# Repeated Measures and Longitudinal Data Analysis II

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#### Welcome and outline - session 10

- Learning objectives
  - ▶ identify hierarchical data
  - define mixed effects models and population average models
  - perform model diagnostics for random effects models
  - interpret random intercepts and random slopes
- ▶ Vittinghoff sections 8.2, 8.3, 8.5

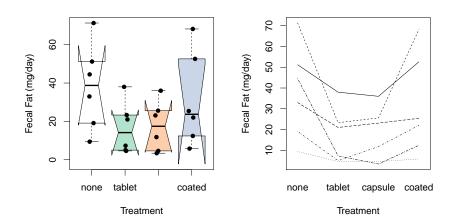
## Recall the simple repeated measures example: fecal fat

- ► Lack of digestive enzymes in the intestine can cause bowel absorption problems.
  - This will be indicated by excess fat in the feces.
  - ▶ Pancreatic enzyme supplements can alleviate the problem.
  - ► fecfat.csv: a study of fecal fat quantity (g/day) for individuals given each of a placebo and 3 types of pills

**Table 7.1** Fecal fat (g/day) for six subjects

Subject	Pill type	Subject			
number	None	Tablet	Capsule	Coated	Average
1	44.5	7.3	3.4	12.4	16.9
2	33.0	21.0	23.1	25.4	25.6
3	19.1	5.0	11.8	22.0	14.5
4	9.4	4.6	4.6	5.8	6.1
5	71.3	23.3	25.6	68.2	47.1
6	51.2	38.0	36.0	52.6	44.5
Pill type					
average	38.1	16.5	17.4	31.1	25.8

#### Fecal fat dataset



### Analysis strategies for hierarchical data

- Analyses for each subgroup
  - e.g., look at each patient independently
  - doesn't work at all in this example, and in general is not an integrated analysis of the whole data
  - could sort of work for an example with many patients per doctor, a few doctors
- Analysis at the highest level in the hierarchy
  - first summarize data to highest level
  - doesn't work at all in this example
  - could sort of work for an example with many patients per doctor, a few doctors
- Analysis on "Derived Variables"
  - consider each treatment type separately, take differences in fat levels between treatment/control for each patient and use paired t-tests
  - can work, but not work for unbalanced groups (e.g. incomplete data for some participants)

### Better analysis strategies for hierarchical data

- Fixed effects models
- Random / mixed effects models
  - model certain regression coefficients (intercept, slopes) as random variables
- ► Population average models
  - using Generalized Estimating Equations (GEE)

### When is hierarchical analysis needed?

- ► Hierarchical analysis strategies are needed:
  - 1. when the correlation structure is of primary interest, *e.g.* familial aggregation of disease, or consistency of treatment within centers,
  - 2. when we wish to "borrow strength" across the levels of a hierarchy in order to improve estimates, and
  - 3. when dealing with unbalanced, correlated data.

### Fixed effects: two-way analysis of variance

► Two-way ANOVA (person *i* with pill type *j*):

$$\textit{FECFAT}_{ij} = \beta_0 + \beta_{\textit{subjecti}} \textit{SUBJECT}_i + \beta_{\textit{pilltypej}} \textit{PILLTYPE}_j + \epsilon_{ij}$$

Assumption:  $\epsilon_i \stackrel{iid}{\sim} N(0, \sigma_{\epsilon}^2)$ 

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
subject	5	5588.38	1117.68	10.45	0.0002
pilltype	3	2008.60	669.53	6.26	0.0057
Residuals	15	1604.98	107.00		

Table 1: Two-way analysis of variance table. Equivalent to subtracting the mean fecal fat content for each individual.

\* Accounts for individual differences in mean fecal fat \* Fits a coefficient for mean fecal fat per individual

#### Mixed effects model

Model looks like two-way ANOVA:

$$\textit{FECFAT}_{ij} = \beta_0 + \beta_{\textit{subjecti}} \textit{SUBJECT}_i + \beta_{\textit{pilltypej}} \textit{PILLTYPE}_j + \epsilon_{ij}$$

- Assumption:  $\epsilon_i \stackrel{iid}{\sim} N(0, \sigma_{\epsilon}^2)$
- ▶ But instead of fitting a  $\beta$  to each individual, we assume that the subject effects are selected from a distribution of possible subject effects:

$$\textit{FECFAT}_{ij} = \beta_0 + \textit{SUBJECT}_i + \beta_{\textit{pilltypej}} \textit{PILLTYPE}_j + \epsilon_{ij}$$

