

Cats Example: Two-Factor Repeated Measures with Baseline as Covariate

A study was done to compare 2 Treatments (A and C) for diabetes in cats. 6 Cats were randomly assigned to each treatment. Hence cat is nested within treatment. Or equivalently, treatment is the between-subjects factor. Blood sugar (Y) measurements were taken at 4 equally spaced Times (0, 1, 2, 3). The first time point (Time=0) was pre-treatment and serves as a baseline.

We consider several approaches:

1. Basic Repeated measures
2. Repeated measures with Time0 (baseline) as covariate
3. Repeated measures on Difference values

Notes:

1. We start by creating unique IDs for each of the cats. Originally labelled as 1-6 for Trt A and 1-6 for Trt C.
2. Different covariance structures were NOT investigated, but certainly could be.

```
library(tidyverse)
library(lme4)
library(lmerTest)
library(pbkrtest)
library(emmeans)
CatsData <- read.csv("C:/hess/STAT512/RNotes/Random3/R3_Cats.csv")
str(CatsData)
```

```
## 'data.frame':   48 obs. of  4 variables:
## $ Trt : Factor w/ 2 levels "A","C": 1 1 1 1 1 1 1 1 1 1 ...
## $ Cat : int  1 1 1 1 2 2 2 2 3 3 ...
## $ Time: int  0 1 2 3 0 1 2 3 0 1 ...
## $ Y : num  70.9 64.9 61.7 56.6 70.4 ...
```

```
#Important: Need to define things as.factor!!!
CatsData$Cat <- as.factor(CatsData$Cat)
CatsData$Time <- as.factor(CatsData$Time)
#Original Cat IDs are NOT Unique
with(table(Cat, Trt), data = CatsData)
```

```
##      Trt
## Cat A C
##   1 4 4
##   2 4 4
##   3 4 4
##   4 4 4
##   5 4 4
##   6 4 4
```

```
#Create unique IDs
CatsData$newID <- paste(CatsData$Trt, CatsData$Cat, sep = "")
with(table(newID, Trt), data = CatsData)
```

