

Module 4: Factorial Treatment Structure

Analyzing a Factorial: Model, ANOVA, and Decision Flow

Example 4.2: Bakery

Treatment Structure: 3×2 Full Factorial

- Shelf Height (Bottom, Middle, Top)
- Shelf Width (Regular, Wide)

Design Structure: CRD with $r = 2$ stores per treatment combination.

Response: Bread Sales



What changes when we add a factor?

- We add:
 - Another main effect
 - An interaction term
- We must decide what effects are present *before* interpreting the treatment means.

The goal of the analysis is to decide which effects matter.

The Treatment Effects Model (Two-way Factorial)

$$y_{ijk} = \mu + \alpha_i + \beta_j + \alpha\beta_{ij} + \epsilon_{ijk} \text{ with } \epsilon_{ijk} \sim \text{iid } N(0, \sigma^2)$$

for $i = 1, 2, 3, \dots, a; j = 1, 2, \dots, b; k = 1, 2, \dots, r$

- y_{ijk} : is the response (sales) from the k^{th} experimental unit (store) using the i^{th} level of Factor A (height) and j^{th} level of Factor B (width) combination
- μ : is the grand/overall mean sales
- α_i : is the effect of the i^{th} level of factor A (height)
- β_j : is the effect of the j^{th} level of factor B (width)
- $\alpha\beta_{ij}$: is the interaction effect between the i^{th} level of A (height) and j^{th} level of B (width)
- ϵ_{ijk} : the experimental error associated with the k^{th} experimental unit (store) using the i^{th} level of Factor A (height) and j^{th} level of Factor B (width) combination

ANOVA for Two-way Factorials

Just like one-way ANOVA: $SST = SST_{rt} + SSE$

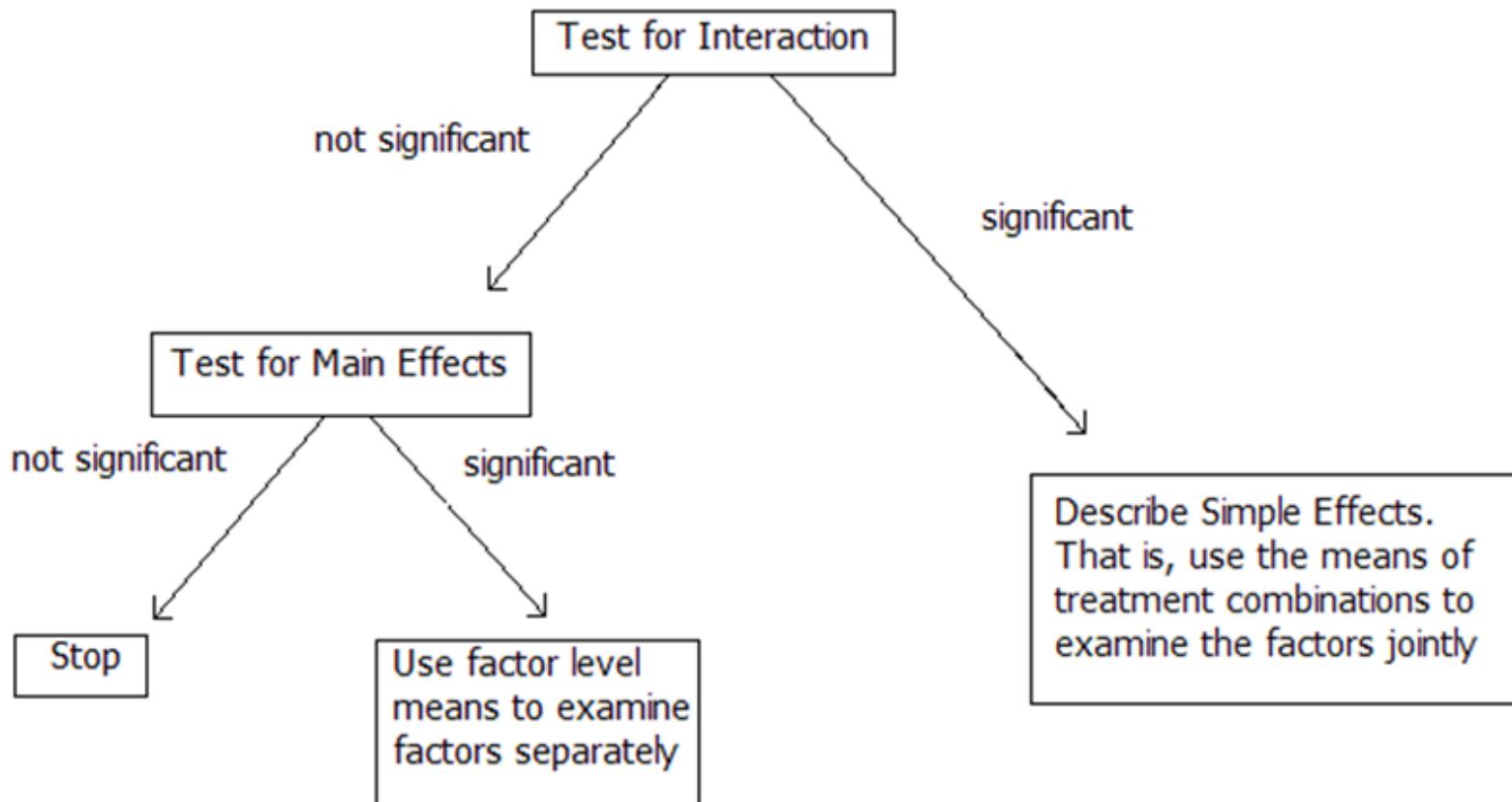
For a factorial: $SST_{rt} = SSA + SSB + SSAB$

