Generalized Linear Mixed Models

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Read in the individual data (or a pairwise dataset)	

```
library(tidyr)
library(dplyr)
#install.packages("lme4")
library(lme4)
acitelli ind <- read.csv("/Users/randigarcia/Desktop/Three-day-workshop/R Workshop/Data/
```

Convert individual data to pairwise. I also create a simhobs variable that will be our binary response, two dummy variables that will be useful for estimating separate random intercepts for men and women, and a count variable cigarettes.

```
tempA <- acitelli ind %>%
 mutate(genderE = gender, partnum = 1) %>%
 mutate(gender = ifelse(gender == 1, "A", "P")) %>%
 gather(variable, value, self_pos:genderE) %>%
 unite(var_gender, variable, gender) %>%
 spread(var gender, value)
tempB <- acitelli ind %>%
 mutate(genderE = gender, partnum = 2) %>%
 mutate(gender = ifelse(gender == 1, "P", "A")) %>%
 gather(variable, value, self_pos:genderE)%>%
 unite(var gender, variable, gender) %>%
 spread(var_gender, value)
```

Logistic Multilevel Modeling (Binary variables)

To account for the nonindependence, we can make use of the glmer() function from the lme4 package. Note that we are asking for the variance of intercepts across dyads, that is the random intercept in traditional multilevel modeling. The gls() function in the nlme package does not have an option for specifying a link function (i.e., there is no family = option). The syntax of glmer() differs a bit from gls() in that the random effects are specified within the formula: + (1/cuplid).

Indistinguishable Dyads

```
apim_bin <- glmer(simhob_bin_A ~ other_pos_A + other_pos_P
                  + (1|cuplid),
                  data = acitelli pair,
                  family = binomial,
                  na.action = na.omit)
summary(apim bin)
## Generalized linear mixed model fit by maximum likelihood (Laplace
##
     Approximation) [glmerMod]
## Family: binomial (logit)
## Formula: simhob bin A ~ other pos A + other pos P + (1 | cuplid)
      Data: acitelli pair
##
##
##
        AIC
                 BIC
                       logLik deviance df.resid
```

```
##
     322.5
              337.3 -157.3
                                314.5
                                           292
##
## Scaled residuals:
##
       Min
                      Median
                  1Q
                                    3Q
                                           Max
## -0.90316 -0.44848 -0.33245 0.02569
                                      2.03565
##
## Random effects:
## Groups Name
                      Variance Std.Dev.
## cuplid (Intercept) 1.659
                                1.288
## Number of obs: 296, groups: cuplid, 148
## Fixed effects:
              Estimate Std. Error z value Pr(>|z|)
                           2.4903 -3.345 0.000821 ***
## (Intercept)
              -8.3310
## other_pos_A
                0.9195
                           0.3894
                                    2.361 0.018217 *
## other pos P
                0.6723
                           0.3756
                                    1.790 0.073481 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
               (Intr) oth A
## other_pos_A -0.738
## other pos P -0.708 0.058
```

Distinguishable Dyads

Interaction approach.

```
## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula:
## simhob_bin_A ~ other_pos_A + other_pos_P + genderE_A + other_pos_A *
## genderE_A + other_pos_P * genderE_A + (man + woman - 1 | cuplid)
## Data: acitelli pair
```

```
##
     314.7
              347.9
                      -148.4
                                296.7
                                           287
##
## Scaled residuals:
##
       Min
                 1Q
                      Median
                                   3Q
                                           Max
## -1.09016 -0.48698 -0.00949 0.03293 2.52186
## Random effects:
## Groups Name Variance Std.Dev. Corr
## cuplid man
                183.8755 13.5601
##
          woman
                  0.1908 0.4368 1.00
## Number of obs: 296, groups: cuplid, 148
##
## Fixed effects:
##
                        Estimate Std. Error z value Pr(>|z|)
                                     15.095 -2.545
## (Intercept)
                         -38.413
                                                     0.0109 *
## other_pos_A
                           6.104
                                     2.453 2.489 0.0128 *
## other_pos_P
                           1.433
                                     2.551 0.562 0.5744
                                 14.859 -2.060 0.0394 *
## genderE A
                         -30.611
## other_pos_A:genderE_A
                                      2.462 2.037 0.0417 *
                          5.014
## other_pos_P:genderE_A 0.937
                                      2.559
                                              0.366
                                                     0.7142
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
              (Intr) oth_A oth_P gndE_A o__A:E
## other_pos_A -0.656
## other_pos_P -0.683 -0.098
## genderE_A
              0.988 -0.647 -0.675
## othr A:E A -0.635 0.985 -0.113 -0.643
## othr P:E A -0.662 -0.114 0.986 -0.670 -0.134
## convergence code: 0
## unable to evaluate scaled gradient
## Model failed to converge: degenerate Hessian with 2 negative eigenvalues
Two-intercept model.
#does not converge
apim_bin_di_two <- glmer(simhob_bin_A ~ gender_A + other_pos_A:gender_A + other_pos_P:ge
                        + (man + woman - 1|cuplid),
                        data = acitelli_pair,
                        family = binomial,
                        na.action = na.omit)
```

