STATLAB WORKSHOP

MIXED EFFECTS MODELING WITH R

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OUTLINE

1 Theory

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- 2 Examples
 - Repeated Measurements
 - Clustered data from random lines
 - Clustering and repeated measurements
 - Longitudinal growth

THEORY

The general formulation of the multiple linear model with iid Normal errors:

$$Y_i = \beta_0 + \beta_1 X_i^{(1)} + \ldots + \beta_d X_i^{(d)} + \epsilon_i$$

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Inference tasks based on the least-squared estimators are straightforward.

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$$Y_i = \beta_0 + \beta_1 x_i^{(1)} + \ldots + \beta_d x_i^{(d)} + b_1 z_i^{(1)} + \ldots + b_m z_i^{(m)} + \epsilon_i$$

with $\epsilon_1, \ldots, \epsilon_n$ iid Normal and the random effects (b_1, \ldots, b_m) are multivariate Normal. (Note x_1, \ldots, x_d and z_1, \ldots, z_m are all explanatory variables.)

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The errors don't actually have to be independent of each other. Our final example will demonstrate this.

EXAMPLES

BLOOD PRESSURE MEASUREMENTS

Suppose that n_f females and n_m males each had the blood pressures measured. Assume that the distributions of female and male blood pressures are both Normal with the same standard deviation σ . Let their expectations be denoted μ_f and μ_m respectively. With x_1,\ldots,x_n representing sex (the only explanatory variable) and Y_1,\ldots,Y_N representing blood pressures, we can express this scenario with the linear model

$$Y_i = \mu_f + (\mu_m - \mu_f) \mathbb{I}(X_i = "M") + \epsilon_i.$$

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Inference for the means fits into the usual linear modeling framework. (In this case, it simplifies to an ordinary two-sample *t*-test.)

REPEATED BLOOD PRESSURE MEASUREMENTS

Suppose alternatively that each of the subjects' blood pressures were measured up to four times, at random moments over the course of a week. The measurements of a single person's blood pressure can vary quite a bit when measured at different times. How could we adapt our model to capture this?

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We might assume that each subject has an *expected* blood pressure and model the specific measurements as iid Normal draws centered at that expectation. Assume further that each subject's distribution of blood pressure measurements shares the same standard deviation σ . Let b_i denote the deviation of subject i from his or her group's mean μ_{x_i} . Assume that these deviations are iid Normal with mean zero and standard deviation σ_s .

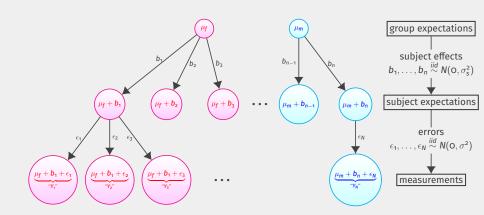
MODEL

Let $n:=n_f+n_m$ be the total number of subjects. Our assumptions fit into the mixed effects modeling framework if you define $z_i^{(1)},\ldots,z_i^{(m)}$ to be the indicator variables that tell us which subject observation j came from. Specifically, $b_i^{(j)}$ is 1 if measurement i is from subject j and it's 0 otherwise. To see how the model equation simplifies, suppose that observation i is from subject j who is a male. Then

$$Y_{i} = \mu_{f} + (\mu_{m} - \mu_{f})\mathbb{I}(x_{i} = "M") + b_{1}z_{i}^{(1)} + \ldots + b_{m}z_{i}^{(m)} + \epsilon_{i}$$

= $\mu_{m} + b_{j} + \epsilon_{i}$.

DIAGRAM



R SCRIPT EXAMPLE 1

Now, let's turn to the first portion of *mixed-effects.R*, where we'll simulate blood pressure data according to the repeated measurements mechanism just described. Then we'll use the *lme4* package to fit the corresponding mixed effects model, and compare the true values to the resulting estimates. Finally, we'll use the *lmertest* package for inference and model selection.

SAT SCORES AT VARIOUS SCHOOLS

Suppose that, at any given school, there's a line relating parents' log income (explanatory variable) and expected SAT score (response variable). The students' actual SAT scores deviate from the line by Normal errors with a common variance σ^2 . Suppose, however, that each school has its own line, that is, its own intercept and slope. Let's assume that the schools' lines are iid draws from a bivariate Normal distribution centered at $(\mu_{\rm int}, \mu_{\rm slope})$.

