtr.1000.Sc -0.057 -0.201  0.025  0.009  0.111  0.199

```

I delted air temperature (A.Temp).

```

mod.Move.5 <- lmer(Move ~ JulianDate.Scaled + Sex + agri.100.Scaled + urban.1
000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData, na.ac
tion=na.exclude, REML = FALSE)

```

```

MixedData.adjust <- MixedData[complete.cases(MixedData$Move, MixedData$Julian
Date.Scaled, MixedData$A.Temp.Scaled, MixedData$Sex, MixedData$agri.100.Scale

```

```
d, MixedData$urban.1000.Scaled, MixedData$water.1000.Scaled),]

mod.full.adjust <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Sex + agri
.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), da
ta = MixedData.adjust, na.action=na.exclude, REML = FALSE)

mod.Move.5.adjust <- lmer(Move ~ JulianDate.Scaled + Sex + agri.100.Scaled +
urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData
.adjust, na.action=na.exclude, REML = FALSE)

anova(mod.Move.5.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.Move.5.adjust: Move ~ JulianDate.Scaled + Sex + agri.100.Scaled + urba
n.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 | Code)
## mod.full.adjust: Move ~ JulianDate.Scaled + A.Temp.Scaled + Sex + agri.100
.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 | Code)
##
##          npar      AIC      BIC  logLik deviance Chisq Df Pr(>Chisq)
## mod.Move.5.adjust      9 4680.3 4717.4 -2331.2   4662.3
## mod.full.adjust      10 4682.1 4723.3 -2331.1   4662.1 0.195  1      0.6588

summary(mod.Move.5)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula:
## Move ~ JulianDate.Scaled + Sex + agri.100.Scaled + urban.1000.Scaled +
## water.1000.Scaled + (1 | Site) + (1 | Code)
## Data: MixedData
##
##      AIC      BIC  logLik deviance df.resid
##  4680.3   4717.4 -2331.2   4662.3      445
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.93033 -0.52757  0.06357  0.55061  1.60079
##
## Random effects:
## Groups   Name      Variance Std.Dev.
## Code     (Intercept) 747.8    27.35
## Site     (Intercept) 128.7    11.35
## Residual                    889.8    29.83
## Number of obs: 454, groups: Code, 430; Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)      89.591      3.897  43.305   22.989   <2e-16 ***
## JulianDate.Scaled    1.146      3.211  21.509    0.357   0.7246
## SexM                5.334      3.981  445.429    1.340   0.1810
```

```
## agri.100.Scaled      7.022      2.932  33.221   2.395   0.0224 *
## urban.1000.Scaled    6.307      3.250  21.924   1.941   0.0652 .
## water.1000.Scaled   -2.936      3.047  20.133  -0.963   0.3468
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) JlnD.S SexM   a.100. u.1000
## JulnDt.Scld -0.067
## SexM         -0.571  0.041
## agr.100.Scl -0.019  0.046 -0.055
## urbn.1000.S -0.100 -0.080  0.085 -0.212
## wtr.1000.Sc -0.061 -0.212  0.012  0.111  0.198
```

I deleted Julian date of testing (JulianDate).

```
mod.Move.6 <- lmer(Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Move, MixedData$JulianDate.Scaled, MixedData$Sex, MixedData$agri.100.Scaled, MixedData$urban.1000.Scaled, MixedData$water.1000.Scaled),]
```

```
mod.full.adjust <- lmer(Move ~ JulianDate.Scaled + Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
mod.Move.6.adjust <- lmer(Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)
```

```
anova(mod.Move.6.adjust, mod.full.adjust)
```

```
## Data: MixedData.adjust
```

```
## Models:
```

```
## mod.Move.6.adjust: Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 | Code)
```

```
## mod.full.adjust: Move ~ JulianDate.Scaled + Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled + (1 | Site) + (1 | Code)
```

```
##              npar      AIC      BIC logLik deviance Chisq Df Pr(>Chisq)
## mod.Move.6.adjust      8 4678.5 4711.4 -2331.2   4662.5
## mod.full.adjust       9 4680.3 4717.4 -2331.2   4662.3 0.1271  1      0.7214
```

```
summary(mod.Move.6)
```

```
## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
```

```
## Formula:
```

```
## Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + water.1000.Scaled +
## (1 | Site) + (1 | Code)
```