- Main effect of factor A (Height): $SSA = rb \sum_i (\bar{y}_{i..} - \bar{y}_{...})^2$
- Main effect of factor B (Width): $SSB = ra \sum_j (\bar{y}_{.j.} - \bar{y}_{...})^2$
- AB Interaction Effect (Height x Width): $SSAB = SST - SSE - SSA - SSB$

Full ANOVA Table (Two-way)

| Source (SV) | df | SS | MS | F | |
|-----------------|------------------|------|----------------------------------|--------------------|--|
| A | $a - 1$ | SSA | $MSA = \frac{SSA}{a-1}$ | $\frac{MSA}{MSE}$ | Do avg sales differ across shelf heights? |
| B | $b - 1$ | SSB | $MSB = \frac{SSB}{b-1}$ | $\frac{MSB}{MSE}$ | Do avg sales differ across shelf widths? |
| AB | $(a - 1)(b - 1)$ | SSAB | $MSAB = \frac{SSAB}{(a-1)(b-1)}$ | $\frac{MSAB}{MSE}$ | Does the effect of width depend on height? |
| Error: e.u.(AB) | $(r - 1)ab$ | SSE | $MSE = \frac{SSE}{(r-1)ab}$ | | |
| Total | $N - 1$ | SST | | | |

Decision Flowchart



What are we testing?

Interaction

$$H_0 : \text{All } \alpha\beta_{ij} = 0 \text{ vs } H_A : \text{At least one } \alpha\beta_{ij} \neq 0$$

Main Effect of A

$$H_0 : \text{All } \alpha_i = 0 \text{ vs } H_A : \text{At least one } \alpha_i \neq 0$$

Main Effect of B

$$H_0 : \text{All } \beta_j = 0 \text{ vs } H_A : \text{At least one } \beta_j \neq 0$$

Example 4.2: Skeleton ANOVA

Source of Variation DF = 12 stores - 1 = 11 total df

R: Fitting the Model

Let's prep the data...

```
1 bakery_data <- read_csv("data/04_bakery_data.csv") %>%
2   mutate(height = factor(height, levels = c("bottom", "middle", "top")),
3         width = factor(width, levels = c("regular", "wide")))
4 head(bakery_data)
```

```
# A tibble: 6 × 4
  height width   sales placement
  <fct>  <fct>   <dbl> <chr>
1 bottom regular     47 bottomregular
2 bottom regular     43 bottomregular
3 bottom wide        46 bottomwide
4 bottom wide        40 bottomwide
5 middle regular    62 middleregular
6 middle regular    68 middleregular
```

```
1 levels(bakery_data$height)
```

```
[1] "bottom" "middle" "top"
```

```
1 levels(bakery_data$width)
```

```
[1] "regular" "wide"
```

