# Chapters 25 & 27: Simple random effects and mixed effects models

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Stat 705: Data Analysis II

#### 25.1 One-way random cell means model

If treatment levels come from a larger population, their effects are best modeled as random. A one-way random cell means model is

$$Y_{ij} = \mu_i + \epsilon_{ij},$$

where

$$\mu_1, \ldots, \mu_r \stackrel{\textit{iid}}{\sim} N(\mu_{\cdot}, \sigma_{\mu}^2) \text{ independent of } \epsilon_{ij} \stackrel{\textit{iid}}{\sim} N(0, \sigma^2).$$

As usual, 
$$i = 1, \ldots, r$$
 and  $j = 1, \ldots, n_i$ .

The test of interest is  $H_0: \sigma_{\mu}^2 = 0.$ 

We can re-express the model as a random effects model, by writing  $\mu_i = \mu + \tau_i$ , where  $\tau_1, \ldots, \tau_r \stackrel{iid}{\sim} N(0, \sigma_{\mu}^2)$ .

 $\tau_1, \ldots, \tau_r$  are called *random effects* and  $\sigma_{\mu}^2$  and  $\sigma^2$  are termed *variance components*. This model is an example of a *random effects* model, because it has only random effects beyond the intercept  $\mu$ . (which is fixed).

The random cell means model has some quite different properties from the fixed cell means model.

• 
$$E(Y_{ij}) = \mu$$
.  
•  $\sigma^2 \{Y_{ij}\} = \sigma^2 + \sigma_{\mu}^2$  (Hence the term variance components)  
•  $\sigma \{Y_{ij}, Y_{ij'}\} = \sigma_{\mu}^2$   
•  $\rho \{Y_{ij}, Y_{ij'}\} = \frac{\sigma_{\mu}^2}{\sigma^2 + \sigma_{\mu}^2}$   
•  $E(\bar{Y}_{..}) = \mu$ .  
•  $\sigma^2(\bar{Y}_{..}) = \frac{\sigma^2 + n\sigma_{\mu}^2}{n}$ 

The MSE and MSTR are defined as they were before. One can show  $E(MSE) = \sigma^2$  and  $E(MSTR) = \sigma^2 + n\sigma_{\mu}^2$  when  $n = n_i$  for all *i*. Most packages provides symbolic forms of expected mean squares for random/mixed models if requested.

If  $\sigma_{\mu} = 0$  we expect  $F^* = MSTR/MSE$  to be somewhat larger than 1. In fact, just like the fixed-effects case,  $F^* \sim F(r-1, n_T - r)$ . This is the test given by proc glm when you add a random A; statement.

One can also fit the model in proc mixed, but this procedure provides a slightly cruder test of  $H_0: \sigma_{\mu} = 0$ .

We can derive estimates for  $\mu$ .,  $\sigma^2$  and  $\frac{\sigma_{\mu}^2}{\sigma^2 + \sigma_{\mu}^2}$  because pivotal quantities are readily available. It is an open question whether we are interested in inference on  $\mu$ . in most practical applications.

Other quantities of interest tended to require moment-based estimates (old school)–e.g., the variance component  $\sigma_{\mu}^2$ . Methods to provide point estimates and/or standard errors include

- Satterthwaite Procedure (still old school)
- Modified Large Sample (still old school)
- Maximum Likelihood (biased)
- Restricted Maximum Likelihood

## Musical competition

r = 4 judges were selected at random to review students' musical performances on trumpet. Four students were randomly selected to have their performances reviewed by a given judge.  $Y_{ij}$  is the rating of the *i*th judge on the *j*th student.