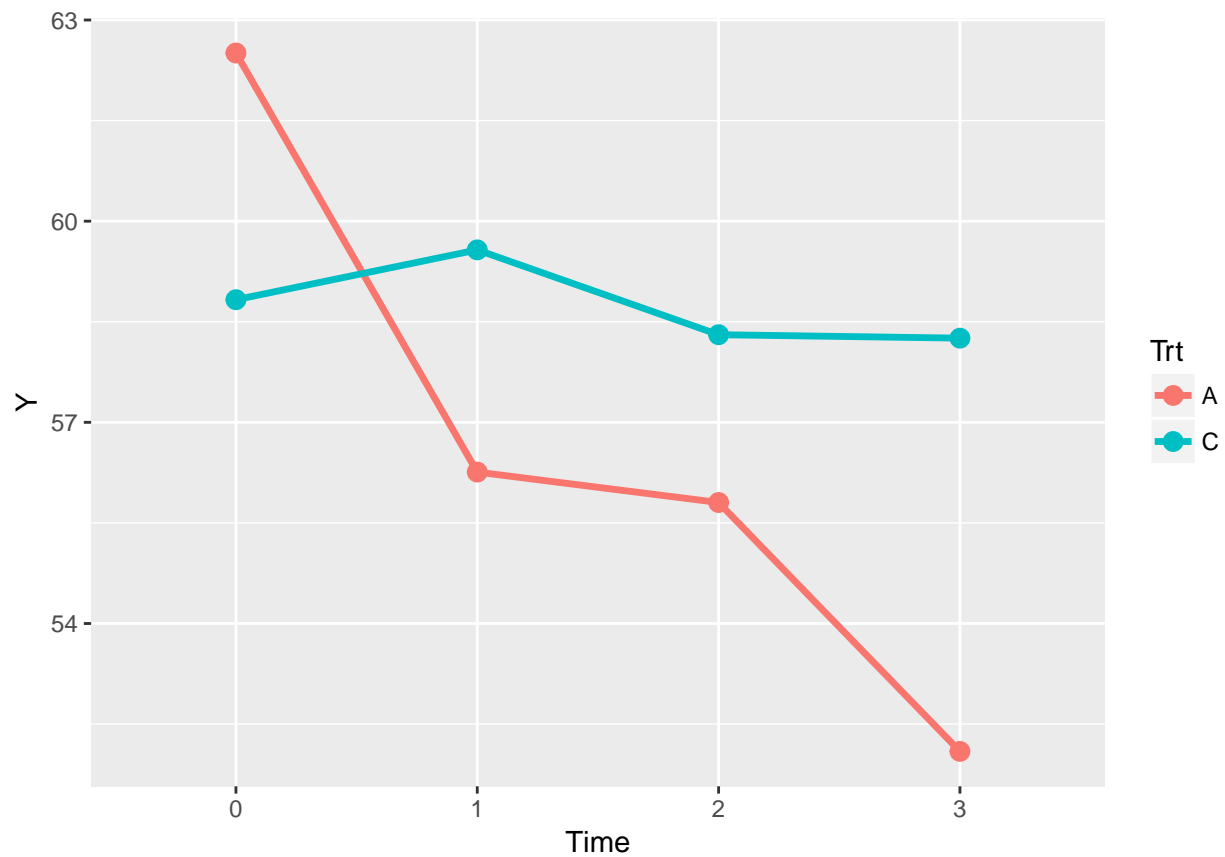
```
##      Trt
## newID A C
##   A1 4 0
```

```
## A2 4 0
## A3 4 0
## A4 4 0
## A5 4 0
## A6 4 0
## C1 0 4
## C2 0 4
## C3 0 4
## C4 0 4
## C5 0 4
## C6 0 4
```

```
head(CatsData)
```

```
## Trt Cat Time Y newID
## 1 A 1 0 70.89283 A1
## 2 A 1 1 64.92599 A1
## 3 A 1 2 61.68156 A1
## 4 A 1 3 56.56081 A1
## 5 A 2 0 70.41208 A2
## 6 A 2 1 67.64846 A2
```

```
SumStats <- aggregate(Y ~ Trt + Time, FUN = mean, data = CatsData)
qplot(x = Time, y = Y, colour = Trt, group = Trt, data = SumStats) + geom_line(size=1.2) +
  geom_point(size=3)
```



Model1A: Basic Model (non-unique IDS)

Since (1) cats are nested within trt and (2) cats were not given unique IDS, we need to explicitly specify the nesting in the model.

```
Model1A <- lmer(Y ~ Trt*Time + (1|Trt:Cat), data = CatsData)
Model1A

## Linear mixed model fit by REML ['merModLmerTest']
## Formula: Y ~ Trt * Time + (1 | Trt:Cat)
## Data: CatsData
## REML criterion at convergence: 249.0432
## Random effects:
## Groups Name Std.Dev.
## Trt:Cat (Intercept) 7.084
## Residual 3.091
## Number of obs: 48, groups: Trt:Cat, 12
## Fixed Effects:
## (Intercept) TrtC Time1 Time2 Time3
## 62.508 -3.679 -6.250 -6.704 -10.415
## TrtC:Time1 TrtC:Time2 TrtC:Time3
## 6.993 6.182 9.842
anova(Model1A, ddf = "Kenward-Roger")

## Analysis of Variance Table of type III with Kenward-Roger
## approximation for degrees of freedom
## Sum Sq Mean Sq NumDF DenDF F.value Pr(>F)
## Trt 2.347 2.347 1 10 0.2457 0.630853
## Time 187.831 62.610 3 30 6.5532 0.001535 **
## Trt:Time 154.615 51.538 3 30 5.3943 0.004328 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Model1B: Basic Model (unique IDS)

Now we rerun the basic model. Since we are now working with unique IDS, the nesting does not need to be explicitly specified in the model. The results are the same as Model1A.