```
## Data: MixedData
##
##      AIC      BIC   logLik deviance df.resid
## 4678.5  4711.4 -2331.2  4662.5      446
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.93595 -0.53469  0.05744  0.55361  1.59762
##
## Random effects:
## Groups   Name      Variance Std.Dev.
## Code     (Intercept) 745.4    27.30
## Site     (Intercept) 127.8    11.31
## Residual                892.9    29.88
## Number of obs: 454, groups: Code, 430; Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)      89.687      3.883  42.667   23.095 <2e-16 ***
## SexM              5.277      3.979  445.904    1.326  0.1854
## agri.100.Scaled   6.971      2.924   33.643    2.384  0.0229 *
## urban.1000.Scaled 6.400      3.233   22.157    1.979  0.0603 .
## water.1000.Scaled -2.705      2.972   20.588   -0.910  0.3732
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) SexM   a.100. u.1000
## SexM          -0.571
## agr.100.Scl   -0.016 -0.057
## urbn.1000.S   -0.106  0.088 -0.209
## wtr.1000.Sc   -0.076  0.021  0.123  0.186
```

I deleted proportion of open water area at 1000m (water.1000).

```
mod.Move.7 <- lmer(Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + (1|Site)
) + (1|Code), data = MixedData, na.action=na.exclude, REML = FALSE)
```

```
MixedData.adjust <- MixedData[complete.cases(MixedData$Move, MixedData$Sex, M
ixedData$agri.100.Scaled, MixedData$urban.1000.Scaled, MixedData$water.1000.S
caled),]
```

```
mod.full.adjust <- lmer(Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + wa
ter.1000.Scaled + (1|Site) + (1|Code), data = MixedData.adjust, na.action=na.
exclude, REML = FALSE)
```

```
mod.Move.7.adjust <- lmer(Move ~ Sex + agri.100.Scaled + urban.1000.Scaled +
(1|Site) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FA
LSE)
```

```

anova(mod.Move.7.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.Move.7.adjust: Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + (1 |
Site) + (1 | Code)
## mod.full.adjust: Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + water.
1000.Scaled + (1 | Site) + (1 | Code)
##
##          npar      AIC      BIC  logLik deviance  Chisq Df Pr(>Chisq)
## mod.Move.7.adjust      7 4677.3 4706.1 -2331.6   4663.3
## mod.full.adjust      8 4678.5 4711.4 -2331.2   4662.5 0.8139  1      0.367

summary(mod.Move.7)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + (1 | Site) +
## (1 | Code)
## Data: MixedData
##
##      AIC      BIC  logLik deviance df.resid
##  4677.3  4706.1 -2331.6   4663.3      447
##
## Scaled residuals:
##      Min      1Q   Median      3Q      Max
## -1.94250 -0.53395  0.06378  0.54919  1.60909
##
## Random effects:
## Groups   Name      Variance Std.Dev.
## Code     (Intercept) 746.7    27.33
## Site     (Intercept) 136.1    11.67
## Residual                891.7    29.86
## Number of obs: 454, groups: Code, 430; Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)      89.393      3.922 42.669  22.792 <2e-16 ***
## SexM              5.346      3.981 445.692   1.343  0.1800
## agri.100.Scaled   7.323      2.944  33.752   2.487  0.0180 *
## urban.1000.Scaled 6.948      3.234  22.414   2.148  0.0427 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) SexM  a.100.
## SexM          -0.565
## agr.100.Scl -0.010 -0.059
## urbn.1000.S -0.093  0.085 -0.238

```

I deleted turtle sex (Sex).

```

mod.Move.8 <- lmer(Move ~ agri.100.Scaled + urban.1000.Scaled + (1|Site) + (1
|Code), data = MixedData, na.action=na.exclude, REML = FALSE)

MixedData.adjust <- MixedData[complete.cases(MixedData$Move, MixedData$Sex, M
ixedData$agri.100.Scaled, MixedData$urban.1000.Scaled),]

mod.full.adjust <- lmer(Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + (1
|Site) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALS
E)

mod.Move.8.adjust <- lmer(Move ~ agri.100.Scaled + urban.1000.Scaled + (1|Sit
e) + (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

