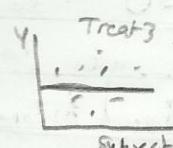
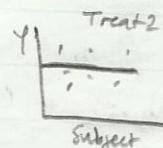
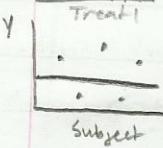
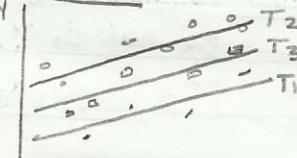


Chapter 22 ANCOVA

ANOVA



ANCOVA



* Reduced Variability

Concomitant variable added (unaffected by treatments) pre study

n_i = number of cases in the i^{th} level

$$n_T = \sum n_i$$

$$Y_{ij} = \mu_0 + T_i + \epsilon_{ij} \quad \text{ANOVA} \quad \leftarrow \text{In text, we fit it differently!}$$

$$Y_{ij} = \mu_0 + T_i + \gamma(X_{ij} - \bar{X}_{..}) + \epsilon_{ij} \downarrow \mu_0 + T_i I_{ij1} + T_2 I_{ij2} \dots + \gamma X_{ij} + \epsilon_{ij} \quad \text{ANCOVA}$$

μ_0 = overall mean

$$I_{ij1} = \frac{1}{\sum \text{treat1}} \quad I_{ij2} = \frac{1}{\sum \text{treat2}}$$

$$T_i = \text{fixed treatment effects} \rightarrow \sum T_i = 0$$

γ = regression coefficient for the relation between Y and X

X_{ij} are constants (concomitant observations)

$$\epsilon_{ij} \sim N(0, \sigma^2)$$

$$i = 1, \dots, r ; j = 1, \dots, n_i$$

$$E(Y_{ij}) = \mu_0 + T_i + \gamma(X_{ij} - \bar{X}_{..})$$

$$\sigma^2(Y_{ij}) = \sigma^2$$

Note: this depends on concomitant variables
so all observations of treatment won't
have the same mean response

We can generalize this model for X 's that are random variables

We can generalize for nonlinear Y/X relationships

ie quadratic: $Y_{ij} = \mu_0 + T_i + \gamma_1(X_{ij1} - \bar{X}_{..1}) + \gamma_2(X_{ij2} - \bar{X}_{..2}) + \epsilon_{ij}$

We can add more concomitant observations

ie two: $Y_{ij} = \mu_0 + T_i + \gamma_1(X_{ij1} - \bar{X}_{..1}) + \gamma_2(X_{ij2} - \bar{X}_{..2}) + \epsilon_{ij}$

Assumptions ① Normality of error terms

② Constant error variance across treatments

③ Equality of slope of different treatment regression lines

④ Linearity of regression lines

⑤ Uncorrelated error terms

Hyp Test: $H_0: \tau_1 = \tau_2 = \dots = \tau_r = 0$

Hai not H_0

$$F^* = \frac{\text{MSR}(x_q, \dots, x_{p-1} | x_1, \dots, x_{q-1})}{\text{MSE}}$$

Reject for large F^*

* If we fail to reject we're interested in looking at linear combinations of regression coefficients to investigate these effects.

* Look at Type III tests of fixed effects in SAS below

SAS

```
proc sgscatter; /* Gives Anova scatterplot linear → no interaction */
  plot y * concomitant / group=x loess;
run;

proc glimmix; /* Get regression equation */
  class x;
  model y = x concomitant / solution; /*  $y_{ij} = \mu_{..} + \tau_i I_{ij..} + \dots + \gamma x_{ij} + \epsilon_{ij}$  */
  lsmeans x / "M1 - M2" l=1 O / adjust=t cl alpha=.05; /* treat diff */
  lsmeans x; /* gives  $\mu_1, \mu_2, \dots$  estimates */
run;

proc glimmix; /* test x * concomitant = 0 ≡ parallel lines in Anova scatter */
  class x;
  model y = x | concomitant;
run;
```

