

Notes 9: Analysis of Variance (N-Way)

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Outline of Notes

1) Balanced Two-Way ANOVA:

- Model Form & Assumptions
- Least-Squares Estimation
- Basic Inference
- Hypertension Example (pt 1)
- Multiple Comparisons
- Hypertension Example (pt 2)

2) Miscellaneous:

- Three-Way ANOVA
- Unbalanced ANOVA
- Kruskal-Wallis Test
(Nonparametric 1-Way ANOVA)

Two-Way ANOVA Model (cell means form)

The Two-Way Analysis of Variance (ANOVA) model has the form

$$y_{ijk} = \mu_{jk} + e_{ijk}$$

for $i \in \{1, \dots, n_{jk}\}$, $j \in \{1, \dots, a\}$, and $k \in \{1, \dots, b\}$ where

- $y_{ijk} \in \mathbb{R}$ is real-valued response for i -th subject in factor cell (j, k)
- $\mu_{jk} \in \mathbb{R}$ is real-valued population mean for factor cell (j, k)
- $e_{ijk} \stackrel{\text{iid}}{\sim} N(0, \sigma^2)$ is Gaussian measurement error
- n_{jk} is number of subjects in cell (j, k) and $n = \sum_{j=1}^a \sum_{k=1}^b n_{jk}$
(note: $n_{jk} = n_*$ $\forall j, k$ in balanced two-way ANOVA)
- a and b are number of levels for first and second factors

Implies that $y_{ijk} \stackrel{\text{ind}}{\sim} N(\mu_{jk}, \sigma^2)$.

Two-Way ANOVA Model (effect coding: interaction)

Using effect coding, the mean for factor cell (j, k) has the form

$$\mu_{jk} = \mu + \alpha_j + \beta_k + \gamma_{jk}$$

for $j \in \{1, \dots, a\}$ and $k \in \{1, \dots, b\}$ where

- μ is overall population mean
- α_j is main effect of first factor such that $\sum_{j=1}^a \alpha_j = 0$
- β_k is main effect of second factor such that $\sum_{k=1}^b \beta_k = 0$
- γ_{jk} is interaction effect such that $\sum_{j=1}^a \gamma_{jk} = 0 \forall k$ and $\sum_{k=1}^b \gamma_{jk} = 0 \forall j$

Two-Way ANOVA Model (effect coding: additive)

Using effect coding, the mean for factor cell (j, k) has the form

$$\mu_{jk} = \mu + \alpha_j + \beta_k$$

for $j \in \{1, \dots, a\}$ and $k \in \{1, \dots, b\}$ where

- μ is overall population mean
- α_j is main effect of first factor such that $\sum_{j=1}^a \alpha_j = 0$
- β_k is main effect of second factor such that $\sum_{k=1}^b \beta_k = 0$

Two-Way ANOVA Model (matrix form: interaction)

In matrix form, the two-way ANOVA model is $\mathbf{y} = \mathbf{X}\mathbf{b} + \mathbf{e}$ where

$$\mathbf{X} = \begin{pmatrix} 1 & x_{11} \cdots x_{1(a-1)} & z_{11} \cdots z_{1(b-1)} & x_{11}z_{11} \cdots x_{1(a-1)}z_{1(b-1)} & \cdots & x_{11}z_{1(b-1)} \cdots x_{1(a-1)}z_{1(b-1)} \\ 1 & x_{21} \cdots x_{2(a-1)} & z_{21} \cdots z_{2(b-1)} & x_{21}z_{21} \cdots x_{2(a-1)}z_{2(b-1)} & \cdots & x_{21}z_{2(b-1)} \cdots x_{2(a-1)}z_{2(b-1)} \\ 1 & x_{31} \cdots x_{3(a-1)} & z_{31} \cdots z_{3(b-1)} & x_{31}z_{31} \cdots x_{3(a-1)}z_{3(b-1)} & \cdots & x_{31}z_{3(b-1)} \cdots x_{3(a-1)}z_{3(b-1)} \\ \vdots & \vdots \quad \ddots \quad \vdots & \vdots \quad \ddots \quad \vdots & \vdots \quad \ddots \quad \vdots & \ddots & \vdots \quad \ddots \quad \vdots \\ 1 & x_{n1} \cdots x_{n(a-1)} & z_{n1} \cdots z_{n(b-1)} & x_{n1}z_{n1} \cdots x_{n(a-1)}z_{n1} & \cdots & x_{n1}z_{n(b-1)} \cdots x_{n(a-1)}z_{n(b-1)} \end{pmatrix}$$

$$\mathbf{b} = (\mu \quad \alpha_1 \cdots \alpha_{a-1} \quad \beta_1 \cdots \beta_{b-1} \quad \gamma_{11} \cdots \gamma_{(a-1)1} \quad \cdots \quad \gamma_{1(b-1)} \cdots \gamma_{(a-1)(b-1)})'$$

where \mathbf{X} has $1 + (a - 1) + (b - 1) + (a - 1)(b - 1) = ab$ columns

- $x_{ij} = \begin{cases} 1 & \text{if } i\text{-th observation is in } j\text{-th level of first factor} \\ -1 & \text{if } i\text{-th observation is in } a\text{-th level of first factor} \\ 0 & \text{otherwise} \end{cases}$
- $z_{ik} = \begin{cases} 1 & \text{if } i\text{-th observation is in } k\text{-th level of second factor} \\ -1 & \text{if } i\text{-th observation is in } b\text{-th level of second factor} \\ 0 & \text{otherwise} \end{cases}$
- $i \in \{1, \dots, n\}$ and additional subscripts on y and e are dropped

Implies that $\mathbf{y} \sim N(\mathbf{X}\mathbf{b}, \sigma^2 \mathbf{I}_n)$.

Two-Way ANOVA Model (matrix form: additive)

In matrix form, the two-way ANOVA model is $\mathbf{y} = \mathbf{X}\mathbf{b} + \mathbf{e}$ where

$$\mathbf{X} = \begin{pmatrix} 1 & x_{11} \cdots x_{1(a-1)} & z_{11} \cdots z_{1(b-1)} \\ 1 & x_{21} \cdots x_{2(a-1)} & z_{21} \cdots z_{2(b-1)} \\ 1 & x_{31} \cdots x_{3(a-1)} & z_{31} \cdots z_{3(b-1)} \\ \vdots & \vdots \quad \ddots \quad \vdots & \vdots \quad \ddots \quad \vdots \\ 1 & x_{n1} \cdots x_{n(a-1)} & z_{n1} \cdots z_{n(b-1)} \end{pmatrix}$$

$$\mathbf{b} = (\mu \quad \alpha_1 \cdots \alpha_{a-1} \quad \beta_1 \cdots \beta_{b-1})'$$

where \mathbf{X} has $1 + (a - 1) + (b - 1) = a + b - 1$ columns

- $x_{ij} = \begin{cases} 1 & \text{if } i\text{-th observation is in } j\text{-th level of first factor} \\ -1 & \text{if } i\text{-th observation is in } a\text{-th level of first factor} \\ 0 & \text{otherwise} \end{cases}$
- $z_{ik} = \begin{cases} 1 & \text{if } i\text{-th observation is in } k\text{-th level of second factor} \\ -1 & \text{if } i\text{-th observation is in } b\text{-th level of second factor} \\ 0 & \text{otherwise} \end{cases}$
- $i \in \{1, \dots, n\}$ and additional subscripts on y and e are dropped

Implies that $\mathbf{y} \sim N(\mathbf{X}\mathbf{b}, \sigma^2 \mathbf{I}_n)$.

Two-Way ANOVA Model (assumptions)

The fundamental assumptions of the two-way ANOVA model are:

- 1 x_{ij} , z_{ik} and y_i are observed random variables (constants)
- 2 $e_i \stackrel{\text{iid}}{\sim} N(0, \sigma^2)$ is an unobserved random variable
- 3 μ_{jk} are unknown constants
- 4 $(y_i | x_{ij}, z_{ik}) \stackrel{\text{ind}}{\sim} N(\mu_{jk}, \sigma^2)$
note: homogeneity of variance

Interpretation of μ_{jk} depends on model form

- Additive: $\mu_{jk} = \mu + \alpha_j + \beta_k$
- Interaction: $\mu_{jk} = \mu + \alpha_j + \beta_k + \gamma_{jk}$

Ordinary Least-Squares (interaction)

We want to find the effect estimates (i.e., $\hat{\mu}$, $\hat{\alpha}_j$, $\hat{\beta}_k$, and $\hat{\gamma}_{jk}$ terms) that minimize the ordinary least squares criterion

$$SSE = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} (y_{ijk} - \mu - \alpha_j - \beta_k - \gamma_{jk})^2$$

If $n_{jk} = n_* \forall j, k$ the least-squares estimates have the form

$$\hat{\mu} = \frac{1}{abn_*} \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} y_{ijk} = \bar{y}_{...}$$

$$\hat{\alpha}_j = \left(\frac{1}{bn_*} \sum_{k=1}^b \sum_{i=1}^{n_*} y_{ijk} \right) - \hat{\mu} = \bar{y}_{\cdot j \cdot} - \bar{y}_{...}$$

$$\hat{\beta}_k = \left(\frac{1}{an_*} \sum_{j=1}^a \sum_{i=1}^{n_*} y_{ijk} \right) - \hat{\mu} = \bar{y}_{\cdot \cdot k} - \bar{y}_{...}$$

$$\hat{\gamma}_{jk} = \left(\frac{1}{n_*} \sum_{i=1}^{n_*} y_{ijk} \right) - \hat{\mu} - \hat{\alpha}_j - \hat{\beta}_k = \bar{y}_{\cdot jk} - \bar{y}_{\cdot j \cdot} - \bar{y}_{\cdot \cdot k} + \bar{y}_{...}$$

which implies that $\hat{y}_{ijk} = \bar{y}_{\cdot jk}$ for all (i, j, k) .