anova(mod.Move.8.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.Move.8.adjust: Move ~ agri.100.Scaled + urban.1000.Scaled + (1 | Site)
+ (1 | Code)
## mod.full.adjust: Move ~ Sex + agri.100.Scaled + urban.1000.Scaled + (1 | S
ite) + (1 | Code)
##
##          npar    AIC    BIC logLik deviance Chisq Df Pr(>Chisq)
## mod.Move.8.adjust      6 4677.1 4701.8 -2332.5    4665.1
## mod.full.adjust       7 4677.3 4706.1 -2331.6    4663.3 1.7987  1      0.1799

summary(mod.Move.8)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Move ~ agri.100.Scaled + urban.1000.Scaled + (1 | Site) + (1 |
## Code)
## Data: MixedData
##
##          AIC          BIC    logLik deviance df.resid
##  4728.9    4753.6   -2358.4    4716.9        453
##
## Scaled residuals:
##      Min         1Q     Median        3Q        Max
## -1.94979 -0.51286  0.06056  0.52536  1.60654
##
## Random effects:
## Groups   Name                Variance Std.Dev.
## Code    (Intercept)    793.9      28.18
## Site    (Intercept)   142.0      11.92
## Residual                    854.7      29.24
## Number of obs: 459, groups: Code, 434; Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    92.524      3.279 20.374  28.216  <2e-16 ***

```

```
## agri.100.Scaled      7.494      2.974 33.220   2.520   0.0167 *
## urban.1000.Scaled   6.698      3.263 22.036   2.053   0.0522 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) a.100.
## agr.100.Scl -0.053
## urbn.1000.S -0.054 -0.235
```

I deleted proportion of urban area at 1000m (urban.1000).

```
mod.Move.9 <- lmer(Move ~ agri.100.Scaled + (1|Site) + (1|Code), data = Mixed
Data, na.action=na.exclude, REML = FALSE)

MixedData.adjust <- MixedData[complete.cases(MixedData$Move, MixedData$agri.1
00.Scaled, MixedData$urban.1000.Scaled),]

mod.full.adjust <- lmer(Move ~ agri.100.Scaled + urban.1000.Scaled + (1|Site)
+ (1|Code), data = MixedData.adjust, na.action=na.exclude, REML = FALSE)

mod.Move.9.adjust <- lmer(Move ~ agri.100.Scaled + (1|Site) + (1|Code), data
= MixedData.adjust, na.action=na.exclude, REML = FALSE)

anova(mod.Move.9.adjust, mod.full.adjust)

## Data: MixedData.adjust
## Models:
## mod.Move.9.adjust: Move ~ agri.100.Scaled + (1 | Site) + (1 | Code)
## mod.full.adjust: Move ~ agri.100.Scaled + urban.1000.Scaled + (1 | Site) +
(1 | Code)
##              npar      AIC      BIC logLik deviance Chisq Df Pr(>Chisq)
## mod.Move.9.adjust      5 4730.7 4751.4 -2360.3   4720.7
## mod.full.adjust       6 4728.9 4753.6 -2358.4   4716.9 3.8449  1    0.0499
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

summary(mod.Move.9)

## Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
## method [lmerModLmerTest]
## Formula: Move ~ agri.100.Scaled + (1 | Site) + (1 | Code)
## Data: MixedData
##
##      AIC      BIC  logLik deviance df.resid
##  4730.7  4751.4 -2360.4   4720.7      454
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.95044 -0.52349  0.05835  0.53308  1.63015
```

```
##
## Random effects:
## Groups      Name          Variance Std.Dev.
## Code        (Intercept) 790.4      28.11
## Site        (Intercept) 188.9      13.74
## Residual                    857.3      29.28
## Number of obs: 459, groups: Code, 434; Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    92.789     3.588 20.914  25.862  <2e-16 ***
## agri.100.Scaled  9.033     3.108 31.860   2.907   0.0066 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr)
## agr.100.Scl -0.079
```

Since the LRT P-value is barely significant (0.0499) after the removal of urban.1000, I decided to calculate the AICc values for mod.Move.8 and mod.Move.9 in order to compare their parsimony.

## AIC test

I had to recreate the initial model for total time spent moving (Move) for the AIC test

```
mod.Move.0 <- lmer(Move ~ JulianDate.Scaled + A.Temp.Scaled + Time.Scaled + P
L.Scaled + Sex + wet.600.Scaled + for.veg.300.Scaled + agri.100.Scaled + urba
n.1000.Scaled + water.1000.Scaled + (1|Site) + (1|Code), data = MixedData, na
.action=na.exclude, REML = FALSE)

