## GEE and GLME

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```
library(tidyverse)
library(geepack)
library(nlme)
library(glme)
library(lme4)
aids = read.csv("aids.csv")
aids = aids %>%
  mutate(occasion = ceiling(week),
         gender = factor(gender, level = c("male", "female")),
         treatment = as.factor(treatment))
aids = aids %>%
  mutate(counts = exp(log_cd4) - 1) %>%
  mutate(counts2 = as.integer(round(counts))) %>%
  mutate(occasion_sqr = I(occasion^2))
log_cd4 ~ occasion + I(occasion^2) + treatment:occasion + treatment:I(occasion^2) + age
model = glmer(counts2 ~ occasion + I(occasion ^ 2) + treatment:occasion + treatment:I(occasion^2) + age
              data = aids,
              family = poisson,
              control = glmerControl(tol = 1e-12),
              nAGQ = 0,
              na.action = na.omit)
summary(model)
## Generalized linear mixed model fit by maximum likelihood (Adaptive
     Gauss-Hermite Quadrature, nAGQ = 0) [glmerMod]
## Family: poisson ( log )
## Formula:
## counts2 ~ occasion + I(occasion^2) + treatment:occasion + treatment:I(occasion^2) +
##
       age + (1 | id)
      Data: aids
##
## Control: glmerControl(tol = 1e-12)
##
                 BIC
                      logLik deviance df.resid
##
  62503.4 62575.2 -31240.7 62481.4
                                            5025
##
## Scaled residuals:
```

```
Median
                 1Q
## -12.0587 -1.3875 -0.2441 1.0653 22.9876
##
## Random effects:
## Groups Name
                      Variance Std.Dev.
          (Intercept) 0.9138 0.9559
## id
## Number of obs: 5036, groups: id, 1309
## Fixed effects:
##
                             Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                            2.478e+00 1.265e-01 19.582 < 2e-16 ***
## occasion
                            1.703e-03 1.646e-03
                                                   1.035 0.300676
## I(occasion^2)
                           -4.932e-04 4.777e-05 -10.325 < 2e-16 ***
                            1.091e-02 3.273e-03
                                                  3.334 0.000856 ***
                            1.506e-03 2.223e-03
                                                   0.677 0.498096
## occasion:treatment2
## occasion:treatment3
                            2.809e-02 2.149e-03
                                                  13.072 < 2e-16 ***
## occasion:treatment4
                            4.880e-02 2.075e-03 23.521 < 2e-16 ***
## I(occasion^2):treatment2 9.251e-05 6.374e-05
                                                   1.451 0.146699
## I(occasion^2):treatment3 -5.108e-04 6.148e-05 -8.309 < 2e-16 ***
## I(occasion^2):treatment4 -7.852e-04 5.878e-05 -13.358 < 2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##
               (Intr) occasn I(c^2) age occs:2 occs:3 occs:4 I(^2):2 I(^2):3
## occasion
               -0.015
## I(occasn^2) 0.011 -0.947
              -0.977 0.001 -0.001
## age
## occsn:trtm2 0.002 -0.738 0.699 -0.001
## occsn:trtm3 0.002 -0.763 0.723 0.000 0.565
## occsn:trtm4 0.004 -0.791 0.749 -0.002 0.585 0.605
## I(ccsn^2):2 -0.002 0.708 -0.748 0.000 -0.947 -0.542 -0.561
## I(ccsn^2):3 -0.001 0.734 -0.776 0.000 -0.543 -0.947 -0.582 0.581
## I(ccsn^2):4 -0.003 0.768 -0.811 0.001 -0.568 -0.588 -0.947 0.608
                                                                        0.630
We noticed that cannot have occasion 2 as a random variable, which doesn't allow us to test our ideal model.
We will instead just use a random intercept and a random occasion slope.
model1 = glmer(counts2 ~ occasion + I(occasion ^ 2) + treatment:occasion + treatment:I(occasion^2) + ag
              data = aids,
              family = poisson,
              control = glmerControl(tol = 1e-12),
             nAGQ = 0,
             na.action = na.omit)
summary(model1)
## Generalized linear mixed model fit by maximum likelihood (Adaptive
     Gauss-Hermite Quadrature, nAGQ = 0) [glmerMod]
## Family: poisson (log)
## counts2 ~ occasion + I(occasion^2) + treatment:occasion + treatment:I(occasion^2) +
##
       age + (1 + occasion | id)
##
      Data: aids
## Control: glmerControl(tol = 1e-12)
##
```