MODEL

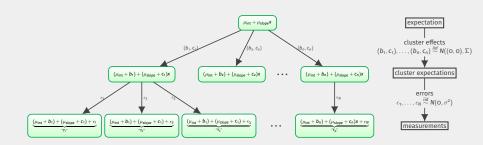
This scenario fits into the linear mixed effects modeling framework described above. Let Y_i and x_i represent the ith student's SAT score and parents' log income. Let $z_i^{(j)}$ be the indicator that student i attends school j. Letting student i attend school k (i.e. $z_i^{(k)} = 1$ and the others are 0), and letting m denote the number of schools in the dataset,

$$Y_{i} = \mu_{\text{int}} + \mu_{\text{slope}} X_{i} + b_{1} Z_{i}^{(1)} + \ldots + b_{m} Z_{i}^{(m)} + c_{1} Z_{i}^{(1)} X_{i} + \ldots + c_{m} Z_{i}^{(m)} X_{i} + \epsilon_{i}$$

$$= \underbrace{(\mu_{\text{int}} + b_{k})}_{\text{school } k' \text{s intercept}} + \underbrace{(\mu_{\text{slope}} + c_{k})}_{\text{school } k' \text{s slope}} X_{i} + \epsilon_{i}$$

where $\epsilon_1, \ldots, \epsilon_n$ are iid $N(0, \sigma^2)$. The Normal deviations (b_j, c_j) of the *j*th school's intercept and slope from the expectations shouldn't be assumed to be independent of each other.

DIAGRAM



R SCRIPT EXAMPLE 2

Now, let's turn to the second portion of *mixed-effects.R*, where we'll simulate SAT scores according to the random lines mechanism just described. Then we'll again fit the corresponding mixed effects model, compare the true values to the resulting estimates, and engage in inference and model selection.

Suppose that the average female SAT score is $\mu_f =$ 1055 and the average male SAT score is $\mu_m =$ 1042.

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Suppose that each school has its own effect on SAT score and that the school's expected males' scores is the same as the effect on female scores. In other words, if a school's effect is b_j , then it's expected female score is $\mu_f + b_j$ and its expected male score is $\mu_m + b_j$. Assume that the effects b_1, b_2, \ldots are iid $N(o, \sigma_{\text{school}}^2)$.

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There are also student effects c_1, c_2, \ldots Suppose that student k is a male who attends school j. His expected SAT score is $\mu_m + b_j + c_k$. Assume the student effects are iid $N(o, \sigma_{\text{student}}^2)$.

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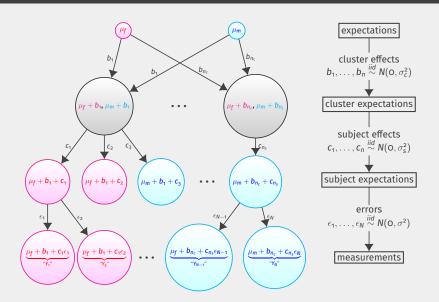
scores differ from the expectation by a Normal error. Assume the errors are iid $N(o, \sigma^2)$.

Model

We can represent this scenario as a mixed effects model with Y_i representing the ith SAT score, $z_i^{(1)}, z_i^{(2)}, \ldots$ indicators representing the student tested, $w_i^{(1)}, w_i^{(2)}, \ldots$ representing that student's school, and x_i representing that student's sex. If Y_i is a test score of student k who is a male at school j,

$$Y_{i} = \mu_{f} + (\mu_{m} - \mu_{f})\mathbb{I}(x_{i} = \text{"M"}) + b_{1}w_{i}^{(1)} + b_{2}w_{i}^{(2)} + \ldots + c_{1}z_{i}^{(1)} + c_{2}z_{i}^{(2)} - \omega_{m} + b_{j} + c_{k} + \epsilon_{i}.$$

DIAGRAM



R SCRIPT EXAMPLE 3

Now, we'll work through the third part of *mixed-effects.R*, where we'll simulate SAT scores according to the mechanism just described: first, generate schools, then generate students who attend those schools, then generate SAT test scores for those students. Then we'll again fit the corresponding mixed effects model, compare the true values to the resulting estimates, and engage in inference and model selection.