Ordinary Least-Squares (interaction proof for μ)

Expanding the first summation produces

$$SSE = \sum_{k=1}^b \sum_{j=1}^a \left[\sum_{i=1}^{n_*} y_{ijk}^2 - 2(\mu + \alpha_j + \beta_k + \gamma_{jk}) \sum_{i=1}^{n_*} y_{ijk} + n_*(\mu + \alpha_j + \beta_k + \gamma_{jk})^2 \right]$$

Taking the derivative with respect to μ we have

$$\begin{aligned} \frac{dSSE}{d\mu} &= \sum_{k=1}^b \sum_{j=1}^a \left[-2 \sum_{i=1}^{n_*} y_{ijk} + 2n_*\mu + 2n_*(\alpha_j + \beta_k + \gamma_{jk}) \right] \\ &= -2 \left(\sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} y_{ijk} \right) + 2abn_*\mu \end{aligned}$$

and setting to zero and solving for μ gives

$$\hat{\mu} = \frac{1}{abn_*} \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} y_{ijk} = \bar{y}_{...}$$

Ordinary Least-Squares (interaction proof for α_j)

Taking the derivative with respect to α_j we have

$$\begin{aligned}\frac{dSSE}{d\alpha_j} &= \sum_{k=1}^b \left[-2 \sum_{i=1}^{n_*} y_{ijk} + 2n_*\alpha_j + 2n_*(\mu + \beta_k + \gamma_{jk}) \right] \\ &= -2 \left(\sum_{k=1}^b \sum_{i=1}^{n_*} y_{ijk} \right) + 2bn_*\alpha_j + 2bn_*\mu\end{aligned}$$

and setting to zero, using $\hat{\mu}$ for μ , and solving for α_j gives

$$\hat{\alpha}_j = \frac{1}{bn_*} \left(\sum_{k=1}^b \sum_{i=1}^{n_*} y_{ijk} \right) - \hat{\mu} = \bar{y}_{\cdot j} - \bar{y}_{\dots}$$

Ordinary Least-Squares (interaction proof for β_k)

Taking the derivative with respect to β_k we have

$$\begin{aligned}\frac{dSSE}{d\beta_k} &= \sum_{j=1}^a [-2 \sum_{i=1}^{n_*} y_{ijk} + 2n_*\beta_k + 2n_*(\mu + \alpha_j + \gamma_{jk})] \\ &= -2 \left(\sum_{j=1}^a \sum_{i=1}^{n_*} y_{ijk} \right) + 2an_*\beta_k + 2an_*\mu\end{aligned}$$

and setting to zero, using $\hat{\mu}$ for μ , and solving for β_k gives

$$\hat{\beta}_k = \frac{1}{an_*} \left(\sum_{j=1}^a \sum_{i=1}^{n_*} y_{ijk} \right) - \hat{\mu} = \bar{y}_{..k} - \bar{y}_{..}$$

Ordinary Least-Squares (interaction proof for γ_{jk})

Taking the derivative with respect to γ_{jk} we have

$$\frac{dSSE}{d\gamma_{jk}} = -2 \sum_{i=1}^{n_*} y_{ijk} + 2n_*\gamma_{jk} + 2n_*(\mu + \alpha_j + \beta_k)$$

and setting to zero, using $(\hat{\mu}, \hat{\alpha}_j, \hat{\beta}_k)$ for (μ, α_j, β_k) , and solving for γ_{jk} gives $\hat{\gamma}_{jk} = \frac{1}{n_*}(\sum_{i=1}^{n_*} y_{ijk}) - \hat{\mu} - \hat{\alpha}_j - \hat{\beta}_k = \bar{y}_{\cdot jk} - \bar{y}_{\cdot j} - \bar{y}_{\cdot \cdot k} + \bar{y}_{\dots}$

Ordinary Least-Squares (additive)

We want to find the effect estimates (i.e., $\hat{\mu}$, $\hat{\alpha}_j$ and $\hat{\beta}_k$ terms) that minimize the ordinary least squares criterion

$$SSE = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} (y_{ijk} - \mu - \alpha_j - \beta_k)^2$$

If $n_{jk} = n_* \forall j, k$ the least-squares estimates have the form

$$\hat{\mu} = \bar{y}_{...} = \frac{1}{abn_*} \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} y_{ijk}$$

$$\hat{\alpha}_j = \bar{y}_{.j.} - \hat{\mu} = \left(\frac{1}{bn_*} \sum_{k=1}^b \sum_{i=1}^{n_*} y_{ijk} \right) - \bar{y}_{...}$$

$$\hat{\beta}_k = \bar{y}_{..k} - \hat{\mu} = \left(\frac{1}{an_*} \sum_{j=1}^a \sum_{i=1}^{n_*} y_{ijk} \right) - \bar{y}_{...}$$

which implies $\hat{y}_{ijk} = \bar{y}_{.j.} + \bar{y}_{..k} - \bar{y}_{...}$ for all (i, j, k) , see previous proofs.

Fitted Values and Residuals

Form of fitted values depends on fit model:

- Additive: $\hat{\mu}_{jk} = \bar{y}_{\cdot j} + \bar{y}_{\cdot \cdot k} - \bar{y}_{\cdot \cdot \cdot}$
- Interaction: $\hat{\mu}_{jk} = \bar{y}_{\cdot jk}$

Residuals have the form

$$\hat{e}_{ijk} = y_{ijk} - \hat{\mu}_{jk}$$

where form of $\hat{\mu}_{jk}$ depends on fit model (additive versus interaction).

ANOVA Sums-of-Squares

In balanced two-way ANOVA model with interaction:

- $SST = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} (y_{ijk} - \bar{y}_{...})^2$ $df = abn_* - 1$
- $SSR = n_* \sum_{k=1}^b \sum_{j=1}^a (\bar{y}_{\cdot jk} - \bar{y}_{...})^2$ $df = ab - 1$
- $SSE = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} (y_{ijk} - \bar{y}_{\cdot jk})^2$ $df = abn_* - ab$

In balanced two-way ANOVA model with no interaction:

- $SST = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} (y_{ijk} - \bar{y}_{...})^2$ $df = abn_* - 1$
- $SSR = n_* \sum_{k=1}^b \sum_{j=1}^a ([\bar{y}_{\cdot j \cdot} + \bar{y}_{\cdot \cdot k} - \bar{y}_{...}] - \bar{y}_{...})^2$ $df = a + b - 2$
- $SSE = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} (y_{ijk} - [\bar{y}_{\cdot j \cdot} + \bar{y}_{\cdot \cdot k} - \bar{y}_{...}])^2$
 $df = abn_* - (a + b - 1)$

Partitioning the Variance

From Notes 5 we know that $SST = SSR + SEE$.

If $n_{jk} = n_* \forall j, k$ can partition $SSR = SSA + SSB + SSAB$ where

- $SSA = bn_* \sum_{j=1}^a (\bar{y}_{\cdot j} - \bar{y}_{\dots})^2$ $df = a - 1$
- $SSB = an_* \sum_{k=1}^b (\bar{y}_{\cdot k} - \bar{y}_{\dots})^2$ $df = b - 1$
- $SSAB = n_* \sum_{k=1}^b \sum_{j=1}^a (\bar{y}_{jk} - \bar{y}_{\cdot j} - \bar{y}_{\cdot k} + \bar{y}_{\dots})^2$ $df = (a - 1)(b - 1)$

Implies that $SSR = SSA + SSB$ for additive model (if $n_{jk} = n_* \forall j, k$).