#define list of models
Move.models <- list(mod.Move.0, mod.Move.1, mod.Move.2, mod.Move.3, mod.Move.
4, mod.Move.5, mod.Move.6, mod.Move.7, mod.Move.8, mod.Move.9)

#specify model names
Move.mod.names <- c('mod.Move.0', 'mod.Move.1', 'mod.Move.2', 'mod.Move.3', '
mod.Move.4', 'mod.Move.5', 'mod.Move.6', 'mod.Move.7', 'mod.Move.8', 'mod.Mov
e.9')
```

*#calculate AICc of each model*

```
aictab(cand.set = Move.models, modnames = Move.mod.names)
```

```
##
## Model selection based on AICc:
##
##              K      AICc Delta_AICc AICcWt Cum.Wt      LL
## mod.Move.2 12 4676.26      0.00   0.41   0.41 -2325.78
```



```
## mod.Move.7  7 4677.52      1.26  0.22  0.62 -2331.63
## mod.Move.1 13 4678.30      2.04  0.15  0.77 -2325.73
## mod.Move.6  8 4678.78      2.52  0.12  0.88 -2331.23
## mod.Move.0 14 4680.43      4.17  0.05  0.93 -2325.73
## mod.Move.5  9 4680.73      4.47  0.04  0.98 -2331.16
## mod.Move.4 10 4682.63      6.37  0.02  0.99 -2331.07
## mod.Move.3 11 4684.54      8.28  0.01  1.00 -2330.97
## mod.Move.8  6 4729.05     52.79  0.00  1.00 -2358.43
## mod.Move.9  5 4730.84     54.58  0.00  1.00 -2360.35
```

mod.Move.8 and mod.Move.9 had an AICc value within 2 of each other which suggests that they are equally parsimonious, so I decided to average them.

## Averaging mod.Move.8 and mod.Move.9

```
# Average the two models using model.avg
averaged.model.move <- model.avg(mod.Move.8, mod.Move.9)

summary(averaged.model.move)

##
## Call:
## model.avg(object = mod.Move.8, mod.Move.9)
##
## Component model call:
## lmer(formula = <2 unique values>, data = MixedData, REML = FALSE,
##       na.action = na.exclude)
##
## Component models:
##    df  logLik   AICc delta weight
## 12  6 -2358.43 4729.05  0.00  0.71
##  1  5 -2360.35 4730.84  1.79  0.29
##
## Term codes:
##   agri.100.Scaled urban.1000.Scaled
##               1                2
##
## Model-averaged coefficients:
## (full average)
##              Estimate Std. Error Adjusted SE z value Pr(>|z|)
## (Intercept)    92.601     3.374      3.383  27.374  <2e-16 ***
## agri.100.Scaled    7.940     3.093      3.101   2.560   0.0105 *
## urban.1000.Scaled  4.756     4.098      4.103   1.159   0.2464
##
## (conditional average)
##              Estimate Std. Error Adjusted SE z value Pr(>|z|)
## (Intercept)    92.601     3.374      3.383  27.374  <2e-16 ***
## agri.100.Scaled    7.940     3.093      3.101   2.560   0.0105 *
## urban.1000.Scaled  6.698     3.263      3.272   2.047   0.0406 *
```

```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

To create the final model I will extract the conditional average coefficients, as they are only averaged over the models where the fixed effects appear.

```
#Extracting the conditional average coefficients from the averaged model
con.average.coefs.move <- averaged.model.move$coefs[,1]
```

## Final model

### Summary statistics

I changed the REML to TRUE to calculate the summary statistics of the final model.

```
#Creating model with just the conditional average coefficients
con.average.model.move <- lmer(Move ~ agri.100.Scaled + urban.1000.Scaled + (
1|Site) + (1|Code),
                                data = MixedData,
                                REML = TRUE,
                                na.action = na.exclude,
                                weights = con.average.coefs.move)
summary(con.average.model.move)

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: Move ~ agri.100.Scaled + urban.1000.Scaled + (1 | Site) + (1 |
##      Code)
##      Data: MixedData
## Weights: con.average.coefs.move
##
## REML criterion at convergence: 4704.3
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.95468 -0.50476  0.06275  0.52497  1.60171
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
##  Code     (Intercept)  794.8      28.19
##  Site     (Intercept)  177.8      13.33
##  Residual                    854.1      29.23
## Number of obs: 459, groups:  Code, 434; Site, 23
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    92.424     3.523 17.918  26.238 9.51e-16 ***
## agri.100.Scaled    7.581     3.148 27.397   2.408  0.0230 *
## urban.1000.Scaled    6.689     3.496 19.258   1.913  0.0707 .
```