```
logLik deviance df.resid
       AIC
                BIC
   52553.6 52638.4 -26263.8 52527.6
##
##
## Scaled residuals:
##
                 1Q
                      Median
                                   3Q
  -10.0615 -1.0129 -0.0878
##
                               0.8022
                                      18.4080
##
## Random effects:
##
   Groups Name
                      Variance Std.Dev. Corr
##
           (Intercept) 0.8678324 0.93158
          occasion
                      0.0009204 0.03034
                                        -0.15
## Number of obs: 5036, groups: id, 1309
## Fixed effects:
                             Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                            2.543e+00 1.228e-01
                                                 20.720 < 2e-16 ***
## occasion
                            5.272e-03
                                       2.493e-03
                                                   2.115
                                                           0.0344 *
## I(occasion^2)
                           -7.719e-04 5.281e-05 -14.617
                                                         < 2e-16 ***
                            9.905e-03 3.173e-03
                                                   3.122
                                                           0.0018 **
## age
## occasion:treatment2
                            5.524e-03
                                      3.445e-03
                                                   1.604
                                                           0.1088
## occasion:treatment3
                            2.737e-02 3.384e-03
                                                   8.088 6.05e-16 ***
                            4.555e-02 3.326e-03
## occasion:treatment4
                                                 13.695
                                                         < 2e-16 ***
## I(occasion^2):treatment2 -7.764e-05 7.076e-05
                                                 -1.097
                                                           0.2725
## I(occasion^2):treatment3 -7.229e-04 6.828e-05 -10.588
                                                         < 2e-16 ***
## I(occasion^2):treatment4 -8.695e-04 6.585e-05 -13.205 < 2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation of Fixed Effects:
##
              (Intr) occasn I(c^2) age occs:2 occs:3 occs:4 I(^2):2 I(^2):3
## occasion
              -0.025
## I(occasn^2) 0.010 -0.625
              -0.976 0.002
                             0.000
## occsn:trtm2 0.001 -0.716
                             0.448 0.000
## occsn:trtm3 0.001 -0.729
                             0.457
                                   0.000 0.527
## occsn:trtm4 0.003 -0.742 0.465 -0.002 0.536 0.546
## I(ccsn^2):2 -0.002 0.463 -0.745 0.001 -0.605 -0.341 -0.347
## I(ccsn^2):3 -0.001 0.480 -0.772 -0.001 -0.347 -0.593 -0.359
## I(ccsn^2):4 -0.003 0.498 -0.801 0.001 -0.360 -0.366 -0.587 0.597
```

The best model that we have so far is:

 $log(count) = \beta_0 + \beta_1 * occasion + \beta_2 * occasion^2 + \beta_3 - 5 * treatment : occasion + \beta_6 - 8 * treatment : occasion^2$ 

## Comparing the Models

```
anova(model, model1)

## Data: aids

## Models:

## model: counts2 ~ occasion + I(occasion^2) + treatment:occasion + treatment:I(occasion^2) +

## model: age + (1 | id)

## model1: counts2 ~ occasion + I(occasion^2) + treatment:occasion + treatment:I(occasion^2) +
```

```
age + (1 + occasion | id)
## model1:
##
                AIC
                     BIC logLik deviance Chisq Df Pr(>Chisq)
         npar
## model
            11 62503 62575 -31241
                                     62481
            13 52554 52638 -26264
                                     52528 9953.8 2 < 2.2e-16 ***
## model1
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
H_0: the reduced model with only a random intercept is adequate.
```

Because the p-value < 2.2e-16, is less than  $\alpha = 0.05$ , we reject the null and conclude that the model with both the random intercept and the random slope fits the data better.

We will also compare these models to the quadratic model previously.