Partitioning the Variance (proof part 1)

To prove $SSR = SSA + SSB + SSAB$ when $n_{jk} = n_* \forall j, k$, note that

$$y_{ijk} - \bar{y}_{...} = (y_{ijk} - \bar{y}_{\cdot jk}) + (\bar{y}_{\cdot jk} - [\bar{y}_{\cdot j} + \bar{y}_{\cdot k} - \bar{y}_{...}]) + (\bar{y}_{\cdot j} - \bar{y}_{...}) + (\bar{y}_{\cdot k} - \bar{y}_{...})$$

Now if we square both sides we have

$$\begin{aligned} (y_{ijk} - \bar{y}_{...})^2 &= (y_{ijk} - \bar{y}_{\cdot jk})^2 + (\bar{y}_{\cdot jk} - [\bar{y}_{\cdot j} + \bar{y}_{\cdot k} - \bar{y}_{...}])^2 + (\bar{y}_{\cdot j} - \bar{y}_{...})^2 + (\bar{y}_{\cdot k} - \bar{y}_{...})^2 \\ &\quad + 2(y_{ijk} - \bar{y}_{\cdot jk}) \{ (\bar{y}_{\cdot jk} - [\bar{y}_{\cdot j} + \bar{y}_{\cdot k} - \bar{y}_{...}]) + (\bar{y}_{\cdot j} - \bar{y}_{...}) + (\bar{y}_{\cdot k} - \bar{y}_{...}) \} \\ &\quad + 2(\bar{y}_{\cdot jk} - [\bar{y}_{\cdot j} + \bar{y}_{\cdot k} - \bar{y}_{...}]) [(\bar{y}_{\cdot j} - \bar{y}_{...}) + (\bar{y}_{\cdot k} - \bar{y}_{...})] \\ &\quad + 2(\bar{y}_{\cdot j} - \bar{y}_{...})(\bar{y}_{\cdot k} - \bar{y}_{...}) \end{aligned}$$

Now if we apply the triple summation we have SST

$$SST = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} (y_{ijk} - \bar{y}_{...})^2$$

Partitioning the Variance (proof part 2)

First, note that we have

$$SSE = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} (y_{ijk} - \bar{y}_{\cdot jk})^2$$

$$SSAB = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} (\bar{y}_{\cdot jk} - [\bar{y}_{\cdot j\cdot} + \bar{y}_{\cdot\cdot k} - \bar{y}_{\cdot\cdot\cdot}])^2$$

$$SSA = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} (\bar{y}_{\cdot j\cdot} - \bar{y}_{\cdot\cdot\cdot})^2$$

$$SSB = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} (\bar{y}_{\cdot\cdot k} - \bar{y}_{\cdot\cdot\cdot})^2$$

so we need to prove that the crossproduct terms are orthogonal.

To prove that the first crossproduct term sums to zero, define

$\delta_{jk} = (\bar{y}_{\cdot jk} - [\bar{y}_{\cdot j\cdot} + \bar{y}_{\cdot\cdot k} - \bar{y}_{\cdot\cdot\cdot}]) + (\bar{y}_{\cdot j\cdot} - \bar{y}_{\cdot\cdot\cdot}) + (\bar{y}_{\cdot\cdot k} - \bar{y}_{\cdot\cdot\cdot})$ and note that

$$\begin{aligned} \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} 2(y_{ijk} - \bar{y}_{\cdot jk})\delta_{jk} &= 2 \sum_{k=1}^b \sum_{j=1}^a \delta_{jk} \sum_{i=1}^{n_{jk}} (y_{ijk} - \bar{y}_{\cdot jk}) \\ &= 2 \sum_{k=1}^b \sum_{j=1}^a \delta_{jk}(0) = 0 \end{aligned}$$

because we are summing mean-centered variable.

Partitioning the Variance (proof part 3)

To prove that the second crossproduct term sums to zero, note that $\hat{\gamma}_{jk} = (\bar{y}_{\cdot jk} - [\bar{y}_{\cdot j} + \bar{y}_{\cdot k} - \bar{y}_{\cdot \cdot}])$, $\hat{\alpha}_j = (\bar{y}_{\cdot j} - \bar{y}_{\cdot \cdot})$, and $\hat{\beta}_k = (\bar{y}_{\cdot k} - \bar{y}_{\cdot \cdot})$, so

$$\sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} 2\hat{\gamma}_{jk}(\hat{\alpha}_j + \hat{\beta}_k) = 2 \sum_{k=1}^b \sum_{j=1}^a n_{jk} \hat{\gamma}_{jk}(\hat{\alpha}_j + \hat{\beta}_k) = 0$$

because

$$n_{jk} \hat{\gamma}_{jk} = \left(\sum_{i=1}^{n_{jk}} y_{ijk} \right) - \left(\frac{1}{b} \sum_{k=1}^b \sum_{i=1}^{n_{jk}} y_{ijk} \right) - \left(\frac{1}{a} \sum_{j=1}^a \sum_{i=1}^{n_{jk}} y_{ijk} \right) + \left(\frac{1}{ab} \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} y_{ijk} \right)$$

which implies that

$$\sum_{k=1}^b \sum_{j=1}^a n_{jk} \hat{\gamma}_{jk} \hat{\alpha}_j = \sum_{j=1}^a \hat{\alpha}_j \left(\sum_{k=1}^b n_{jk} \hat{\gamma}_{jk} \right) = \sum_{j=1}^a \hat{\alpha}_j(0) = 0$$

$$\sum_{k=1}^b \sum_{j=1}^a n_{jk} \hat{\gamma}_{jk} \hat{\beta}_k = \sum_{k=1}^b \hat{\beta}_k \left(\sum_{j=1}^a n_{jk} \hat{\gamma}_{jk} \right) = \sum_{k=1}^b \hat{\beta}_k(0) = 0$$

Partitioning the Variance (proof part 4)

To prove that the third crossproduct term sums to zero, note that

$$\sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} 2(\bar{y}_{\cdot j} - \bar{y}_{\dots})(\bar{y}_{\cdot k} - \bar{y}_{\dots}) = 2 \sum_{k=1}^b \sum_{j=1}^a n_{jk} \hat{\alpha}_j \hat{\beta}_k$$

and if $n_{jk} = n_* \forall j, k$ we have that

$$\begin{aligned} 2 \sum_{k=1}^b \sum_{j=1}^a n_{jk} \hat{\alpha}_j \hat{\beta}_k &= 2n_* \sum_{k=1}^b \sum_{j=1}^a \hat{\alpha}_j \hat{\beta}_k \\ &= 2n_* \sum_{k=1}^b \hat{\beta}_k \left(\sum_{j=1}^a \hat{\alpha}_j \right) \\ &= 2n_* \sum_{k=1}^b \hat{\beta}_k(0) = 0 \end{aligned}$$

which completes the proof; note that this is the **ONLY** part of the proof that requires the balanced assumption.

Extended ANOVA Table and F Tests

We typically organize the SS information into an ANOVA table:

| Source | SS | df | MS | F | p-value |
|--------|--|------------------|--------|------------|------------|
| SSR | $n_* \sum_{k=1}^b \sum_{j=1}^a (\bar{y}_{\cdot jk} - \bar{y}_{\cdot \cdot})^2$ | $ab - 1$ | MSR | F^* | p^* |
| SSA | $bn_* \sum_{j=1}^a (\bar{y}_{\cdot j} - \bar{y}_{\cdot \cdot})^2$ | $a - 1$ | MSA | F_a^* | p_a^* |
| SSB | $an_* \sum_{k=1}^b (\bar{y}_{\cdot \cdot k} - \bar{y}_{\cdot \cdot})^2$ | $b - 1$ | MSB | F_b^* | p_b^* |
| SSAB | $n_* \sum_{k=1}^b \sum_{j=1}^a (\bar{y}_{\cdot jk} - \bar{y}_{\cdot j} - \bar{y}_{\cdot \cdot k} + \bar{y}_{\cdot \cdot})^2$ | $(a - 1)(b - 1)$ | $MSAB$ | F_{ab}^* | p_{ab}^* |
| SSE | $\sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} (y_{ijk} - \bar{y}_{\cdot jk})^2$ | $ab(n_* - 1)$ | MSE | | |
| SST | $\sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} (y_{ijk} - \bar{y}_{\cdot \cdot})^2$ | $abn_* - 1$ | | | |

$$MSR = \frac{SSR}{ab-1}, MSA = \frac{SSA}{a-1}, MSB = \frac{SSB}{b-1}, MSAB = \frac{SSAB}{(a-1)(b-1)}, MSE = \frac{SSE}{ab(n_*-1)},$$

$$F^* = \frac{MSR}{MSE} \sim F_{ab-1, ab(n_*-1)} \quad \text{and} \quad p^* = P(F_{ab-1, ab(n_*-1)} > F^*),$$

$$F_a^* = \frac{MSA}{MSE} \sim F_{a-1, ab(n_*-1)} \quad \text{and} \quad p_a^* = P(F_{a-1, ab(n_*-1)} > F_a^*),$$

$$F_b^* = \frac{MSB}{MSE} \sim F_{b-1, ab(n_*-1)} \quad \text{and} \quad p_b^* = P(F_{b-1, ab(n_*-1)} > F_b^*),$$

$$F_{ab}^* = \frac{MSAB}{MSE} \sim F_{(a-1)(b-1), ab(n_*-1)} \quad \text{and} \quad p_{ab}^* = P(F_{(a-1)(b-1), ab(n_*-1)} > F_{ab}^*),$$

ANOVA Table F Tests

F^* statistic and p^* -value are testing $H_0 : \alpha_j = \beta_k = \gamma_{jk} = 0 \ \forall j, k$ versus $H_1 : (\exists j, k \in \{1, \dots, a\} \times \{1, \dots, b\})(\alpha_j = \beta_k = \gamma_{jk} = 0 \text{ is false})$

- Equivalent to $H_0 : \mu_{jk} = \mu \ \forall j, k$ versus $H_1 : \text{not all } \mu_{jk} \text{ are equal}$

F_a^* statistic and p_a^* -value are testing $H_0 : \alpha_j = 0 \ \forall j$ versus $H_1 : (\exists j \in \{1, \dots, a\})(\alpha_j \neq 0)$

- Testing main effect of first factor

F_b^* statistic and p_b^* -value are testing $H_0 : \beta_k = 0 \ \forall k$ versus $H_1 : (\exists k \in \{1, \dots, b\})(\beta_k \neq 0)$

- Testing main effect of second factor

F_{ab}^* statistic and p_{ab}^* -value are testing $H_0 : \gamma_{jk} = 0 \ \forall j, k$ versus $H_1 : (\exists j, k \in \{1, \dots, a\} \times \{1, \dots, b\})(\gamma_{jk} \neq 0)$

- Testing interaction effect

Hypertension Example: Data Description

Hypertension example from Maxwell & Delany (2003).