Where we assume:  $SUBJECT_i \stackrel{iid}{\sim} N(0, \tau_{00}^2)$ 

- ▶ This is a *mixed effects* model because:
  - ▶ the "true" intercept varies randomly from patient to patient
  - the "true" (population) coefficient of treatment is fixed (the same for everyone)

### Mixed effects model coefficients, variances, ICC

```
## Linear mixed-effects model fit by REML
     Data: fecfat
   Log-restricted-likelihood: -84.55594
   Fixed: fecfat ~ pilltype
##
       (Intercept) pilltypetablet pilltypecapsule pilltypecoated
         38.083334
                        -21.550001
                                        -20.666667
##
                                                          -7.016668
##
## Random effects:
  Formula: ~1 | subject
           (Intercept) Residual
              15 89557 10 34403
## StdDev:
##
## Number of Observations: 24
## Number of Groups: 6
ICC = 15.9^{2}/(15.9^{2} + 10.34^{2}) = 0.7 = 0.7.
```

- Recall ICC is a measure of how large the subject effect is, in relation to the error term
- Variances were estimated directly by the model!

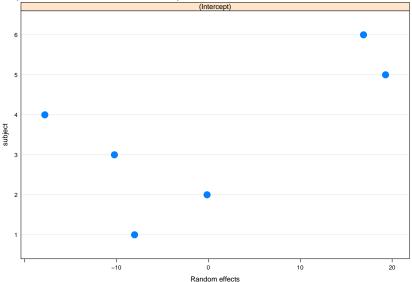
## Assumptions of the mixed model

$$\textit{FECFAT}_{ij} = \beta_0 + \textit{SUBJECT}_i + \beta_{\textit{pilltypej}} \textit{PILLTYPE}_j + \epsilon_{ij}$$

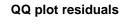
- Normally distributed residuals as in fixed effects model:
- Normally distributed latent variable:
  - SUBJECT<sub>i</sub>  $\stackrel{iid}{\sim} N(0, \tau_{00}^2)$

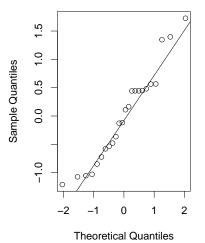
# Mixed effects model results (cont'd)

A plot of the random intercept:

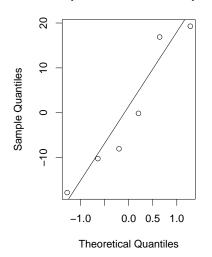


# Mixed effects model diagnostics





#### QQ plot random intercepts



#### Mixed effects model results

```
## Linear mixed-effects model fit by REML
   Data: fecfat
         ATC
                  BIC logLik
##
    181.1119 187.0863 -84.55594
##
## Random effects:
## Formula: ~1 | subject
          (Intercept) Residual
## StdDev: 15.89557 10.34403
##
## Fixed effects: fecfat ~ pilltype
                      Value Std.Error DF t-value p-value
##
## (Intercept) 38.08333 7.742396 15 4.918805 0.0002
## pilltypetablet -21.55000 5.972127 15 -3.608430 0.0026
## pilltypecapsule -20.66667 5.972127 15 -3.460521 0.0035
## pilltypecoated -7.01667 5.972127 15 -1.174903 0.2583
## Correlation:
                  (Intr) plltypt plltypcp
##
## pilltypetablet -0.386
## pilltypecapsule -0.386 0.500
## pilltypecoated -0.386 0.500 0.500
##
## Standardized Within-Group Residuals:
           Min
## -1.210052934 -0.615068039 -0.002727166 0.457105344 1.725618643
##
## Number of Observations: 24
## Number of Groups: 6
```

- Note: correlation of the estimator of the fixed effects
  - ▶ high correlations may (but not necessarily) be due to collinearity

### Mixed effects model results (cont'd)

Inference for variance terms (and fixed effects):

```
## Approximate 95% confidence intervals
##
   Fixed effects:
##
                       lower
                                            upper
## (Intercept)
                  21.58081 38.083334 54.585860
## pilltypetablet -34.27929 -21.550001 -8.820714
## pilltypecapsule -33.39595 -20.666667 -7.937381
## pilltypecoated -19.74595 -7.016668 5.712618
## attr(,"label")
## [1] "Fixed effects:"
##
   Random Effects:
    Level: subject
##
                     lower
                              est.
                                        upper
## sd((Intercept)) 8.001171 15.89557 31.57904
##
   Within-group standard error:
##
       lower
  7.232404 10.344027 14.794374
```