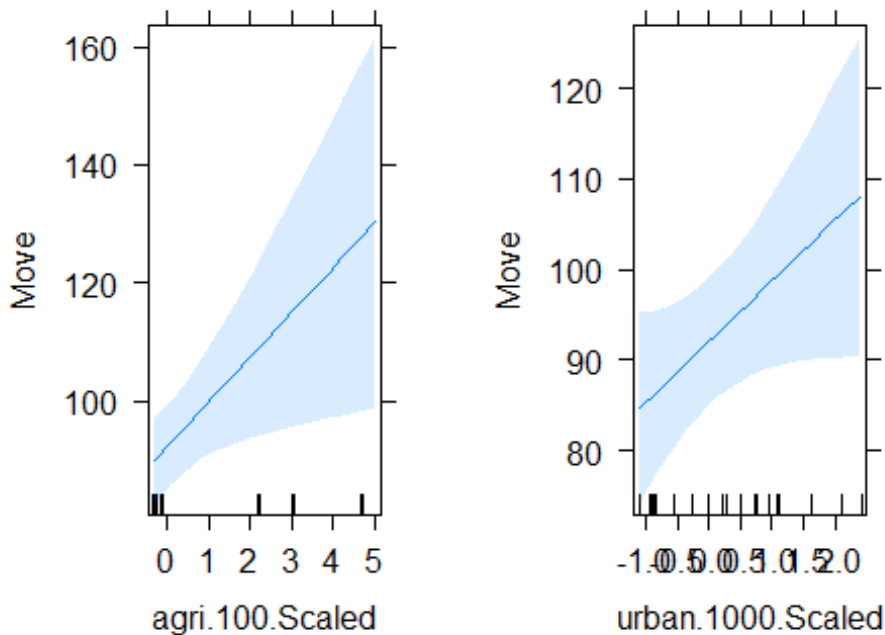
```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) a.100.
## agr.100.Scl -0.061
## urbn.1000.S -0.057 -0.237
```

Proportion of urban land area at 1000m does not have a statistically significant effect on total time spent moving.

## Visualization of the predictor effects

```
plot(allEffects(con.average.model.move))
```

### agri.100.Scaled effect ploturban.1000.Scaled effect plot



## Calculation of the marginal and conditional variance explained by the final model

```
r.squaredGLMM(con.average.model.move)
```

```
##              R2m              R2c
## [1,] 0.05853838 0.5597985
```

Marginal R2: fixed effects R2. Conditional R2: fixed and random effects R2.

## Calculation of the 95% confidence intervals

```
confint(con.average.model.move, level = 0.95, method = "Wald")
```

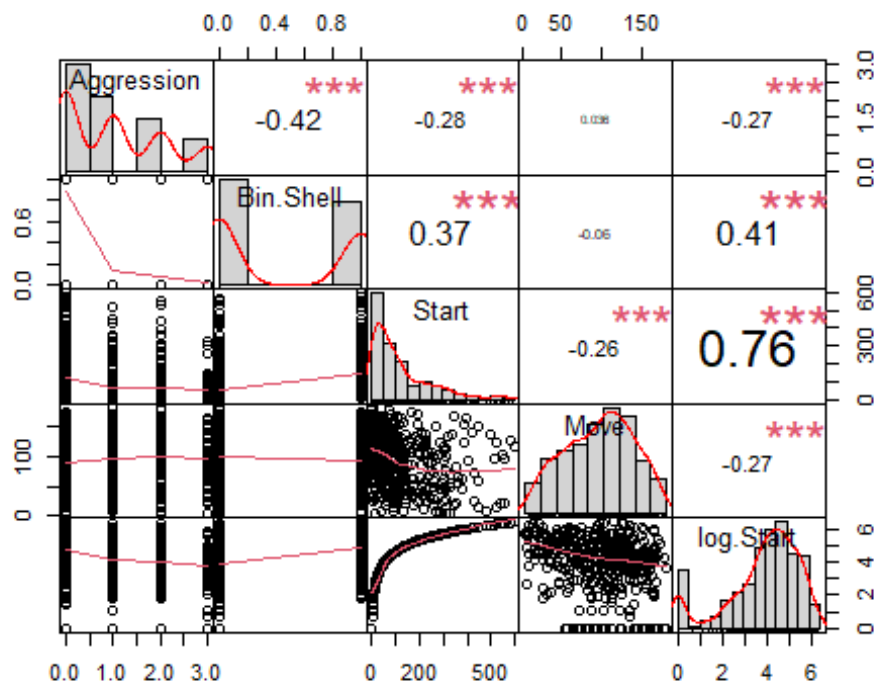
```
##                2.5 %   97.5 %
## .sig01          NA      NA
## .sig02          NA      NA
## .sigma          NA      NA
## (Intercept)    85.5197263 99.32801
## agri.100.Scaled  1.4108529 13.75138
## urban.1000.Scaled -0.1628494 13.54157
```