```
Model1B <- lmer(Y ~ Trt*Time + (1|newID), data = CatsData)
summary(Model1B)

## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Y ~ Trt * Time + (1 | newID)
## Data: CatsData
##
## REML criterion at convergence: 249
##
## Scaled residuals:
## Min 1Q Median 3Q Max
## -1.9770 -0.6500 0.0400 0.6279 1.6428
##
## Random effects:
```

```

## Groups Name Variance Std.Dev.
## newID (Intercept) 50.182 7.084
## Residual 9.554 3.091
## Number of obs: 48, groups: newID, 12
##
## Fixed effects:
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) 62.508 3.155 12.832 19.810 5.30e-11 ***
## TrtC -3.679 4.462 12.832 -0.825 0.424685
## Time1 -6.250 1.785 30.000 -3.502 0.001468 **
## Time2 -6.704 1.785 30.000 -3.756 0.000742 ***
## Time3 -10.415 1.785 30.000 -5.836 2.21e-06 ***
## TrtC:Time1 6.993 2.524 30.000 2.771 0.009503 **
## TrtC:Time2 6.182 2.524 30.000 2.449 0.020364 *
## TrtC:Time3 9.842 2.524 30.000 3.900 0.000503 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
## (Intr) TrtC Time1 Time2 Time3 TrC:T1 TrC:T2
## TrtC -0.707
## Time1 -0.283 0.200
## Time2 -0.283 0.200 0.500
## Time3 -0.283 0.200 0.500 0.500
## TrtC:Time1 0.200 -0.283 -0.707 -0.354 -0.354
## TrtC:Time2 0.200 -0.283 -0.354 -0.707 -0.354 0.500
## TrtC:Time3 0.200 -0.283 -0.354 -0.354 -0.707 0.500 0.500

anova(Model1B, ddf = "Kenward-Roger")

## Analysis of Variance Table of type III with Kenward-Roger
## approximation for degrees of freedom
## Sum Sq Mean Sq NumDF DenDF F.value Pr(>F)
## Trt 2.347 2.347 1 10 0.2457 0.630853
## Time 187.831 62.610 3 30 6.5532 0.001535 **
## Trt:Time 154.615 51.538 3 30 5.3943 0.004328 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

emmeans(Model1B, pairwise ~ Trt|Time)

## $emmeans
## Time = 0:
## Trt emmean SE df lower.CL upper.CL
## A 62.50801 3.155318 12.83 55.68230 69.33372
## C 58.82859 3.155318 12.83 52.00288 65.65430
##
## Time = 1:
## Trt emmean SE df lower.CL upper.CL
## A 56.25795 3.155318 12.83 49.43224 63.08366
## C 59.57181 3.155318 12.83 52.74610 66.39752
##
## Time = 2:
## Trt emmean SE df lower.CL upper.CL
## A 55.80448 3.155318 12.83 48.97877 62.63019

```

```
## C 58.30690 3.155318 12.83 51.48119 65.13261
##
## Time = 3:
## Trt emmean SE df lower.CL upper.CL
## A 52.09279 3.155318 12.83 45.26708 58.91849
## C 58.25529 3.155318 12.83 51.42958 65.08100
##
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
##
## $contrasts
## Time = 0:
## contrast estimate SE df t.ratio p.value
## A - C 3.679417 4.462293 12.83 0.825 0.4247
##
## Time = 1:
## contrast estimate SE df t.ratio p.value
## A - C -3.313859 4.462293 12.83 -0.743 0.4711
##
## Time = 2:
## contrast estimate SE df t.ratio p.value
## A - C -2.502422 4.462293 12.83 -0.561 0.5846
##
## Time = 3:
## contrast estimate SE df t.ratio p.value
## A - C -6.162502 4.462293 12.83 -1.381 0.1909
```

Reformatting

In order to use Time0 (baseline) as a covariate or use Difference as the response, we need to do some reformatting. Specifically, we need to add a column with the Time0 values. This column can be used to calculate a difference (downstream time points versus Time 0). Since we are using Time0 as a covariate (or calculating difference versus Time0), we “drop” that period from the analysis.

Note: The `rename()` function from `dplyr` is very handy!