- Would conclude that variation of the intercept between subjects is non-zero
  - not attributable to within-subject variation

### Longitudinal data

- Interested in the change in the value of a variable within a "subject"
- Collect data repeatedly through time.
- For hierarchical longitudinal analysis to be effective,
   before/after measurements need to be positively correlated

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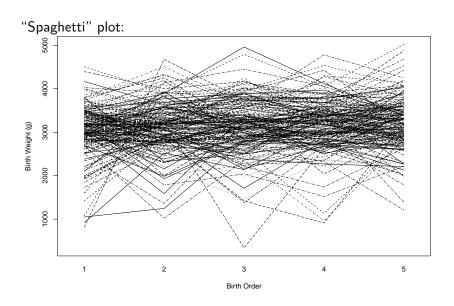
### Longitudinal data examples

- ► Example 1: a measure of sleepiness before and after administration of treatment or placebo
- ► Example 2: Study of Osteoporotic Fractores (SOF dataset)
  - ▶ 9,704 women tracked with clinical visits every two years
  - Bone Mineral Density (BMD), Body Mass Index (BMI), many other variables
- Questions for Example 2:
  - Is change in BMD related to age at menopause? This is a time-invariant predictor, age at menopause, with time-dependent changes in the outcome, BMD.
  - 2. Is change in BMD related to change in BMI? This is an analysis relating a time-varying predictor, BMI, with changes in the outcome, BMD. BMI varies quite a lot between women, but also varies within a woman over time.

# Longitudinal data examples (cont'd)

- birthweight and birth order
- provides birthweights and order of infants from mothers who had 5 children in Georgia
  - interested in whether birthweight of babies changes with order
  - whether this difference depends on the mother's age at first childbirth or on the weight of initial baby.

## Georgia Birthweights dataset

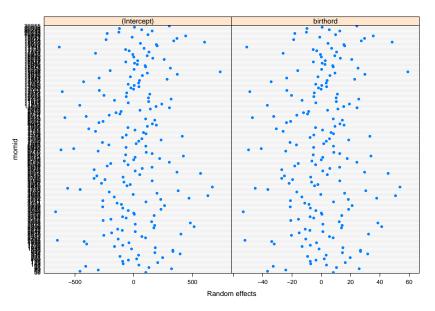


### Georgia Birthweights dataset

- ▶ Do birth weights or the effect of birth order vary by mother?
  - random intercept, random slope

Note: the control argument increases the max # iterations to allow convergence. If msVerbose=TRUE, it produces verbose output.

# Georgia Birthweights dataset (cont'd)



## Georgia Birthweights dataset (cont'd)

summary(gafit1)

```
## Linear mixed-effects model fit by REML
## Data: ga
##
          ATC
                  BIC
                        logLik
   15321.23 15350.67 -7654.616
##
## Random effects:
## Formula: ~birthord | momid
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                        Corr
## (Intercept) 296.36304 (Intr)
## birthord 23,82664 0.996
## Residual
              443 43313
##
## Fixed effects: bweight ~ birthord
##
                 Value Std.Error DF t-value p-value
## (Intercept) 2995.640 38.99534 799 76.82046
## birthord
                46.608 10.05758 799 4.63411
## Correlation:
           (Intr)
##
## birthord -0.662
##
## Standardized Within-Group Residuals:
##
           Min
                       Q1
                                  Med
                                                          Max
## -5.32850276 -0.42195082 0.04103012 0.53113344 3.25512201
##
## Number of Observations: 1000
## Number of Groups: 200
```

## Georgia Birthweights dataset (cont'd)

```
intervals(gafit1)
```

```
## Approximate 95% confidence intervals
##
   Fixed effects:
##
                    lower
                              est.
                                        upper
## (Intercept) 2919.09459 2995.640 3072.18541
## birthord
                 26.86559
                            46.608
                                     66.35041
## attr(,"label")
## [1] "Fixed effects:"
##
   Random Effects:
   Level: momid
##
                                  lower
                                                         upper
## sd((Intercept))
                             227.421283 296.3630360 386.20418
## sd(birthord)
                               9.351472 23.8266383 60.70795
## cor((Intercept),birthord) -1.000000 0.9964047 1.00000
##
   Within-group standard error:
      lower
                est.
                        upper
## 422 2138 443 4331 465 7189
```

- Do birth weights or the effect of birth order vary by mother?
  - yes: both standard deviations are non-zero

### Population Average Models

- ► An alternative to random / mixed-effects models that is more robust to assumptions of:
  - distribution of random effects
  - correlation structure
- Estimates correlation structure from the data rather than assuming normality
  - Requires a fair bit more clusters than observations per cluster
- Estimates regression coefficients and robust standard errors
  - commonly by Generalized Estimating Equations (GEE)