Total of $n = 72$ subjects participate in hypertension experiment.

- Factor A: `drug` type ($a = 3$ levels: X, Y, Z)
- Factor B: `diet` type ($b = 2$ levels: yes, no)

Randomly assign $n_{jk} = 12$ subjects to each treatment cell:

- Note there are $(ab) = (3)(2) = 6$ treatment cells
- Observations are independent within and between cells

Hypertension Example: Descriptive Statistics

Sum of blood pressure for each treatment cell ($\sum_{i=1}^{12} y_{ijk}$):

| Drug | Diet | | Total |
|---------------|----------------|-----------------|-------|
| | No ($k = 1$) | Yes ($k = 2$) | |
| X ($j = 1$) | 2136 | 2052 | 4188 |
| Y ($j = 2$) | 2424 | 2154 | 4578 |
| Z ($j = 3$) | 2388 | 2130 | 4518 |
| Total | 6948 | 6336 | 13284 |

Sum-of-squares of blood pressure for each treatment cell ($\sum_{i=1}^{12} y_{ijk}^2$):

| Drug | Diet | | Total |
|---------------|----------------|-----------------|---------|
| | No ($k = 1$) | Yes ($k = 2$) | |
| X ($j = 1$) | 382368 | 352518 | 734886 |
| Y ($j = 2$) | 491008 | 388898 | 879906 |
| Z ($j = 3$) | 478238 | 380462 | 858700 |
| Total | 1351614 | 1121878 | 2473492 |

Hypertension Example: OLS Estimation (by hand)

Least-squares estimates are cell means: $\hat{\mu}_{jk} = \bar{y}_{.jk}$ and

$$\hat{\mu} = \frac{1}{abn_*} \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_*} y_{ijk} = \bar{y}_{...} = \frac{13284}{72} = 184.5$$

$$\hat{\alpha}_1 = \left(\frac{1}{bn_*} \sum_{k=1}^b \sum_{i=1}^{n_*} y_{i1k} \right) - \hat{\mu} = \bar{y}_{.1.} - \bar{y}_{...} = \frac{4188}{24} - 184.5 = -10$$

$$\hat{\alpha}_2 = \left(\frac{1}{bn_*} \sum_{k=1}^b \sum_{i=1}^{n_*} y_{i2k} \right) - \hat{\mu} = \bar{y}_{.2.} - \bar{y}_{...} = \frac{4578}{24} - 184.5 = 6.25$$

$$\hat{\alpha}_3 = \left(\frac{1}{bn_*} \sum_{k=1}^b \sum_{i=1}^{n_*} y_{i3k} \right) - \hat{\mu} = \bar{y}_{.3.} - \bar{y}_{...} = \frac{4518}{24} - 184.5 = 3.75$$

$$\hat{\beta}_1 = \left(\frac{1}{an_*} \sum_{j=1}^a \sum_{i=1}^{n_*} y_{ij1} \right) - \hat{\mu} = \bar{y}_{..1} - \bar{y}_{...} = \frac{6948}{36} - 184.5 = 8.5$$

$$\hat{\beta}_2 = \left(\frac{1}{an_*} \sum_{j=1}^a \sum_{i=1}^{n_*} y_{ij2} \right) - \hat{\mu} = \bar{y}_{..2} - \bar{y}_{...} = \frac{6336}{36} - 184.5 = -8.5$$

Hypertension Example: OLS Estimation (by hand)

Continuing from the previous slide...

$$\hat{\gamma}_{11} = \bar{y}_{.11} - \bar{y}_{.1.} - \bar{y}_{..1} + \bar{y}_{...} = \frac{2136}{12} - \frac{4188}{24} - \frac{6948}{36} + 184.5 = -5$$

$$\hat{\gamma}_{12} = \bar{y}_{.12} - \bar{y}_{.1.} - \bar{y}_{..2} + \bar{y}_{...} = \frac{2052}{12} - \frac{4188}{24} - \frac{6336}{36} + 184.5 = 5$$

$$\hat{\gamma}_{21} = \bar{y}_{.21} - \bar{y}_{.2.} - \bar{y}_{..1} + \bar{y}_{...} = \frac{2424}{12} - \frac{4578}{24} - \frac{6948}{36} + 184.5 = 2.75$$

$$\hat{\gamma}_{22} = \bar{y}_{.22} - \bar{y}_{.2.} - \bar{y}_{..2} + \bar{y}_{...} = \frac{2154}{12} - \frac{4578}{24} - \frac{6336}{36} + 184.5 = -2.75$$

$$\hat{\gamma}_{31} = \bar{y}_{.31} - \bar{y}_{.3.} - \bar{y}_{..1} + \bar{y}_{...} = \frac{2388}{12} - \frac{4518}{24} - \frac{6948}{36} + 184.5 = 2.25$$

$$\hat{\gamma}_{32} = \bar{y}_{.32} - \bar{y}_{.3.} - \bar{y}_{..2} + \bar{y}_{...} = \frac{2130}{12} - \frac{4518}{24} - \frac{6336}{36} + 184.5 = -2.25$$

Hypertension Example: Enter Data (in R)

```
> bp=scan("/Users/Nate/Desktop/hypertension.dat")
Read 72 items
> diet=factor(rep(rep(c("no", "yes"), each=6), 6))
> drug=factor(rep(rep(c("X", "Y", "Z"), each=12), 2))
> biof=factor(rep(c("present", "absent"), each=36))
> hyper=data.frame(bp=bp, diet=diet, drug=drug, biof=biof)
> hyper[1:20,]
```

| | bp | diet | drug | biof |
|----|-----|------|------|---------|
| 1 | 170 | no | X | present |
| 2 | 175 | no | X | present |
| 3 | 165 | no | X | present |
| 4 | 180 | no | X | present |
| 5 | 160 | no | X | present |
| 6 | 158 | no | X | present |
| 7 | 161 | yes | X | present |
| 8 | 173 | yes | X | present |
| 9 | 157 | yes | X | present |
| 10 | 152 | yes | X | present |
| 11 | 181 | yes | X | present |
| 12 | 190 | yes | X | present |
| 13 | 186 | no | Y | present |
| 14 | 194 | no | Y | present |
| 15 | 201 | no | Y | present |
| 16 | 215 | no | Y | present |
| 17 | 219 | no | Y | present |
| 18 | 209 | no | Y | present |
| 19 | 164 | yes | Y | present |
| 20 | 166 | yes | Y | present |

Hypertension Example: OLS Estimation (in R)

Effect coding for drug and diet:

```
> contrasts(hyper$drug) <- contr.sum(3)
> contrasts(hyper$drug)
      [,1] [,2]
X       1    0
Y       0    1
Z      -1   -1
> contrasts(hyper$diet) <- contr.sum(2)
> contrasts(hyper$diet)
      [,1]
no       1
yes      -1
> mymod=lm(bp~drug*diet, data=hyper)
> summary(mymod)      # I deleted some output
```

Coefficients:

| | Estimate | Std. Error | t value | Pr(> t) |
|-------------|----------|------------|---------|--------------|
| (Intercept) | 184.500 | 1.642 | 112.355 | < 2e-16 *** |
| drug1 | -10.000 | 2.322 | -4.306 | 5.64e-05 *** |
| drug2 | 6.250 | 2.322 | 2.691 | 0.00901 ** |
| diet1 | 8.500 | 1.642 | 5.176 | 2.30e-06 *** |
| drug1:diet1 | -5.000 | 2.322 | -2.153 | 0.03498 * |
| drug2:diet1 | 2.750 | 2.322 | 1.184 | 0.24059 |

```
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
Residual standard error: 13.93 on 66 degrees of freedom
Multiple R-squared:  0.4329, Adjusted R-squared:  0.3899
F-statistic: 10.07 on 5 and 66 DF,  p-value: 3.385e-07
```

Hypertension Example: Sums-of-Squares (by hand 1)