```
model_quadratic2 = lme(log_cd4 ~ occasion + I(occasion^2) +
             treatment:occasion + treatment:I(occasion^2)
               + age, data = aids,
             random = ~ occasion + I(occasion^2) | id,
             method = "ML")
summary(model_quadratic2)
## Linear mixed-effects model fit by maximum likelihood
```

```
Data: aids
##
          AIC
                   {\tt BIC}
##
                          logLik
##
     11888.81 11999.73 -5927.406
##
## Random effects:
## Formula: ~occasion + I(occasion^2) | id
  Structure: General positive-definite, Log-Cholesky parametrization
##
                 StdDev
                             Corr
## (Intercept)
                 0.759158207 (Intr) occasn
## occasion
                 0.041653870 0.327
## I(occasion^2) 0.000946246 -0.393 -0.921
## Residual
                 0.547194358
##
## Fixed effects: log_cd4 ~ occasion + I(occasion^2) + treatment:occasion + treatment:I(occasion^2) +
                                 Value Std.Error
##
                                                    \mathsf{DF}
                                                         t-value p-value
                             2.5752308 0.11579256 3719 22.240037 0.0000
## (Intercept)
                            -0.0121732 0.00490191 3719 -2.483360 0.0131
## occasion
## I(occasion^2)
                            -0.0001247 0.00013385 3719 -0.931732 0.3515
                             0.0096564 0.00299342 1307
                                                        3.225870
                                                                   0.0013
## age
## occasion:treatment2
                             0.0073144 0.00686356 3719
                                                        1.065682
                                                                   0.2866
                             0.0186455 0.00685665 3719 2.719328
## occasion:treatment3
                                                                  0.0066
## occasion:treatment4
                             0.0438280 0.00679153 3719 6.453332 0.0000
## I(occasion^2):treatment2 -0.0001423 0.00018672 3719 -0.762371
                                                                   0.4459
## I(occasion^2):treatment3 -0.0003629 0.00018812 3719 -1.928913 0.0538
## I(occasion^2):treatment4 -0.0008380 0.00018513 3719 -4.526468 0.0000
## Correlation:
##
                            (Intr) occasn I(c^2) age
                                                        occs:2 occs:3 occs:4
## occasion
                            -0.026
## I(occasion^2)
                             0.016 - 0.939
## age
                            -0.975 0.003 -0.002
## occasion:treatment2
                             0.002 -0.707 0.666 -0.001
```

0.002 -0.707 0.667 -0.002 0.505

0.002 -0.714 0.673 -0.001 0.510 0.510

## occasion:treatment3

## occasion:treatment4

```
## I(occasion^2):treatment3 -0.002 0.663 -0.709 0.002 -0.474 -0.939 -0.479
## I(occasion^2):treatment4 0.000 0.674 -0.720 -0.001 -0.481 -0.482 -0.939
##
                            I(^2):2\ I(^2):3
## occasion
## I(occasion^2)
## age
## occasion:treatment2
## occasion:treatment3
## occasion:treatment4
## I(occasion^2):treatment2
## I(occasion^2):treatment3  0.508
## I(occasion^2):treatment4  0.516
                                     0.512
## Standardized Within-Group Residuals:
##
                        Q1
                                   Med
                                                QЗ
                                                           Max
## -4.48500432 -0.41811557 0.02420634 0.47119729 3.88851463
##
## Number of Observations: 5036
## Number of Groups: 1309
```

Comparing the AIC of the GLME model with our previous LME model:

- GLME AIC = 52553.6
- LME AIC = 11888.81

Because the GLME AIC is greater than the LME AIC, we can conclude that the LME model we created previously is better (fits the data better) than the GLME.