R: Fitting the Model

```
1 options(contrasts = c("contr.sum", "contr.poly"))
2 bakery_mod <- lm(sales ~ height + width + height*width, data = bakery_data)
```

```
1 anova(bakery_mod)
```

Analysis of Variance Table

Response: sales

| | Df | Sum Sq | Mean Sq | F value | Pr(>F) |
|--------------|----|--------|---------|---------|---------------|
| height | 2 | 1544 | 772.00 | 74.7097 | 5.754e-05 *** |
| width | 1 | 12 | 12.00 | 1.1613 | 0.3226 |
| height*width | 2 | 24 | 12.00 | 1.1613 | 0.3747 |
| Residuals | 6 | 62 | 10.33 | | |

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

```
1 summary(bakery_mod)
```

Call:
lm(formula = sales ~ height + width + height*width, data = bakery_data)

Residuals:

| Min | 1Q | Median | 3Q | Max |
|-----|----|--------|----|-----|
| -3 | -2 | 0 | 2 | 3 |

Coefficients:

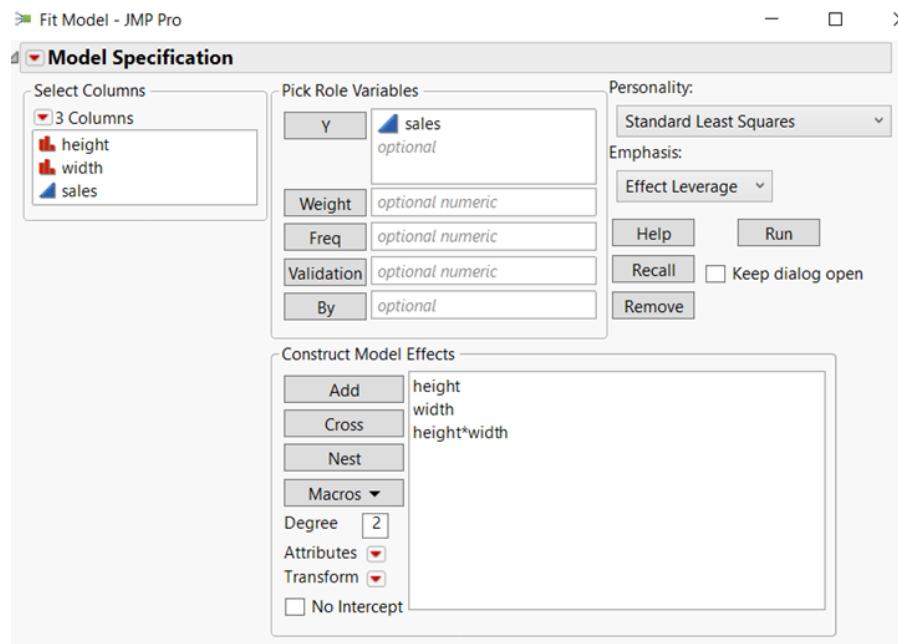
| | Estimate | Std. Error | t value | Pr(> t) |
|----------------|----------|------------|---------|--------------|
| (Intercept) | 51.000 | 0.928 | 54.959 | 2.44e-09 *** |
| height1 | -7.000 | 1.312 | -5.334 | 0.00177 ** |
| height2 | 16.000 | 1.312 | 12.192 | 1.85e-05 *** |
| width1 | -1.000 | 0.928 | -1.078 | 0.32261 |
| height1*width1 | 2.000 | 1.312 | 1.524 | 0.17835 |
| height2*width1 | -1.000 | 1.312 | -0.762 | 0.47494 |

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

Residual standard error: 3.215 on 6 degrees of freedom
Multiple R-squared: 0.9622, Adjusted R-squared: 0.9308
F-statistic: 30.58 on 5 and 6 DF, p-value: 0.0003384