Defining $n = \sum_{k=1}^b \sum_{j=1}^a n_{jk}$, the relevant sums-of-squares are

$$\begin{aligned} SST &= \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} (y_{ijk} - \bar{y}_{\dots})^2 = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} y_{ijk}^2 - \frac{1}{n} \left(\sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} y_{ijk} \right)^2 \\ &= 2473492 - \frac{1}{72} (13284)^2 = 22594 \end{aligned}$$

$$\begin{aligned} SSE &= \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} (y_{ijk} - \bar{y}_{\cdot jk})^2 = \sum_{k=1}^b \sum_{j=1}^a \sum_{i=1}^{n_{jk}} y_{ijk}^2 - \sum_{k=1}^b \sum_{j=1}^a \frac{\left(\sum_{i=1}^{n_{jk}} y_{ijk} \right)^2}{n_{jk}} \\ &= 2473492 - \left[2136^2 + 2052^2 + 2424^2 + 2154^2 + 2388^2 + 2130^2 \right] / 12 = 12814 \end{aligned}$$

$$SSR = SST - SSE = 22594 - 12814 = 9780$$

Hypertension Example: Sums-of-Squares (by hand 2)

The sums-of-squares for the main and interaction effects are given by

$$\begin{aligned} SSA &= bn_* \sum_{j=1}^a (\bar{y}_{\cdot j} - \bar{y}_{\dots})^2 = bn_* \sum_{j=1}^a \hat{\alpha}_j^2 \\ &= (2)(12) \left[(-10)^2 + 6.25^2 + 3.75^2 \right] = 3675 \end{aligned}$$

$$\begin{aligned} SSB &= an_* \sum_{k=1}^b (\bar{y}_{\cdot k} - \bar{y}_{\dots})^2 = an_* \sum_{k=1}^b \hat{\beta}_k^2 \\ &= (3)(12) \left[(-8.5)^2 + 8.5^2 \right] = 5202 \end{aligned}$$

$$\begin{aligned} SSAB &= n_* \sum_{k=1}^b \sum_{j=1}^a (\bar{y}_{jk} - \bar{y}_{\cdot j} - \bar{y}_{\cdot k} + \bar{y}_{\dots})^2 = n_* \sum_{k=1}^b \sum_{j=1}^a \hat{\gamma}_{jk}^2 \\ &= 12 \left[(-5)^2 + 5^2 + 2.75^2 + (-2.75)^2 + 2.25^2 + (-2.25)^2 \right] = 903 \end{aligned}$$

and since $n_{jk} = n_* = 12 \forall j, k$, we have

$$\begin{aligned} SSR &= SSA + SSB + SSAB \\ 9780 &= 3675 + 5202 + 903 \end{aligned}$$

Hypertension Example: ANOVA Table (by hand)

Putting things together in ANOVA table:

| Source | SS | df | MS | F | p-value |
|--------|-------|----|--------|-------|---------|
| SSR | 9780 | 5 | 1956.0 | 10.07 | < .0001 |
| SSA | 3675 | 2 | 1837.5 | 9.46 | 0.0002 |
| SSB | 5202 | 1 | 5202.0 | 26.79 | < .0001 |
| SSAB | 903 | 2 | 451.5 | 2.33 | 0.1057 |
| SSE | 12814 | 66 | 194.2 | | |
| SST | 22594 | 71 | | | |

Hypertension Example: ANOVA Table (in R)

```
> anova(mymod)
```

```
Analysis of Variance Table
```

```
Response: bp
```

| | Df | Sum Sq | Mean Sq | F value | Pr(>F) |
|-----------|----|--------|---------|---------|---------------|
| drug | 2 | 3675 | 1837.5 | 9.4643 | 0.0002433 *** |
| diet | 1 | 5202 | 5202.0 | 26.7935 | 2.305e-06 *** |
| drug:diet | 2 | 903 | 451.5 | 2.3255 | 0.1056925 |
| Residuals | 66 | 12814 | 194.2 | | |

```
---
```

```
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Multiple Comparisons Overview

Still have multiple comparison problem:

- Overall test is not very informative
- Can examine effect estimates for group differences
- Need follow-up tests to examine linear combinations of means

Still can use the same tools as before:

- Bonferroni
- Tukey (Tukey-Kramer)
- Scheffé

Two-Way ANOVA Linear Combinations

Assuming interaction model, we now have

$$\hat{L} = \sum_{k=1}^b \sum_{j=1}^a c_{jk} \bar{y}_{.jk} \quad \text{and} \quad \hat{V}(\hat{L}) = \hat{\sigma}^2 \sum_{k=1}^b \sum_{j=1}^a c_{jk}^2 / n_{jk}$$

where c_{jk} are the coefficients and $\hat{\sigma}^2$ is the MSE.

Assuming the additive model, we have

$$\begin{aligned} \hat{L}_a &= \sum_{j=1}^a c_j \bar{y}_{.j} & \text{and} & & \hat{V}(\hat{L}_a) &= \hat{\sigma}^2 \sum_{j=1}^a c_j^2 / n_{j.} \\ \hat{L}_b &= \sum_{k=1}^b c_k \bar{y}_{..k} & \text{and} & & \hat{V}(\hat{L}_b) &= \hat{\sigma}^2 \sum_{k=1}^b c_k^2 / n_{.k} \end{aligned}$$

where c_j and c_k are main effect coefficients, $\hat{\sigma}^2$ is the MSE, and $n_{j.} = \sum_{k=1}^b n_{jk}$ and $n_{.k} = \sum_{j=1}^a n_{jk}$ are the marginal sample sizes.

Two-Way Multiple Comparisons in Practice

For interaction model, you follow-up on $\hat{\mu}_{jk} = \bar{y}_{.jk}$

- Bonferroni for any f tests (independent or not)
- Tukey (Tukey-Kramer) for all pairwise comparisons
- Scheffé for all possible contrasts

For additive model, you follow-up on $\hat{\mu}_j = \bar{y}_{.j}$ and $\hat{\mu}_k = \bar{y}_{..k}$

- Bonferroni for any f tests (independent or not)
- Tukey (Tukey-Kramer) for all pairwise comparisons
- Scheffé for all possible contrasts

For additive model, Tukey and Scheffé control FWER for each main effect family separately.

- Use Bonferroni in combination with Tukey/Scheffé to control FWER for both families simultaneously

Hypertension Example: Interaction (by hand)

All $ab(ab - 1)/2 = 15$ possible pairwise comparisons of $\hat{\mu}_{jk}$:

$$\hat{L} = \bar{y}_{.jk} - \bar{y}_{.j'k'} \quad \text{and} \quad \hat{V}(\hat{L}) = 194.2(2/12) = 32.36667$$

and we know that $\frac{\sqrt{2}(\hat{L})}{\sqrt{\hat{V}(\hat{L})}} \sim q_{ab, abn_* - ab}$, so $100(1 - \alpha)\%$ CI is given by

$$\hat{L} \pm \frac{1}{\sqrt{2}} q_{ab, abn_* - ab}^{(\alpha)} \sqrt{\hat{V}(\hat{L})}$$

where $q_{ab, abn_* - ab}^{(\alpha)}$ is critical value from studentized range.

For example, 95% CI for $\mu_{21} - \mu_{11}$ is given by:

$$\begin{aligned} & (\hat{\mu}_{21} - \hat{\mu}_{11}) \pm \frac{1}{\sqrt{2}} q_{6,66}^{(.05)} \sqrt{\hat{V}(\hat{L})} \\ & \left(\frac{2424}{12} - \frac{2136}{12} \right) \pm \frac{1}{\sqrt{2}} (4.150851) \sqrt{32.36667} = [7.303829; 40.69617] \end{aligned}$$

Hypertension Example: Interaction (in R)

All $ab(ab - 1)/2 = 15$ possible pairwise comparisons of $\hat{\mu}_{jk}$:

```
> mymod=aov(bp~drug*diet,data=hyper)
```

```
> TukeyHSD(mymod,"drug:diet")
```

```
  Tukey multiple comparisons of means
    95% family-wise confidence level
```