Covariance Model for Two Factors

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

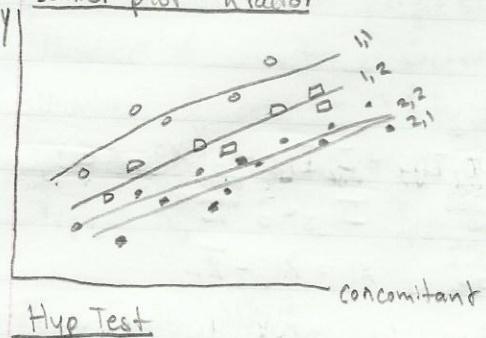
$$Y_{ijk} = \mu_{..} + \alpha_i + \beta_j + (\alpha\beta)_{ij} + \gamma(X_{ijk} - \bar{X}_{...}) + \epsilon_{ijk}$$

In fact, we fit it differently $\rightarrow = \mu_{..} + \alpha_1 I_{ijk1} + \beta_1 I_{ijk2} + (\alpha\beta)_{11} I_{ijk1} I_{ijk2} + \gamma X_{ijk} + \epsilon_{ijk}$

$$I_1 = \begin{cases} 1 & \text{level 1 factor A} \\ 0 & \text{level 2 factor A} \end{cases}$$

$$I_2 = \begin{cases} 1 & \text{level 1 factor B} \\ 0 & \text{level 2 factor B} \end{cases}$$

Scatter plot - 2 factor



Hyp Test

concomitant

H_0 : no an interaction

H_1 : \exists interaction

$$F^* = \frac{\text{SSR}(I, I_2 | X, I_1, I_2)}{\text{MSE}}$$

MSE

Reject for large F^*

Linear Combinations $\alpha_1 - \alpha_2$ $\beta_1 - \beta_2$ } Comparing main effects factor levels

SAS data twoFactor;

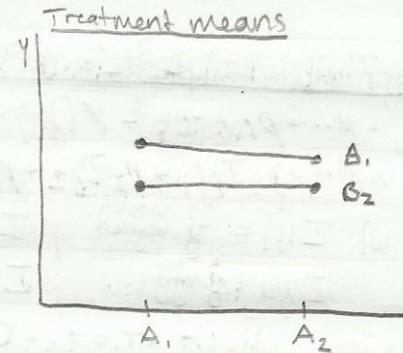
```
set ____;
group = '11';
if x1=1 and x2=2 then group = '12';
if x1=2 and x2=1 then group = '21';
if x1=2 and x2=2 then group = '22';
run;
```

```
proc sgscatter data=twoFactor;
plot y*concomitant / group=group loess;
run;
```

```
proc glm; /* Check type 3 test for interaction */
class x1 x2;
model y=concomitant x1|x2 / solution;
run;
```

```
proc glimmix; class x1 x2; /* Comparing main effects */
model y=concomitant x1 x2 / solution;
lsmeans x1 "α1-α2" l=1 / cl α=.05;
lsmeans x2 "β1-β2" l=1 / cl α=.05;
run;
```

Treatment means



Covariance Analysis of RCB Design

$$\begin{aligned} Y_{ij} &= \mu_{..} + \rho_1 + I_{ij} + \gamma(X_{ij} - \bar{X}_{..}) + \epsilon_{ij} \\ &= \mu_{..} + \rho_1 I_{ij1} + \rho_2 I_{ij2} + \rho_3 I_{ij3} + \tau_i I_{ij4} + I_{ij5} + \gamma X_{ij} + \epsilon_{ij} \\ \text{w/ } I_{ij1} &= \begin{cases} 1 & \text{block 1} \\ 0 & \text{block 2} \\ 0 & \text{block 3} \\ 0 & \text{block 4} \end{cases}, \quad I_{ij2} = \begin{cases} 0 & \text{block 1} \\ 1 & \text{block 2} \\ 0 & \text{block 3} \\ 0 & \text{block 4} \end{cases}, \quad I_{ij3} = \begin{cases} 0 & \text{block 1} \\ 0 & \text{block 2} \\ 1 & \text{block 3} \\ 0 & \text{block 4} \end{cases} \\ I_{ij4} &= \begin{cases} 1 & \text{Treat 1} \\ 0 & \text{Treat 2} \\ 0 & \text{Treat 3} \end{cases}, \quad I_{ij5} = \begin{cases} 0 & \text{Treat 1} \\ 0 & \text{Treat 2} \\ 0 & \text{Treat 3} \end{cases}, \quad X_{ij} = X_{ij} - \bar{X}_{..} \end{aligned}$$

Hyp Test $H_0: \tau_1 = \tau_2 = \tau_3 = 0$

$H_a: \text{not } H_0$

Use p value in type III table