JMP: Fitting the Model

► Analyze > Fit Model > Assign Y = Response + Highlight both treatment factors and click Macros > Full Factorial



| Source | DF | Sum of Squares | | |
|----------|----|----------------|---------|----------|
| | | Mean Square | F Ratio | Prob > F |
| Model | 5 | 1580.0000 | 316.000 | 30.5806 |
| Error | 6 | 62.0000 | 10.333 | 0.0003* |
| C. Total | 11 | 1642.0000 | | |

| Source | Nparm | DF | Sum of Squares | | |
|--------------|-------|----|----------------|----------|---------|
| | | | F Ratio | Prob > F | |
| height | 2 | 2 | 1544.0000 | 74.7097 | <.0001* |
| width | 1 | 1 | 12.0000 | 1.1613 | 0.3226 |
| height*width | 2 | 2 | 24.0000 | 1.1613 | 0.3747 |

JMP: Fitting the Model

▼ Response > Expanded Estimates

| Expanded Estimates | | | | |
|--|----------|-----------|---------|---------|
| Nominal factors expanded to all levels | | | | |
| Term | Estimate | Std Error | t Ratio | Prob> t |
| Intercept | 51 | 0.927961 | 54.96 | <.0001* |
| height[bottom] | -7 | 1.312335 | -5.33 | 0.0018* |
| height[middle] | 16 | 1.312335 | 12.19 | <.0001* |
| height[top] | -9 | 1.312335 | -6.86 | 0.0005* |
| width[regular] | -1 | 0.927961 | -1.08 | 0.3226 |
| width[wide] | 1 | 0.927961 | 1.08 | 0.3226 |
| height[bottom]*width[regular] | 2 | 1.312335 | 1.52 | 0.1783 |
| height[bottom]*width[wide] | -2 | 1.312335 | -1.52 | 0.1783 |
| height[middle]*width[regular] | -1 | 1.312335 | -0.76 | 0.4749 |
| height[middle]*width[wide] | 1 | 1.312335 | 0.76 | 0.4749 |
| height[top]*width[regular] | -1 | 1.312335 | -0.76 | 0.4749 |
| height[top]*width[wide] | 1 | 1.312335 | 0.76 | 0.4749 |

Recall, "sum to zero"