```
Fit: aov(formula = bp ~ drug * diet, data = hyper)
```

```
$`drug:diet`
```

| | diff | lwr | upr | p adj |
|-------------|-------|------------|------------|-----------|
| Y:no-X:no | 24.0 | 7.303829 | 40.696171 | 0.0010415 |
| Z:no-X:no | 21.0 | 4.303829 | 37.696171 | 0.0058124 |
| X:yes-X:no | -7.0 | -23.696171 | 9.696171 | 0.8203137 |
| Y:yes-X:no | 1.5 | -15.196171 | 18.196171 | 0.9998189 |
| Z:yes-X:no | -0.5 | -17.196171 | 16.196171 | 0.9999992 |
| Z:no-Y:no | -3.0 | -19.696171 | 13.696171 | 0.9948741 |
| X:yes-Y:no | -31.0 | -47.696171 | -14.303829 | 0.0000117 |
| Y:yes-Y:no | -22.5 | -39.196171 | -5.803829 | 0.0025081 |
| Z:yes-Y:no | -24.5 | -41.196171 | -7.803829 | 0.0007710 |
| X:yes-Z:no | -28.0 | -44.696171 | -11.303829 | 0.0000856 |
| Y:yes-Z:no | -19.5 | -36.196171 | -2.803829 | 0.0128988 |
| Z:yes-Z:no | -21.5 | -38.196171 | -4.803829 | 0.0044123 |
| Y:yes-X:yes | 8.5 | -8.196171 | 25.196171 | 0.6690751 |
| Z:yes-X:yes | 6.5 | -10.196171 | 23.196171 | 0.8616371 |
| Z:yes-Y:yes | -2.0 | -18.696171 | 14.696171 | 0.9992610 |

Hypertension Example: Additive (by hand A part 1)

All $a(a-1)/2 = 3$ possible pairwise comparisons of $\hat{\mu}_j$:

$$Y - X : \quad \hat{L}_{a_1} = \frac{4578}{24} - \frac{4188}{24} = 16.25$$

$$Z - X : \quad \hat{L}_{a_2} = \frac{4518}{24} - \frac{4188}{24} = 13.75$$

$$Z - Y : \quad \hat{L}_{a_3} = \frac{4518}{24} - \frac{4578}{24} = -2.5$$

and the variance is given by

$$\hat{V}(\hat{L}_{a_j}) = \hat{\sigma}^2 \sum_{j=1}^a c_j^2 / n_j = (201.7206)(2/24) = 16.81005$$

$$\text{where } \hat{\sigma}^2 = \frac{SSE + SSAB}{abn_* - (a+b-1)} = \frac{12814 + 903}{68} = 201.7206$$

Hypertension Example: Additive (by hand A part 2)

Note $\frac{\sqrt{2}(\hat{L}_{a_j})}{\sqrt{\hat{V}(\hat{L}_{a_j})}} \sim q_{a,abn_*(a+b-1)}$, so $100(1 - \alpha)\%$ CI is given by

$$\hat{L}_{a_j} \pm \frac{1}{\sqrt{2}} q_{a,abn_*(a+b-1)}^{(\alpha)} \sqrt{\hat{V}(\hat{L}_{a_j})}$$

where $q_{a,abn_*(a+b-1)}^{(\alpha)}$ is critical value from studentized range.

The 95% CI for all three pairwise comparisons is given by

$$\begin{aligned} \hat{L}_{a_1} \pm \frac{1}{\sqrt{2}} q_{3,68}^{(.05)} \sqrt{\hat{V}(\hat{L}_{a_1})} &= 16.25 \pm \frac{1}{\sqrt{2}} (3.388576) \sqrt{16.81005} = [6.426037; 26.07396] \\ \hat{L}_{a_2} \pm \frac{1}{\sqrt{2}} q_{3,68}^{(.05)} \sqrt{\hat{V}(\hat{L}_{a_2})} &= 13.75 \pm \frac{1}{\sqrt{2}} (3.388576) \sqrt{16.81005} = [3.926037; 23.57396] \\ \hat{L}_{a_3} \pm \frac{1}{\sqrt{2}} q_{3,68}^{(.05)} \sqrt{\hat{V}(\hat{L}_{a_3})} &= -2.5 \pm \frac{1}{\sqrt{2}} (3.388576) \sqrt{16.81005} = [-12.32396; 7.323963] \end{aligned}$$

Hypertension Example: Additive (by hand B part 1)

All $b(b-1)/2 = 1$ possible pairwise comparison of $\hat{\mu}_k$:

$$\text{yes} - \text{no} : \quad \hat{L}_b = \frac{6336}{36} - \frac{6948}{36} = -17$$

and the variance is given by

$$\hat{V}(\hat{L}_b) = \hat{\sigma}^2 \sum_{k=1}^b c_k^2 / n_{.k} = (201.7206)(2/36) = 11.2067$$

$$\text{where } \hat{\sigma}^2 = \frac{SSE + SSAB}{abn_* - (a+b-1)} = \frac{12814 + 903}{68} = 201.7206$$

Hypertension Example: Additive (by hand B part 2)

Note $\frac{\sqrt{2}(\hat{L}_b)}{\sqrt{\hat{V}(\hat{L}_b)}} \sim q_{b,abn_*(a+b-1)}$, so $100(1 - \alpha)\%$ CI is given by

$$\hat{L}_b \pm \frac{1}{\sqrt{2}} q_{b,abn_*(a+b-1)}^{(\alpha)} \sqrt{\hat{V}(\hat{L}_b)}$$

where $q_{b,abn_*(a+b-1)}^{(\alpha)}$ is critical value from studentized range.

The 95% CI for pairwise comparison is given by

$$\begin{aligned} \hat{L}_b \pm \frac{1}{\sqrt{2}} q_{2,68}^{(.05)} \sqrt{\hat{V}(\hat{L}_b)} &= -17 \pm \frac{1}{\sqrt{2}} (2.822019) \sqrt{11.2067} \\ &= [-23.68011; -10.31989] \end{aligned}$$

Hypertension Example: Additive (in R)

All $a(a - 1)/2 = 3$ possible pairwise comparisons of $\hat{\mu}_j$:

```
> mymod=aov(bp~drug+diet,data=hyper)
> TukeyHSD(mymod,"drug")
  Tukey multiple comparisons of means
    95% family-wise confidence level
```

```
Fit: aov(formula = bp ~ drug + diet, data = hyper)
```

```
$drug
      diff      lwr      upr      p adj
Y-X 16.25    6.426037 26.073963 0.0005220
Z-X 13.75    3.926037 23.573963 0.0036941
Z-Y -2.50 -12.323963  7.323963 0.8152941
```

All $b(b - 1)/2 = 1$ possible pairwise comparison of $\hat{\mu}_k$:

```
> mymod=aov(bp~drug+diet,data=hyper)
> TukeyHSD(mymod,"diet")
  Tukey multiple comparisons of means
    95% family-wise confidence level
```

```
Fit: aov(formula = bp ~ drug + diet, data = hyper)
```

```
$diet
      diff      lwr      upr      p adj
yes-no -17 -23.68011 -10.31989 3.2e-06
```

3-Way ANOVA Model (cell means form)

The 3-Way Analysis of Variance (ANOVA) model has the form

$$y_{ijkl} = \mu_{jkl} + e_{ijkl}$$

for $i \in \{1, \dots, n_{jkl}\}$, $j \in \{1, \dots, a\}$, $k \in \{1, \dots, b\}$, $l \in \{1, \dots, c\}$, where

- $y_{ijkl} \in \mathbb{R}$ is response for i -th subject in factor cell (j, k, l)
- $\mu_{jkl} \in \mathbb{R}$ is population mean for factor cell (j, k, l)
- $e_{ijkl} \stackrel{\text{iid}}{\sim} N(0, \sigma^2)$ is Gaussian measurement error
- n_{jkl} is number of subjects in cell (j, k, l)
(note: $n_{jkl} = n_* \forall j, k, l$ in balanced 3-way ANOVA)
- (a, b, c) is number of factor levels for Factors (A, B, C)

Implies that $y_{ijkl} \stackrel{\text{ind}}{\sim} N(\mu_{jkl}, \sigma^2)$.

3-Way ANOVA Model (all interactions)

The 3-Way ANOVA with all interactions assumes that

$$\mu_{jkl} = \mu + \alpha_j + \beta_k + \gamma_l + \delta_{jk} + \zeta_{jl} + \eta_{kl} + \theta_{jkl}$$

for $j \in \{1, \dots, a\}$, $k \in \{1, \dots, b\}$, and $l \in \{1, \dots, c\}$ where

- μ is overall population mean
- α_j is main effect of first factor such that $\sum_{j=1}^a \alpha_j = 0$
- β_k is main effect of second factor such that $\sum_{k=1}^b \beta_k = 0$
- γ_l is main effect of third factor such that $\sum_{l=1}^c \gamma_l = 0$
- δ_{jk} is interaction between factors A and B such that $\sum_{j=1}^a \delta_{jk} = 0 \forall k$ and $\sum_{k=1}^b \delta_{jk} = 0 \forall j$
- ζ_{jl} is interaction between factors A and C such that $\sum_{j=1}^a \zeta_{jl} = 0 \forall l$ and $\sum_{l=1}^c \zeta_{jl} = 0 \forall j$
- η_{kl} is interaction between factors B and C such that $\sum_{k=1}^b \zeta_{kl} = 0 \forall l$ and $\sum_{l=1}^c \eta_{kl} = 0 \forall k$
- θ_{jkl} is 3-way interaction such that $\sum_{j=1}^a \theta_{jkl} = 0 \forall k, l$ and $\sum_{k=1}^b \theta_{jkl} = 0 \forall j, l$ and $\sum_{l=1}^c \theta_{jkl} = 0 \forall j, k$

3-Way ANOVA Model (all 2-way interactions)

The 3-Way ANOVA with all two-way interactions assumes that

$$\mu_{jkl} = \mu + \alpha_j + \beta_k + \gamma_l + \delta_{jk} + \zeta_{jl} + \eta_{kl}$$

for $j \in \{1, \dots, a\}$, $k \in \{1, \dots, b\}$, and $l \in \{1, \dots, c\}$ where

- μ is overall population mean
- α_j is main effect of first factor such that $\sum_{j=1}^a \alpha_j = 0$
- β_k is main effect of second factor such that $\sum_{k=1}^b \beta_k = 0$
- γ_l is main effect of third factor such that $\sum_{l=1}^c \gamma_l = 0$
- δ_{jk} is interaction between factors A and B such that $\sum_{j=1}^a \delta_{jk} = 0 \forall k$ and $\sum_{k=1}^b \delta_{jk} = 0 \forall j$
- ζ_{jl} is interaction between factors A and C such that $\sum_{j=1}^a \zeta_{jl} = 0 \forall l$ and $\sum_{l=1}^c \zeta_{jl} = 0 \forall j$
- η_{kl} is interaction between factors B and C such that $\sum_{k=1}^b \zeta_{kl} = 0 \forall l$ and $\sum_{l=1}^c \eta_{kl} = 0 \forall k$

3-Way ANOVA Model (additive)

The 3-Way ANOVA with no interactions assumes that

$$\mu_{jkl} = \mu + \alpha_j + \beta_k + \gamma_l$$

for $j \in \{1, \dots, a\}$, $k \in \{1, \dots, b\}$, and $l \in \{1, \dots, c\}$ where

- μ is overall population mean
- α_j is main effect of first factor such that $\sum_{j=1}^a \alpha_j = 0$
- β_k is main effect of second factor such that $\sum_{k=1}^b \beta_k = 0$
- γ_l is main effect of third factor such that $\sum_{l=1}^c \gamma_l = 0$

Memory Example: Data Description (revisited)

Hypertension example from Maxwell & Delany (2003).

Total of $n = 72$ subjects participate in hypertension experiment.

- Factor A: `drug` type ($a = 3$ levels: X, Y, Z)
- Factor B: `diet` type ($b = 2$ levels: yes, no)
- Factor C: `biof` type ($c = 2$ levels: present, absent)

Randomly assign $n_{jkl} = 6$ subjects to each treatment cell:

- Note there are $(abc) = (3)(2)(2) = 12$ treatment cells
- Observations are independent within and between cells

Hypertension Example: Look at Data