Since the judges are chosen randomly from a large population of available judges, the random-effects one-way model applies.

```
data music;
input rating judge @@;
datalines;
    76 1 65 1 85 1 74 1 59 2 75 2 81 2 67 2
    49 3 63 3 61 3 46 3 74 4 71 4 85 4 89 4
;
run;
proc glm; class judge; * Chapter 16, fixed-effects approach;
model rating=judge; run;
proc glm; class judge; * Chapter 25, mixed-effects approach;
model rating=judge;
random judge; run;
proc glimmix; class judge; * Tim prefers method=mle;
model rating= / s cl;
random judge / g cl;
covtest zerog; * tests H0: sigma_mu=0 vs. H0: sigma_mu>0;
```

In the two-factor random effects model, we specify

$$Y_{ijk} = \mu_{\cdot\cdot} + \alpha_i + \beta_j + (\alpha\beta)_{ij} + \epsilon_{ijk}$$

where  $\mu_{..}$  is a constant,  $\alpha_i \stackrel{iid}{\sim} N(0, \sigma_{\alpha}^2), \beta_j \stackrel{iid}{\sim} N(0, \sigma_{\beta}^2), (\alpha\beta)_{ij} \stackrel{iid}{\sim} N(0, \sigma_{\alpha\beta}^2)$  and  $\epsilon_{ijk} \stackrel{iid}{\sim} N(0, \sigma^2)$  are pairwise independent.

As mentioned in class, these models can be useful when studying sources of measurement error, e.g., in an industrial R & R study (Repeatability and Reproducibility–I'll bet those weren't your first two guesses).

The Expected Mean Squares suggests a different set of F tests to test variance components.

<u>Source</u>	<u>df</u>	EMS
А	a-1	$\sigma^2 + n\sigma_{\alpha\beta}^2 + nb\sigma_{\alpha}^2$
В	b-1	$\sigma^2 + n\sigma^2_{\alpha\beta} + na\sigma^2_{\beta}$
AB	(a-1)(b-1)	$\sigma^2 + n\sigma^2_{\alpha\beta}$
Error	ab(n-1)	$\sigma^2$
Total	abn-1	

On-line prices for several drugs were checked at multiple local pharmacies. Is this a random effects two-factor study? Are model assumptions satisfied?

	Medicine								
	Walmart	Mart	Target	Longs	Agape	CVS			
Lasix	29.76	31.52	31.52	31.71	32.22	31.92			
Nardil	18.50	23.13	24.54	23.23	24.89	17.62			
Procardia	23.25	25.53	32.88	25.53	29.62	23.72			
Naprosyn	187.58	184.75	185.81	185.45	185.45	188.90			
Danazol	139.39	135.30	148.19	135.40	251.02	153.30			
Lidocaine	239.75	157.24	216.36	157.14	157.14	145.07			

# 25.2 Drug case study

```
data pharmacy;
length Drug $10;
input Drug $ Pharmacy $ Price;
datalines;
Lasix Walmart 29.76 Lasix MedMart 31.52 & Lasix Target 31.52 Lasix Longs 31.71
Lasix Agape 32.22 Lasix CVS 31.92
Lidocaine Walmart 239.75 Lidocaine MedMart 157.24 Lidocaine Target 216.36
Lidocaine Longs 157.14 Lidocaine Agape 157.14 Lidocaine CVS 145.07
;
run;
proc glm data=pharmacy; class drug pharmacy;
model price=drug pharmacy;
random drug pharmacv/test; run;
proc glimmix data=pharmacy; class drug pharmacy;
model price= / s cl:
random drug pharmacy / g cl;
covtest zerog;
 run;
```

The book focuses on a *restricted* version of the factor effects model. This approach is rarely used any more, perhaps because an intellectual inconsistency–it is never generalized to models with more fixed and random effects.

$$Y_{ijk} = \mu_{..} + \alpha_i + \beta_j + (\alpha\beta)_{ij} + \epsilon_{ijk}$$

where  $\mu_{\cdots}$  is a constant,  $\alpha_i$  are fixed effects,  $\beta_j \stackrel{iid}{\sim} N(0, \sigma_{\beta}^2), (\alpha\beta)_{ij} \stackrel{iid}{\sim} N(0, \sigma_{\alpha\beta}^2)$  and  $\epsilon_{ijk} \stackrel{iid}{\sim} N(0, \sigma^2)$  are pairwise independent. The Expected Mean Squares for the unrestricted model uses MSAB for the error term for both random and fixed effects. This has important consequences for analysis of the fixed effect.