$$\hat{\mu} =$$

Height

Bottom $\hat{\alpha}_1 = 7$

Middle $\hat{\alpha}_2 = 16$

Top $\hat{\alpha}_3 =$

Width

Regular $\hat{\beta}_1 = -1$

Wide $\hat{\beta}_2 =$

Regular

Bottom $\widehat{\alpha\beta}_{11} = 2$

Middle $\widehat{\alpha\beta}_{21} = -1$

Top $\widehat{\alpha\beta}_{31} =$

Wide

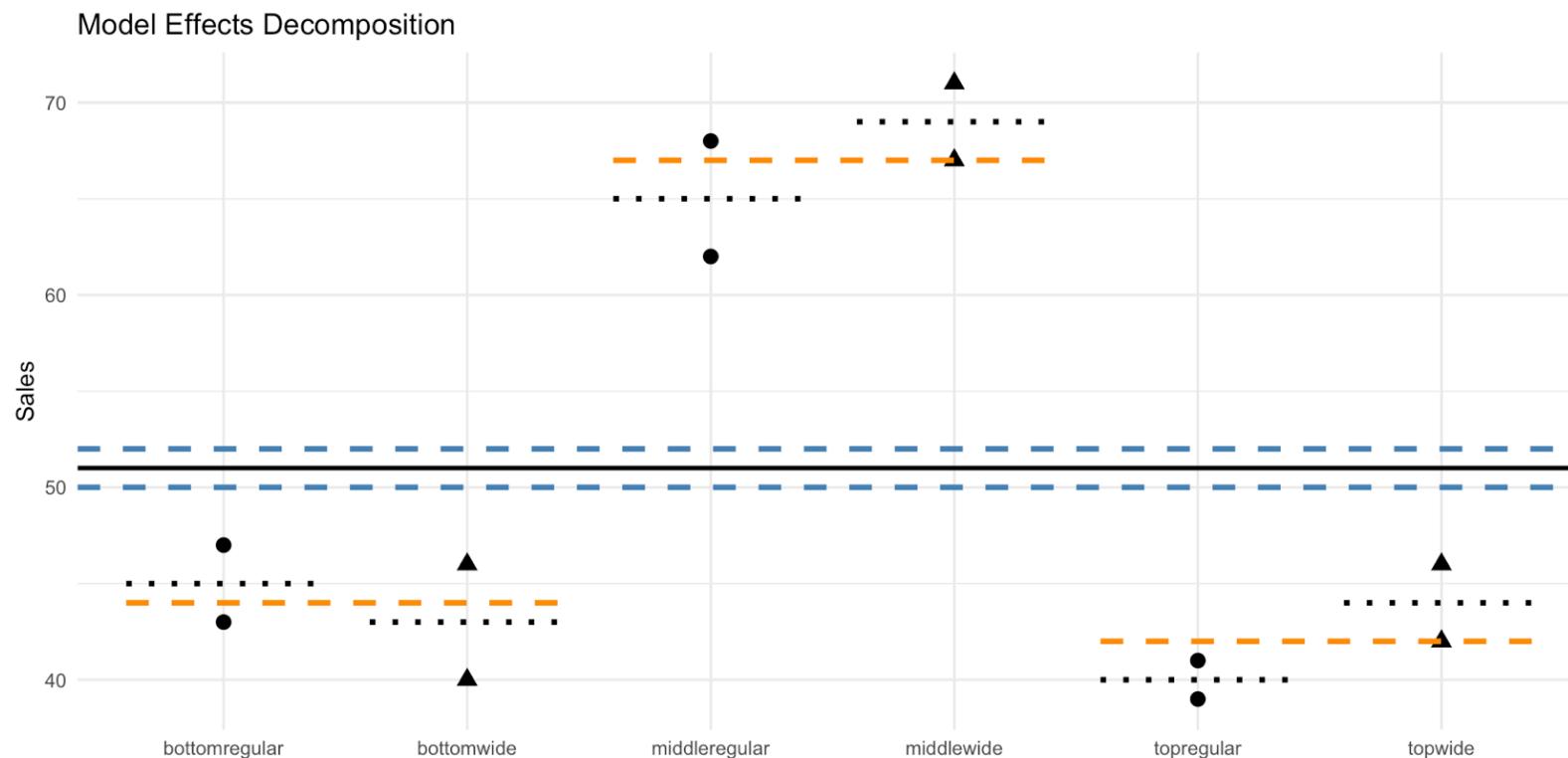
$\widehat{\alpha\beta}_{12} =$

$\widehat{\alpha\beta}_{22} =$

$\widehat{\alpha\beta}_{32} =$

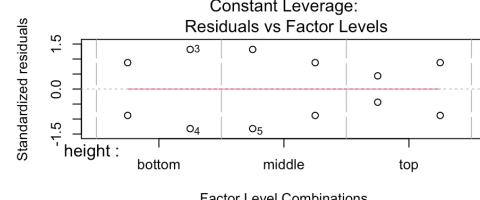
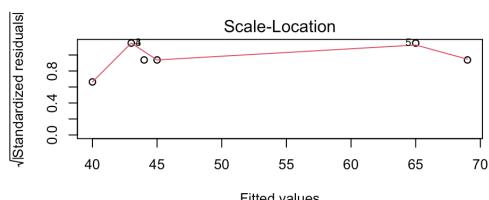
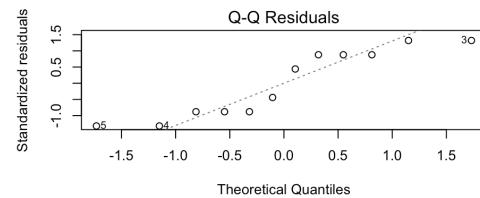
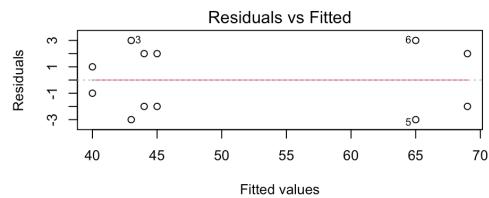
Decomposition of Model Effects

Recall our statistical effects model: $y_{ijk} = \mu + \alpha_i + \beta_j + \alpha\beta_{ij} + \epsilon_{ijk}$



Model Diagnostics

```
1 par(mfrow = c(2,2))
2 plot(bakery_mod)
```



```
1 par(mfrow = c(1,1))
```