MEASUREMENTS OF TREES OVER TIME WITH DIFFERENT FERTILIZERS

Pretend that three different fertilizers are being studied. For each fertilizer, a number of trees will be planted and their heights will be measured after 1 year, 2 years, 3 years, and 4 years. Assume that for each fertilizer, the expected amount growth (increased height) is the same every year for the first four years. A tree with fertilizer A is expected to grow $\mu_A = 2.7$ feet per year, with B it's expected to grow 3 feet per year, and with C it's expected to grow 3.6 feet per year.

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Any particular tree will have its own effect. In particular, suppose it's expected growth per year deviates from the treatment group's expectation by some Normal draw. Furthermore, during any particular year, this tree's growth will differ from its expected growth by a Normal draw. This is a special type of repeated measurements scenario in which the timing of the measurements is important; it's called "longitudinal data."

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Any particular tree will have its own effect. In particular, suppose it's expected growth per year deviates from the treatment group's expectation by some Normal draw. Furthermore, during any particular year, this tree's growth will differ from its expected growth by a Normal draw. This is a special type of repeated measurements scenario in which the timing of the measurements is important; it's called "longitudinal data."

Why does that matter? Each tree has a sequence of deviations from its expected line. If a tree grows more than it was expected to during the first year, then most likely it will still be taller than its expectation after the second year. With longitudinal data, we typically don't assume that an individual's repeated errors (deviations from the expectation) are independent of each other; dependence among neighboring errors is called *autocorrelation*.

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MODEL

We can represent this scenario as a mixed effects model; this time we'll use double subscripts to index the observations. Let $Y_{i,j}$ represent the jth measurement of tree i (which is the height of tree i after j years), $z_i^{(1)}, z_i^{(2)}, \ldots$ indicators representing the tree tested, and b_i representing the ith tree's deviation from the expected growth rate of its treatment group. Assume b_1, b_2, \ldots are iid $N(0, \sigma_t^2)$. If, for example, tree i got fertilizer B, then

$$\begin{aligned} Y_{i,j} &= \eta_{\mathsf{A}} j + (\eta_{\mathsf{B}} - \eta_{\mathsf{A}}) \mathbb{I}(\mathsf{X}_i = \text{``B''}) j + (\mu_{\mathsf{C}} - \mu_{\mathsf{A}}) \mathbb{I}(\mathsf{X}_i = \text{``C''}) j \\ &+ b_1 z_i^{(1)} j + b_2 z_i^{(2)} j + \ldots + \epsilon_{i,j} \\ &= \underbrace{(\mu_{\mathsf{B}} + b_i)}_{\mathsf{Slope of the tree } i} j + \epsilon_{i,j}. \end{aligned}$$

Recall that $\epsilon_{i,1}, \epsilon_{i,2}, \ldots$ aren't independent of each other. Rather each year there is an independent Normal deviation from the expected amount of growth, and $\epsilon_{i,j}$ is the sum of draw 1 through draw j.

MODEL

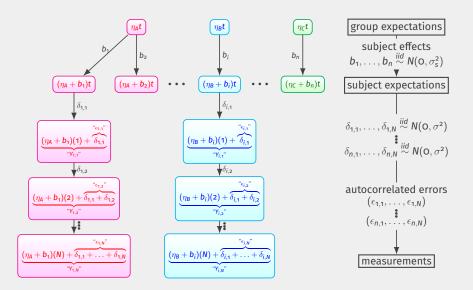
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(We know that all trees had a height of zero at time zero, so I've built that into the model by not including an intercept term.)

DIAGRAM



R SCRIPT EXAMPLE 4

Now, we'll work through the fourth and final part of *mixed-effects.R*, where we'll simulate SAT scores according to the mechanism just described: first, generate trees, then generate a sequence of heights for those trees. We'll again fit the corresponding mixed effects model, compare the true values to the resulting estimates, and engage in inference and model selection. Following this example, you'll find an exercise to attempt on your own. My solution is provided in *mixed-effects-solutions.R*.