In the restricted model, the error term for the random effect B is MSE!

Inference on the marginal mean  $\mu_i$  is more difficult since there is no ready estimate for the variance of  $\hat{\mu}_i = \bar{Y}_i$ ... However, inference on the fixed effect  $\alpha_i$  is straightforward. We can show  $\hat{\alpha}_i = \bar{Y}_{i..} - \bar{Y}_{...}$  has variance  $\sigma^2(\hat{\alpha}_i) = \frac{\sigma^2 + n\sigma_{\alpha\beta}^2}{bn}$ , which is readily estimated by MSAB/bn. This result leads to ready results for LSMEANS and pairwise differences, since  $\hat{\mu}_i - \hat{\mu}_{i'} = \hat{\alpha}_i - \hat{\alpha}_{i'}$ . When block levels come from a large population, we can consider a complete randomized block design with random block effects. One very important example of this is the repeated measures design, where each block is an experimental unit in which all treatment levels are randomly applied. In fact, the blocks are retermed "subjects" and we consider a sample of subjects from their population.

$$Y_{ij} = \mu_{\cdot\cdot} + \underbrace{\rho_i}_{\text{subject}} + \underbrace{\tau_j}_{\text{level}} + \epsilon_{ij},$$

where

$$ho_1,\ldots,
ho_{n_b}\stackrel{iid}{\sim} N(0,\sigma_
ho^2)$$
 independent of  $\epsilon_{ij}\stackrel{iid}{\sim} N(0,\sigma^2).$ 

There are  $i = 1, ..., n_b$  subjects receiving each of j = 1, ..., r treatments.

Note that the model can be extended to factorial treatment structure, e.g.

$$Y_{ijk} = \mu_{..} + \rho_i + \alpha_j + \beta_k + (\alpha\beta)_{jk} + \epsilon_{ijk},$$

Examples of subjects include people, animals, families, cities, and clinics.

This is an example of a mixed effects model; there is a mix of random ( $\rho_i$ 's) and fixed ( $\alpha_j$ 's,  $\beta_k$ 's, and ( $\alpha\beta$ )<sub>*jk*</sub>'s) effects in the model.

# Random blocks, comments

- We assume subject effects and treatment effects do not interact. We can check via Tukey's 1 df test for additivity. Also look at interaction plots as in fixed-effects RCBD designs.
- ANOVA table and sums of squares are exactly the same, except now the F-test for blocks tests H<sub>0</sub> : σ<sub>ρ</sub> = 0 instead of H<sub>0</sub> : ρ<sub>1</sub> = ··· = ρ<sub>n<sub>b</sub></sub> = 0.
- Test for treatment is same  $H_0: \tau_1 = \cdots = \tau_r = 0$ .
- Every treatment is given to every experimental unit in *randomized order*.
- Two sets of residuals to consider. Both should be normal; *e<sub>ij</sub>* should have constant variance.

**1** 
$$e_{ij} = Y_{ij} - {\hat{\mu} + \hat{\rho}_i + \hat{\tau}_j}$$
, and  
**2**  $\hat{\rho}_i$ .

- corr $(Y_{ij_1}, Y_{ij_2}) = \sigma_{\rho}^2/(\sigma^2 + \sigma_{\rho}^2)$  for  $j_1 \neq j_2$  tells you how correlated the repeated measures are.
- We will use proc glimmix to fit these models.

# Road paint wear (p. 1082)

These are problems 25.19 and 25.20.

A state highway department studied wear of five paints at eight randomly picked locations. The standard is paint 1. Paints 1, 3, and 5 are white; paints 2 and 4 are yellow. At each location a random ordering of the paints were applied to the road. After an exposure period, a combined measure of wear  $Y_{ij}$  was recorded. The higher the score, the better the wearing characteristics.