```
> bp=scan("/Users/Nate/Desktop/hypertension.dat")
Read 72 items
> diet=factor(rep(rep(c("no", "yes"), each=6), 6))
> drug=factor(rep(rep(c("X", "Y", "Z"), each=12), 2))
> biof=factor(rep(c("present", "absent"), each=36))
> hyper=data.frame(bp=bp, diet=diet, drug=drug, biof=biof)
> hyper[1:20,]
```

| | bp | diet | drug | biof |
|----|-----|------|------|---------|
| 1 | 170 | no | X | present |
| 2 | 175 | no | X | present |
| 3 | 165 | no | X | present |
| 4 | 180 | no | X | present |
| 5 | 160 | no | X | present |
| 6 | 158 | no | X | present |
| 7 | 161 | yes | X | present |
| 8 | 173 | yes | X | present |
| 9 | 157 | yes | X | present |
| 10 | 152 | yes | X | present |
| 11 | 181 | yes | X | present |
| 12 | 190 | yes | X | present |
| 13 | 186 | no | Y | present |
| 14 | 194 | no | Y | present |
| 15 | 201 | no | Y | present |
| 16 | 215 | no | Y | present |
| 17 | 219 | no | Y | present |
| 18 | 209 | no | Y | present |
| 19 | 164 | yes | Y | present |
| 20 | 166 | yes | Y | present |

Hypertension Example: All Interactions

```
> contrasts(hyper$drug) <- contr.sum(3)
> contrasts(hyper$diet) <- contr.treatment(2, base=1)
> contrasts(hyper$biof) <- contr.treatment(2, base=1)
> mymod = lm(bp ~ drug * diet * biof, data = hyper)
> anova(mymod)
```

Analysis of Variance Table

Response: bp

| | Df | Sum Sq | Mean Sq | F value | Pr(>F) | |
|----------------|----|--------|---------|---------|-----------|-----|
| drug | 2 | 3675 | 1837.5 | 11.7287 | 5.019e-05 | *** |
| diet | 1 | 5202 | 5202.0 | 33.2043 | 3.053e-07 | *** |
| biof | 1 | 2048 | 2048.0 | 13.0723 | 0.0006151 | *** |
| drug:diet | 2 | 903 | 451.5 | 2.8819 | 0.0638153 | . |
| drug:biof | 2 | 259 | 129.5 | 0.8266 | 0.4424565 | |
| diet:biof | 1 | 32 | 32.0 | 0.2043 | 0.6529374 | |
| drug:diet:biof | 2 | 1075 | 537.5 | 3.4309 | 0.0388342 | * |
| Residuals | 60 | 9400 | 156.7 | | | |

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Hypertension Example: All 2-Way Interactions

```
> contrasts(hyper$drug) <- contr.sum(3)
> contrasts(hyper$diet) <- contr.treatment(2, base=1)
> contrasts(hyper$biof) <- contr.treatment(2, base=1)
> mymod = lm(bp ~ drug * diet + drug * biof + diet * biof, data = hyper)
> anova(mymod)
```

Analysis of Variance Table

Response: bp

| | Df | Sum Sq | Mean Sq | F value | Pr(>F) | |
|-----------|----|--------|---------|---------|-----------|-----|
| drug | 2 | 3675 | 1837.5 | 10.8759 | 8.940e-05 | *** |
| diet | 1 | 5202 | 5202.0 | 30.7899 | 6.345e-07 | *** |
| biof | 1 | 2048 | 2048.0 | 12.1218 | 0.000919 | *** |
| drug:diet | 2 | 903 | 451.5 | 2.6724 | 0.077043 | . |
| drug:biof | 2 | 259 | 129.5 | 0.7665 | 0.468992 | |
| diet:biof | 1 | 32 | 32.0 | 0.1894 | 0.664925 | |
| Residuals | 62 | 10475 | 169.0 | | | |

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Hypertension Example: Additive Model

```
> contrasts(hyper$drug) <- contr.sum(3)
> contrasts(hyper$diet) <- contr.treatment(2, base=1)
> contrasts(hyper$biof) <- contr.treatment(2, base=1)
> mymod = lm(bp ~ drug + diet + biof, data = hyper)
> anova(mymod)
```

Analysis of Variance Table

Response: bp

| | Df | Sum Sq | Mean Sq | F value | Pr(>F) | |
|-----------|----|--------|---------|---------|-----------|-----|
| drug | 2 | 3675 | 1837.5 | 10.550 | 0.0001039 | *** |
| diet | 1 | 5202 | 5202.0 | 29.868 | 7.346e-07 | *** |
| biof | 1 | 2048 | 2048.0 | 11.759 | 0.0010403 | ** |
| Residuals | 67 | 11669 | 174.2 | | | |

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Hypertension Example: Multiple Comparisons

Assuming we chose the additive model, we would perform follow-up tests on the marginal means.

- Factor A: $\hat{\mu}_{a_j} = \hat{\mu} + \hat{\alpha}_j$
- Factor B: $\hat{\mu}_{b_k} = \hat{\mu} + \hat{\beta}_k$
- Factor C: $\hat{\mu}_{c_l} = \hat{\mu} + \hat{\gamma}_l$

If we chose the 3-way interaction model, we would perform follow-up tests on the individual cell means.

$$\hat{\mu}_{jkl} = \hat{\mu} + \hat{\alpha}_j + \hat{\beta}_k + \hat{\gamma}_l + \hat{\delta}_{jk} + \hat{\zeta}_{jl} + \hat{\eta}_{kl} + \hat{\theta}_{jkl}$$

Hypertension Example: Multiple Comparisons

```
> mymod=aov(bp~drug+diet+biof,data=hyper)

> TukeyHSD(mymod,"drug")      # I deleted some output
  Tukey multiple comparisons of means
    95% family-wise confidence level

$drug
      diff          lwr          upr      p adj
Y-X 16.25    7.118642 25.381358 0.0001874
Z-X 13.75    4.618642 22.881358 0.0016810
Z-Y -2.50  -11.631358  6.631358 0.7894946

> TukeyHSD(mymod,"diet")      # I deleted some output
  Tukey multiple comparisons of means
    95% family-wise confidence level

$diet
      diff          lwr          upr p adj
yes-no  -17 -23.20877 -10.79123 7e-07

> TukeyHSD(mymod,"biof")      # I deleted some output
  Tukey multiple comparisons of means
    95% family-wise confidence level