##

AIC

BIC

logLik deviance df.resid

apim_bin_di_two <- glmer(simhob_bin_A ~ gender_A + other_pos_A:gender_A + other_pos_P:ge

```
+ (1|cuplid),
                        data = acitelli pair,
                        family = binomial,
                        na.action = na.omit)
apim_bin_di_two <- glmer(simhob_bin_A ~ gender_A + other_pos_A:gender_A + other_pos_P:ge
                        + (man + woman - 1|cuplid),
                        data = acitelli_pair,
                        family = binomial,
                        na.action = na.omit,
                        nAGQ = 0) #Adaptive Gauss-Hermite Quadrature
summary(apim bin di two)
## Generalized linear mixed model fit by maximum likelihood (Adaptive
    Gauss-Hermite Quadrature, nAGQ = 0) [glmerMod]
##
## Family: binomial (logit)
## Formula:
## simhob_bin_A ~ gender_A + other_pos_A:gender_A + other_pos_P:gender_A -
       1 + (man + woman - 1 | cuplid)
##
##
     Data: acitelli_pair
##
##
       AIC
                BIC
                      logLik deviance df.resid
##
      330.4
              363.6
                      -156.2
                                312.4
                                            287
##
## Scaled residuals:
                                    3Q
                  1Q
                      Median
                                            Max
## -0.97710 -0.52321 -0.39856 0.00841
                                      1.90671
##
## Random effects:
## Groups Name Variance Std.Dev. Corr
## cuplid man
                1.257
                         1.121
          woman 1.038
                          1.019
                                   1.00
## Number of obs: 296, groups: cuplid, 148
##
## Fixed effects:
##
                           Estimate Std. Error z value Pr(>|z|)
## gender_Ahus
                            -7.1637
                                        2.7924 - 2.566
                                                        0.0103 *
## gender Awife
                            -6.0917
                                        2.4851 -2.451
                                                         0.0142 *
## gender_Ahus:other_pos_A
                                        0.5214 0.817 0.4141
                           0.4259
## gender_Awife:other_pos_A 1.0173
                                        0.4539 2.241 0.0250 *
## gender_Ahus:other_pos_P
                             0.9160
                                         0.5003 1.831 0.0671 .
## gender Awife:other pos P
                             0.1722
                                        0.4655 0.370 0.7114
## ---
```

Log-Linear Multilevel Modeling (Count variables)

Indistinguishable Dyads

```
apim_poi <- glmer(cigarettes_A ~ other_pos_A + other_pos_P</pre>
                  + (1|cuplid),
                  data = acitelli pair,
                  family = poisson,
                  na.action = na.omit)
summary(apim poi)
## Generalized linear mixed model fit by maximum likelihood (Laplace
     Approximation) [glmerMod]
## Family: poisson (log)
## Formula: cigarettes_A ~ other_pos_A + other_pos_P + (1 | cuplid)
      Data: acitelli pair
##
##
##
                       logLik deviance df.resid
        AIC
                 BIC
                       -440.8
##
      889.5
               904.3
                                 881.5
                                            292
##
## Scaled residuals:
      Min
                1Q Median
                                3Q
                                       Max
## -1.2675 -0.4623 -0.1371 0.4359 2.1573
## Random effects:
## Groups Name
                       Variance Std.Dev.
## cuplid (Intercept) 0.1776
                                0.4215
## Number of obs: 296, groups: cuplid, 148
##
## Fixed effects:
##
               Estimate Std. Error z value Pr(>|z|)
```

```
## (Intercept) 0.16159
                          0.66097
                                    0.244
                                              0.807
## other pos A 0.10373
                                              0.344
                           0.10972
                                     0.945
## other_pos_P -0.07413
                           0.10762 -0.689
                                              0.491
##
## Correlation of Fixed Effects:
##
               (Intr) oth A
## other pos A -0.715
## other pos P -0.696 0.006
## convergence code: 0
## Model failed to converge with max|grad| = 0.00120471 (tol = 0.001, component 1)
```