```
head(CatsData)
```

```
## Trt Cat Time Y newID
## 1 A 1 0 70.89283 A1
## 2 A 1 1 64.92599 A1
## 3 A 1 2 61.68156 A1
## 4 A 1 3 56.56081 A1
## 5 A 2 0 70.41208 A2
## 6 A 2 1 67.64846 A2
```

```
Time0 <- CatsData %>%
  filter(Time == 0) %>%
  rename(Time0 = Y) %>%
  select(-Trt, -Cat, -Time)
head(Time0)
```

```
## Time0 newID
## 1 70.89283 A1
## 2 70.41208 A2
```

```
## 3 69.79313    A3
## 4 54.96961    A4
## 5 55.04248    A5
## 6 53.93793    A6

nrow(Time0)

## [1] 12

WithTime0 <- full_join(CatsData, Time0) %>%
  mutate(Diff = Y - Time0) %>%
  filter(Time != 0)

## Joining, by = "newID"

head(WithTime0)

##   Trt Cat Time      Y newID   Time0    Diff
## 1   A   1   1 64.92599   A1 70.89283 -5.966840
## 2   A   1   2 61.68156   A1 70.89283 -9.211274
## 3   A   1   3 56.56081   A1 70.89283 -14.332024
## 4   A   2   1 67.64846   A2 70.41208 -2.763622
## 5   A   2   2 68.98425   A2 70.41208 -1.427830
## 6   A   2   3 62.08641   A2 70.41208 -8.325670

nrow(WithTime0)

## [1] 36
```

Model2: Using baseline (Time0) as a covariate

After reformatting, it is easy to include Time0 as a covariate!

```
Model2 <- lmer(Y ~ Time0 + Trt*Time + (1|newID), data = WithTime0)
summary(Model2)

## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Y ~ Time0 + Trt * Time + (1 | newID)
## Data: WithTime0
##
## REML criterion at convergence: 175.9
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.03044 -0.60779 -0.01715  0.68440  1.46115
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
## newID    (Intercept)    4.588     2.142
## Residual                    10.269     3.205
## Number of obs: 36, groups: newID, 12
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)   7.9260      6.8494  9.4520  1.157   0.2756
```

```

## Time0          0.7732      0.1066  9.0000   7.250 4.81e-05 ***
## TrtC           6.1588      2.2597 22.7390   2.725  0.0121 *
## Time2         -0.4535      1.8501 20.0000  -0.245  0.8089
## Time3         -4.1652      1.8501 20.0000  -2.251  0.0358 *
## TrtC:Time2     -0.8114      2.6165 20.0000  -0.310  0.7597
## TrtC:Time3      2.8486      2.6165 20.0000   1.089  0.2892
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) Time0  TrtC   Time2  Time3  TrC:T2
## Time0      -0.973
## TrtC        -0.329  0.174
## Time2       -0.135  0.000  0.409
## Time3       -0.135  0.000  0.409  0.500
## TrtC:Time2   0.095  0.000 -0.579 -0.707 -0.354
## TrtC:Time3   0.095  0.000 -0.579 -0.354 -0.707  0.500
anova(Model2, ddf = "Kenward-Roger")

## Analysis of Variance Table of type III with Kenward-Roger
## approximation for degrees of freedom
##          Sum Sq Mean Sq NumDF DenDF F.value    Pr(>F)
## Time0      539.80  539.80     1     9  52.566 4.815e-05 ***
## Trt        170.00  170.00     1     9  16.555  0.002805 **
## Time       47.16   23.58     2    20   2.296  0.126533
## Trt:Time   22.17   11.08     2    20   1.079  0.358776
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
emmeans(Model2, pairwise ~ Trt|Time)

## $emmeans
## Time = 1:
##   Trt    emmean      SE    df lower.CL upper.CL
##   A    54.83547  1.585782 23.01  51.55514  58.11579
##   C    60.99430  1.585782 23.01  57.71397  64.27462
##
## Time = 2:
##   Trt    emmean      SE    df lower.CL upper.CL
##   A    54.38199  1.585782 23.01  51.10167  57.66232
##   C    59.72939  1.585782 23.01  56.44906  63.00971
##
## Time = 3:
##   Trt    emmean      SE    df lower.CL upper.CL
##   A    50.67030  1.585782 23.01  47.38998  53.95062
##   C    59.67777  1.585782 23.01  56.39745  62.95810
##
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
##
## $contrasts
## Time = 1:
##   contrast estimate      SE    df t.ratio p.value
##   A - C    -6.158829  2.259733 22.74  -2.725  0.0121

```