$biof
      diff          lwr          upr      p adj
present-absent -10.66667 -16.87544 -4.457897 0.0010403
```

Unbalanced ANOVA: Model Form

Unbalanced ANOVA has same model form as balanced, but unequal sample sizes in each cell.

- 1-way: $n_j \neq n_{j'}$ for some j, j'
- 2-way: $n_{jk} \neq n_{j'k'}$ for some $(jk), (j'k')$
- 3-way: $n_{jkl} \neq n_{j'k'l'}$ for some $(jkl), (j'k'l')$

Consequences for 2-way (and higher way) unbalanced design:

- Parameter estimates are not simple cell means
- Non-orthogonal SS (e.g., $SSR \neq SSA + SSB + SSAB$)

Unbalanced ANOVA: Testing Effects

Because of non-orthogonality, cannot test effects using $F = \frac{MS?}{MSE}$.

Consider 2-way ANOVA and all 7 possible models

$$y_{ijk} = \mu + \alpha_j + \beta_k + \gamma_{jk} + \mathbf{e}_{ijk} \quad (1)$$

$$y_{ijk} = \mu + \alpha_j + \beta_k + \mathbf{e}_{ijk} \quad (2)$$

$$y_{ijk} = \mu + \alpha_j + \gamma_{jk} + \mathbf{e}_{ijk} \quad (3)$$

$$y_{ijk} = \mu + \beta_k + \gamma_{jk} + \mathbf{e}_{ijk} \quad (4)$$

$$y_{ijk} = \mu + \alpha_j + \mathbf{e}_{ijk} \quad (5)$$

$$y_{ijk} = \mu + \beta_k + \mathbf{e}_{ijk} \quad (6)$$

$$y_{ijk} = \mu + \mathbf{e}_{ijk} \quad (7)$$

Unbalanced ANOVA: Testing Effects (continued)

To test effect, use F test comparing full and reduced models.

To test each effect there are multiple choices we could use for full and reduced models:

- A: $F=1$ and $R=4$ or $F=2$ and $R=6$ or $F=5$ and $R=7$
- B: $F=1$ and $R=3$ or $F=2$ and $R=5$ or $F=6$ and $R=7$
- AB: $F=1$ and $R=2$ or $F=3$ and $R=5$ or $F=4$ and $R=6$

Types of Sum-of-Squares

Type I SS

- Amount of additional variation explained by the model when a term is added to the model (aka *sequential sum-of-squares*).
- In two-way ANOVA, type I SS would compare:
 - (a) Main Effect A: $F=5$ and $R=7$
 - (b) Main Effect B: $F=2$ and $R=5$
 - (c) Interaction Effect: $F=1$ and $R=2$

Type II SS

- Amount of additional variation explained by the model when a term and all associated interactions are added to the model.
- In two-way ANOVA, type II SS would compare:
 - (a) Main Effect A: $F=2$ and $R=6$
 - (b) Main Effect B: $F=2$ and $R=5$
 - (c) Interaction Effect: $F=1$ and $R=2$

Type III SS

- Amount of variation a term adds to the model when all other terms are included, which is sometimes called *partial sum-of-squares*.
- In two-way ANOVA, type III SS would compare:
 - (a) Main Effect A: $F=1$ and $R=4$
 - (b) Main Effect B: $F=1$ and $R=3$
 - (c) Interaction Effect: $F=1$ and $R=2$

Types of Sum-of-Squares (in R)

When fitting multi-way ANOVAs, `anova` function gives Type I SS.

- Order matters in unbalanced design!
- `bp=drug+diet` produces different Type I SS tests than `bp=diet+drug` if design is unbalanced

Use `Anova` function in `car` package for Type II and Type III SS.

- Function performs Type II SS tests by default
- Use `type=3` option for Type III SS tests

Unbalanced ANOVA: Estimation and Inference

To estimate parameters, just use MLR approach:

$$\hat{\mathbf{b}} = (\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{y}$$

where \mathbf{X} is design matrix and \mathbf{b} contains effects.

To perform multiple comparisons, same approach but use least-squares means. For example, 3-way additive would use

- Factor A: $\hat{\mu}_{a_j} = \hat{\mu} + \hat{\alpha}_j$
- Factor B: $\hat{\mu}_{b_k} = \hat{\mu} + \hat{\beta}_k$
- Factor C: $\hat{\mu}_{c_l} = \hat{\mu} + \hat{\gamma}_l$

where $\hat{\mu}$, $\hat{\alpha}_j$, $\hat{\beta}_k$, and $\hat{\gamma}_l$ are least-squares estimates.

Hypertension Example: Type I

```
> contrasts(hyper$drug) <- contr.sum(3)
> contrasts(hyper$diet) <- contr.treatment(2, base=1)
> contrasts(hyper$biof) <- contr.treatment(2, base=1)
> mymod = lm(bp ~ drug * diet * biof, data = hyper[1:71,])
> anova(mymod)
```

Analysis of Variance Table

Response: bp

| | Df | Sum Sq | Mean Sq | F value | Pr(>F) | |
|----------------|----|--------|---------|---------|-----------|-----|
| drug | 2 | 3733.6 | 1866.8 | 11.7306 | 5.138e-05 | *** |
| diet | 1 | 5113.3 | 5113.3 | 32.1311 | 4.558e-07 | *** |
| biof | 1 | 2087.2 | 2087.2 | 13.1154 | 0.0006101 | *** |
| drug:diet | 2 | 879.5 | 439.8 | 2.7633 | 0.0712569 | . |
| drug:biof | 2 | 280.5 | 140.3 | 0.8813 | 0.4196123 | |
| diet:biof | 1 | 24.2 | 24.2 | 0.1522 | 0.6978384 | |
| drug:diet:biof | 2 | 1055.8 | 527.9 | 3.3172 | 0.0431275 | * |
| Residuals | 59 | 9389.2 | 159.1 | | | |

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Hypertension Example: Type II

```
> library(car)
> Anova(mymod,type=2)
Anova Table (Type II tests)
```

Response: bp

| | Sum Sq | Df | F value | Pr(>F) | |
|----------------|--------|----|---------|-----------|-----|
| drug | 3704.1 | 2 | 11.6378 | 5.491e-05 | *** |
| diet | 4975.9 | 1 | 31.2676 | 6.085e-07 | *** |
| biof | 2061.8 | 1 | 12.9561 | 0.0006541 | *** |
| drug:diet | 872.5 | 2 | 2.7413 | 0.0727049 | . |
| drug:biof | 277.7 | 2 | 0.8726 | 0.4231893 | |
| diet:biof | 24.2 | 1 | 0.1522 | 0.6978384 | |
| drug:diet:biof | 1055.8 | 2 | 3.3172 | 0.0431275 | * |
| Residuals | 9389.2 | 59 | | | |

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Hypertension Example: Type III

```
> library(car)
> Anova(mymod, type=3)
Anova Table (Type III tests)
```

Response: bp

| | Sum Sq | Df | F value | Pr(>F) | |
|----------------|--------|----|-----------|-----------|-----|
| (Intercept) | 712818 | 1 | 4479.2168 | < 2.2e-16 | *** |
| drug | 1332 | 2 | 4.1850 | 0.019969 | * |
| diet | 2864 | 1 | 17.9962 | 7.92e-05 | *** |
| biof | 1296 | 1 | 8.1438 | 0.005951 | ** |
| drug:diet | 294 | 2 | 0.9247 | 0.402336 | |
| drug:biof | 1152 | 2 | 3.6195 | 0.032907 | * |
| diet:biof | 27 | 1 | 0.1692 | 0.682287 | |
| drug:diet:biof | 1056 | 2 | 3.3172 | 0.043127 | * |
| Residuals | 9389 | 59 | | | |

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Kruskal-Wallis Test: Overview

Suppose data from one-way ANOVA situation, but $y_{ij} \not\sim N(\mu_j, \sigma^2)$

- Maybe Y is not normally distributed
- And/or maybe Y has heterogeneous variance

Can still test location differences between groups.

- $H_0 : \tilde{\mu}_j = \tilde{\mu} \forall j$ versus $H_1 : \text{not all } \tilde{\mu}_j = \tilde{\mu}$
- $\tilde{\mu}_j$ is population for j -th factor level

Analyze rank data instead of raw magnitude data.

Kruskal-Wallis Test: Test Statistic

Kruskal-Wallis test statistic is given by

$$K = (n - 1) \frac{\sum_{j=1}^g n_j (\bar{r}_{\cdot j} - \bar{r}_{..})^2}{\sum_{j=1}^g \sum_{i=1}^{n_j} (r_{ij} - \bar{r}_{..})^2}$$

where

- $r_{ij} \in \{1, \dots, n\}$ is rank of y_{ij}
- $n = \sum_{j=1}^g n_j$ is total sample size
- $\bar{r}_{\cdot j} = \frac{1}{n_j} \sum_{i=1}^{n_j} r_{ij}$ is average rank for j -th group
- $\bar{r}_{..} = \frac{1}{n} \sum_{j=1}^g \sum_{i=1}^{n_j} r_{ij}$

Asymptotically $K \sim \chi_{g-1}^2$ so use $\chi_{g-1(\alpha)}^2$ critical values for CIs.

Kruskal-Wallis Test: Memory Example

Revisiting the memory example:

```
> sync=c(23,27,23,22,28,24,18,33,21,15,
         19,25,29,25,19,30,29,24,23,36,
         22,16,30,17,19,26,20,17,21,23)
> cond=factor(rep(c("fast","normal","slow"),10))
> smod=lm(sync~cond,contrasts=list(cond=contr.sum))
> tapply(sync,cond,sd)
      fast      normal      slow
5.202563 5.103376 2.529822
> kruskal.test(sync~cond)
```

```
Kruskal-Wallis rank sum test
```

```
data:  sync by cond
Kruskal-Wallis chi-squared = 8.1309, df = 2, p-value = 0.01716
```

```
> srnk=rank(sync)
> top=(tapply(srnk,cond,mean)-mean(srnk))^2
> bot=(srnk-mean(srnk))^2
> 29*sum(10*top)/sum(bot)
[1] 8.13089
```