Distinguishable Dyads

Interaction approach.

```
## Generalized linear mixed model fit by maximum likelihood (Laplace
     Approximation) [glmerMod]
## Family: poisson (log)
## Formula:
## cigarettes A ~ other pos A + other pos P + genderE A + other pos A *
       genderE A + other pos P * genderE A + (man + woman - 1 |
##
                                                                     cuplid)
##
      Data: acitelli_pair
##
##
        AIC
                 BIC
                       logLik deviance df.resid
##
      898.1
               931.3
                       -440.1
                                 880.1
##
## Scaled residuals:
       Min
                1Q Median
                                3Q
                                       Max
## -1.2346 -0.4408 -0.1400 0.4723 2.2872
##
## Random effects:
## Groups Name Variance Std.Dev. Corr
##
   cuplid man
                 0.1651
                          0.4063
##
           woman 0.1828
                          0.4275
                                   1.00
## Number of obs: 296, groups: cuplid, 148
```

```
##
## Fixed effects:
##
                         Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                         0.09390
                                    0.66378
                                               0.142
                                                        0.888
## other pos A
                         0.10208
                                    0.11035
                                               0.925
                                                       0.355
## other pos P
                                  0.10877 -0.526
                         -0.05726
                                                       0.599
## genderE A
                         0.49516
                                  0.53704 0.922
                                                       0.357
## other pos A:genderE A 0.01354 0.11607
                                              0.117
                                                      0.907
## other_pos_P:genderE_A -0.12865
                                    0.11448 -1.124
                                                       0.261
##
## Correlation of Fixed Effects:
##
               (Intr) oth_A oth_P gndE_A o__A:E
## other pos A -0.710
## other_pos_P -0.695 -0.003
## genderE A -0.026 -0.083
                             0.117
## othr A:E A -0.092 0.096 0.029 -0.552
## othr__P:E_A 0.119 -0.010 -0.150 -0.527 -0.411
## convergence code: 0
## Model failed to converge with max|grad| = 0.0342649 (tol = 0.001, component 1)
Two-intercept model.
apim poi di two <- glmer(cigarettes A ~ gender A + other pos A:gender A + other pos P:ge
                         + (man + woman - 1|cuplid),
                         data = acitelli_pair,
                         family = poisson,
                         na.action = na.omit)
summary(apim_poi_di_two)
## Generalized linear mixed model fit by maximum likelihood (Laplace
##
    Approximation) [glmerMod]
## Family: poisson (log)
## Formula:
## cigarettes_A ~ gender_A + other_pos_A:gender_A + other_pos_P:gender_A -
##
       1 + (man + woman - 1 | cuplid)
##
     Data: acitelli_pair
##
##
                      logLik deviance df.resid
       AIC
                BIC
##
      898.1
              931.3
                      -440.1
                                 880.1
##
## Scaled residuals:
      Min
                1Q Median
                                3Q
                                      Max
## -1.2355 -0.4398 -0.1403 0.4725
                                   2.2843
##
## Random effects:
```

```
Groups Name Variance Std.Dev. Corr
##
   cuplid man
                0.1654
                         0.4067
##
##
          woman 0.1837
                         0.4286
                                  1.00
## Number of obs: 296, groups: cuplid, 148
## Fixed effects:
##
                           Estimate Std. Error z value Pr(>|z|)
## gender Ahus
                            0.58286
                                       0.84298
                                                 0.691
                                                          0.489
## gender Awife
                                       0.86511 - 0.454
                           -0.39262
                                                          0.650
## gender Ahus:other pos A
                            0.11710
                                       0.16773 0.698
                                                          0.485
## gender_Awife:other_pos_A 0.08651
                                       0.15230
                                                0.568
                                                          0.570
## gender Ahus:other pos P -0.18596
                                       0.14563 - 1.277
                                                          0.202
## gender Awife:other_pos_P 0.07131
                                       0.16943
                                                0.421
                                                          0.674
##
## Correlation of Fixed Effects:
              gndr_Ah gndr_Aw gndr_Ah:__A gndr_Aw:__A gndr_Ah:__P
##
## gender Awif 0.209
## gndr Ah: A -0.696 -0.137
## gndr_Aw:__A -0.120 -0.565
                              -0.051
## gndr Ah: P -0.544 -0.124
                              -0.216
                                           0.223
## gndr_Aw:__P -0.140 -0.673
                               0.209
                                          -0.223
                                                      -0.052
## convergence code: 0
## Model failed to converge with max|grad| = 0.00401621 (tol = 0.001, component 1)
```

Generalized Estimating Equations (GEE)

Indistinguishable Dyads

```
##
## GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA
## gee S-function, version 4.13 modified 98/01/27 (1998)
##
## Model:
## Link:
                              Logit
## Variance to Mean Relation: Binomial
## Correlation Structure:
                              Unstructured
##
## Call:
## gee(formula = simhob_bin_A ~ other_pos_A + other_pos_P, id = cuplid,
      data = acitelli pair, na.action = na.omit, family = binomial,
       corstr = "unstructured")
##
##
## Summary of Residuals:
          Min
                       1Q
                               Median
                                               3Q
                                                          Max
## -0.43310561 -0.27624377 -0.19297150 0.07522507 0.86606888
##
##
## Coefficients:
                Estimate Naive S.E. Naive z Robust S.E. Robust z
## (Intercept) -6.2729871 1.8140043 -3.458088 1.7415077 -3.602044
## other pos A 0.6982148 0.2901570 2.406334
                                                0.2851282 2.448775
## other pos P 0.5025443 0.2840654 1.769115 0.2605414 1.928846
##
## Estimated Scale Parameter:
                              0.9978716
## Number of Iterations: 2
##
## Working Correlation
             [,1]
                       [,2]
## [1,] 1.0000000 0.2703261
## [2,] 0.2703261 1.0000000
```