## Population Average Models

Compare mixed model multiple linear regression:

$$E[Y_{ij}|X_{ij}] = \beta_0 + \alpha_{0j} + \beta_1 X_{ij}, \alpha_{0j} \sim N(0, \sigma)$$

for subject i in group j.

to a population average model:

$$E[Y_{ij}|X_{ij}] = \beta_0^* + \beta_1^* X_{ij}$$

- ▶ Interpretations of  $\beta^*$  and  $\beta$  are equivalent
- Numerically equivalent for linear and log-linear models (if specification of mixed model is correct), but not for logistic link.

### Fit a population average model

```
gafit.gee <- gee::gee(bweight ~ birthord, corstr="exchangeable",
                     id=momid, data=ga)
summary(gafit.gee)
##
## GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA
   gee S-function, version 4.13 modified 98/01/27 (1998)
##
## Model:
## Link:
                              Identity
## Variance to Mean Relation: Gaussian
## Correlation Structure:
                              Exchangeable
##
## Call:
## gee::gee(formula = bweight ~ birthord, id = momid, data = ga,
       corstr = "exchangeable")
##
##
## Summary of Residuals:
##
        Min
                         Median
                                       30
                                                Max
                   10
## -2795 464 -299 126 48 840 341 144 1824 536
##
##
## Coefficients:
              Estimate Naive S.E. Naive z Robust S.E. Robust z
## (Intercept) 2995.640 41.973695 71.369462 38.808066 77.191170
## hirthord 46 608 9 958128 4 680398 9 996256 4 662546
##
## Estimated Scale Parameter: 332525.3
## Number of Iterations: 1
##
## Working Correlation
##
            [.1]
                      [,2]
                                Γ.31
                                          Γ.47
                                                    Γ.51
## [1.] 1.0000000 0.4035684 0.4035684 0.4035684 0.4035684
```

## GEE working correlation types

- Must make some assumption about form correlation among observations from the same subject, same hospital, etc
- Independence
  - no correlation between measurements within group
- Exchangeable
  - all correlations (except those variables with themselves) are a common value
  - nothing to distinguish one member of a cluster from another
  - appropriate in the absence of other data structures such as measurements taken through time or space
- Auto-regressive
  - observations taken more closely in time are more highly correlated

### GEE working correlation types

- Unstructured
  - estimates a separate correlation between observations taken on each pair of "times"
- Non-stationary
  - similar to unstructured, but assumes all correlations for pairs separated far enough in time are zero
- Stationary
  - e.g. stationary of order 2: observations taken at time points 1 and 3 have the same correlation as time points 2 and 4
  - but this might be different from the correlation between observations taken at times 2 and 3
  - correlations for observations 3 or more time periods apart assumed to be zero

Fewer assumptions requires more data, and good assumptions improve results

# Help in choosing a method

Characteristic	Marginal	Fixed-effect	Mixed-effect
Distinguishes observations belonging to the same or different subjects	Yes <sup>a</sup>	Yes	Yes
Reliant on distribution of subject-specific effects	No	No	Yes
Subjects considered a sample from a population larger than the sample itself	Yes <sup>a</sup>	No	Yes
Computation handles few subjects well	No	Yes	No
Computation handles a very large number of subjects well	Yes	No	Yes
Noisy for few observations per subject	No	Yes	No
Computation handles a large number of observations per subject	Depends <sup>b</sup>	Yes	Yes
Accommodates variable observations per subject	Yes	Yes	Yes

Note: aOnly for calculation of standard errors.

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Figure 2: decision table

<sup>&</sup>lt;sup>b</sup>Problems can arise under some specifications of the working covariance structure and depending on the estimation method used.

#### Conclusions

- Ignoring within-subject correlations can produce very wrong results, and is not always "conservative"
- Hierarchical analysis strategies are needed for any of:
  - When the correlation structure is of primary interest, e.g. familial aggregation of disease, or consistency of treatment within centers,
  - 2. When we wish to "borrow strength" across the levels of a hierarchy in order to improve estimates, and
  - 3. When dealing with unbalanced correlated data. E.g., no requirement that each Georgia mother have exactly 5 children.
- Population average models provide a robust alternative to mixed models
  - for one level of hierarchy

#### Lab exercise

- Continue the MLM\_R.pdf lab from session 9
- ► Fit population average models using several different working correlation structures, and compare the results to each other and to the mixed model