Recall the model

$$Y_{ij} = \mu + \underbrace{\rho_i}_{\text{location}} + \underbrace{\tau_j}_{\text{paint}} + \epsilon_{ij},$$

where

$$\rho_1, \ldots, \rho_{n_b} \stackrel{iid}{\sim} N(0, \sigma_{\rho}^2) \text{ independent of } \epsilon_{ij} \stackrel{iid}{\sim} N(0, \sigma^2).$$
Here  $r = 5$  and  $n = 8$ .

#### Road paint wear in SAS

data road;														
input wea	r locat	ion pa	aint @@;											
datalines;														
11.0	1	1	13.0	1	2	10.0	1	3	18.0	1	4	15.0	1	5
20.0	2	1	28.0	2	2	15.0	2	3	30.0	2	4	18.0	2	5
8.0	3	1	10.0	3	2	8.0	3	3	16.0	3	4	12.0	3	5
30.0	4	1	35.0	4	2	27.0	4	3	41.0	4	4	28.0	4	5
14.0	5	1	16.0	5	2	13.0	5	3	22.0	5	4	16.0	5	5
25.0	6	1	27.0	6	2	26.0	6	3	33.0	6	4	25.0	6	5
43.0	7	1	46.0	7	2	41.0	7	3	55.0	7	4	42.0	7	5
13.0	8	1	14.0	8	2	12.0	8	3	20.0	8	4	13.0	8	5
:														

proc glm plots=all; class location paint; \* interaction plot to check additivity; model wear=location|paint;

r = 4 Chardonnary wines of the same vintage were judged by n = 6 judges. Each wine was blinded and given to each judge in randomized order. The wines were scored on a 40-point scale  $Y_{ij}$ , with higher scores meaning better wine.

The six judges are considered to come from a large population of wine-tasting judges and so a repeated measures model is appropriate.

The analysis of these data are carried out in your textbook on pp. 1132–1137.

## Wine tasting in SAS proc glm

```
data wine:
input rating judge wine @@;
datalines:
 20 1 1 24 1 2 28 1 3 28 1 4 15 2 1 18 2 2 23 2 3 24 2 4
 18 3 1 19 3 2 24 3 3 23 3 4 26 4 1 26 4 2 30 4 3 30 4 4
 22 5 1 24 5 2 28 5 3 26 5 4 19 6 1 21 6 2 27 6 3 25 6 4
* spaghetti plot figure 27.2 on p. 1133;
proc sgplot noautolegend;
series x=wine y=rating / group=judge;
scatter x=wine y=rating / group=judge markerchar=judge;
 run;
* glm works, but is not really designed for repeated measures:
* this duplicates what is in your book;
proc glm plots=all; * gives figure 27.3 on p. 1133;
class wine judge;
model rating=wine judge;
random judge; * need to include 'judge' in model using glm;
lsmeans wine / pdiff adjust=tukev alpha=0.05 cl;
 run:
```

### Analysis in proc glimmix

```
* proc mixed or proc glimmix is a better choice overall;
* note that Tukey intervals are essentially the same;
* conditional residuals are r_ij;
proc glimmix plots=all;
class wine judge:
model rating=wine / s chisg: * model includes only 'fixed' effects:
                              * random includes only 'random' effects;
random judge;
lsmeans wine / pdiff adjust=tukev alpha=0.05;
covtest zerog; * tests H0: sigma_rho=0 vs. H0: sigma_rho>0;
 run;
* obtain estimates of rho i:
ods listing close; ods output SolutionR=rand; * sends the rho_i to 'rand';
proc glimmix data=wine;
class wine judge;
model rating=wine;
random judge / s; * ask for rho_i;
run:
ods output close: ods listing:
* check that rho_i estimates are approximately normal;
proc print data=rand; run;
proc univariate data=rand normal; var estimate;
run:
```