```
##
## Time = 2:
## contrast estimate SE df t.ratio p.value
## A - C -5.347393 2.259733 22.74 -2.366 0.0269
##
## Time = 3:
## contrast estimate SE df t.ratio p.value
## A - C -9.007472 2.259733 22.74 -3.986 0.0006
```

Model3: Using Diff (from Baseline) as the Response

```
Model3 <- lmer(Diff ~ Trt*Time + (1|newID), data = WithTime0)
summary(Model3)
```

```
## Linear mixed model fit by REML t-tests use Satterthwaite approximations
## to degrees of freedom [lmerMod]
## Formula: Diff ~ Trt * Time + (1 | newID)
## Data: WithTime0
##
## REML criterion at convergence: 177.3
##
## Scaled residuals:
## Min 1Q Median 3Q Max
## -1.78233 -0.60578 -0.00503 0.50279 1.70926
##
## Random effects:
## Groups Name Variance Std.Dev.
## newID (Intercept) 7.41 2.722
## Residual 10.27 3.205
## Number of obs: 36, groups: newID, 12
##
## Fixed effects:
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) -6.2501 1.7165 22.2000 -3.641 0.00143 **
## TrtC 6.9933 2.4275 22.2000 2.881 0.00863 **
## Time2 -0.4535 1.8501 20.0000 -0.245 0.80887
## Time3 -4.1652 1.8501 20.0000 -2.251 0.03577 *
## TrtC:Time2 -0.8114 2.6165 20.0000 -0.310 0.75967
## TrtC:Time3 2.8486 2.6165 20.0000 1.089 0.28922
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
## (Intr) TrtC Time2 Time3 TrC:T2
## TrtC -0.707
## Time2 -0.539 0.381
## Time3 -0.539 0.381 0.500
## TrtC:Time2 0.381 -0.539 -0.707 -0.354
## TrtC:Time3 0.381 -0.539 -0.354 -0.707 0.500
```

```
anova(Model3, ddf = "Kenward-Roger")
```

```
## Analysis of Variance Table of type III with Kenward-Roger
```



```
## approximation for degrees of freedom
##          Sum Sq Mean Sq NumDF DenDF F.value   Pr(>F)
## Trt      167.400 167.400     1    10 16.3015 0.002371 **
## Time      47.164  23.582     2    20  2.2964 0.126533
## Trt:Time   22.169  11.085     2    20  1.0794 0.358776
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

emmeans(Model3, pairwise ~ Trt|Time)

## $emmeans
## Time = 1:
## Trt      emmean      SE    df  lower.CL upper.CL
## A      -6.250588 1.716537 22.2  -9.808082 -2.692036
## C       0.7432166 1.716537 22.2  -2.814806  4.301239
##
## Time = 2:
## Trt      emmean      SE    df  lower.CL upper.CL
## A      -6.7035326 1.716537 22.2 -10.261555 -3.145510
## C      -0.5216935 1.716537 22.2  -4.079716  3.036329
##
## Time = 3:
## Trt      emmean      SE    df  lower.CL upper.CL
## A     -10.4152255 1.716537 22.2 -13.973248 -6.857203
## C      -0.5733070 1.716537 22.2  -4.131330  2.984716
##
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
##
## $contrasts
## Time = 1:
## contrast estimate      SE    df t.ratio p.value
## A - C      -6.993275 2.42755 22.2  -2.881  0.0086
##
## Time = 2:
## contrast estimate      SE    df t.ratio p.value
## A - C      -6.181839 2.42755 22.2  -2.547  0.0183
##
## Time = 3:
## contrast estimate      SE    df t.ratio p.value
## A - C     -9.841918 2.42755 22.2  -4.054  0.0005
```