## Calculation of Pearson and Spearman correlation coefficients between behaviours

### Pearson correlation coefficients

*# Visualization of the correlations*

```
cor.behaviour <- MixedData[,c(11,13,14,15,68)]
chart.Correlation(cor.behaviour, histogram=TRUE, pch=19)
```



*# Creation of the correlation table*

```
table.corr.pearson.behaviour <- rcorr(as.matrix(MixedData[,c(11,13,14,15,68)]),
  type="pearson")
table.cor.pearson.behaviour.r <- table.corr.pearson.behaviour$r # Pearson correlation coefficients
table.cor.pearson.behaviour.p <- table.corr.pearson.behaviour$P # P values of the correlations
table.cor.pearson.behaviour.r
```

```
##           Aggression  Bin.Shell      Start      Move  log.Start
## Aggression  1.00000000 -0.42367744 -0.2759465  0.03613673 -0.2663096
## Bin.Shell  -0.42367744  1.00000000  0.3699742 -0.06016433  0.4081813
## Start      -0.27594651  0.36997418  1.0000000 -0.26028477  0.7558242
## Move        0.03613673 -0.06016433 -0.2602848  1.00000000 -0.2732679
## log.Start  -0.26630957  0.40818132  0.7558242 -0.27326794  1.0000000
```

```
table.cor.pearson.behaviour.p
```

```
##           Aggression  Bin.Shell      Start      Move  log.Star
t
## Aggression          NA 0.000000e+00 1.748938e-09 4.399102e-01 6.582143e-0
9
## Bin.Shell  0.000000e+00          NA 2.220446e-16 1.982237e-01 0.000000e+0
0
## Start      1.748938e-09 2.220446e-16          NA 1.522212e-08 0.000000e+0
0
## Move        4.399102e-01 1.982237e-01 1.522212e-08          NA 2.644465e-0
9
## log.Start   6.582143e-09 0.000000e+00 0.000000e+00 2.644465e-09          N
A
```

### *Spearman correlation coefficients*

*# Creation of the correlation table*

```
table.corr.spearman.behaviour <- rcorr(as.matrix(MixedData[,c(11,13,14,15,68)
]), type="spearman")
```

```
table.cor.spearman.behaviour.r <- table.corr.spearman.behaviour$r # Pearson c
orrelation coefficients
```

```
table.cor.spearman.behaviour.p <- table.corr.spearman.behaviour$P # P values
of the correlations
```

```
table.cor.spearman.behaviour.r
```

```
##           Aggression  Bin.Shell      Start      Move  log.Start
## Aggression  1.00000000 -0.4416514 -0.3028358  0.05072592 -0.3028358
## Bin.Shell  -0.44165143  1.0000000  0.4269960 -0.06471060  0.4269960
## Start      -0.30283584  0.4269960  1.0000000 -0.28682170  1.0000000
## Move        0.05072592 -0.0647106 -0.2868217  1.00000000 -0.2868217
## log.Start  -0.30283584  0.4269960  1.0000000 -0.28682170  1.0000000
```

```
table.cor.spearman.behaviour.p
```

```
##           Aggression  Bin.Shell      Start      Move  log.Start
## Aggression          NA 0.00000000 3.270673e-11 2.781425e-01 3.270673e-11
## Bin.Shell  0.000000e+00          NA 0.000000e+00 1.663435e-01 0.000000e+00
## Start      3.270673e-11 0.00000000          NA 3.844765e-10 0.000000e+00
## Move        2.781425e-01 0.1663435 3.844765e-10          NA 3.844765e-10
## log.Start   3.270673e-11 0.00000000 0.000000e+00 3.844765e-10          NA
```