

STAT 516 Lec 06

Two-way factorial design (balanced)

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Tensile strength example from Kuehl (2000)

$$n_{ij} = 3 \text{ for all } i, j.$$

$$y_{111} = 68$$

$$y_{112} = 63$$

Two-way

Table 6.3 Tensile strength (psi) of asphaltic concrete specimens for two aggregate types with each of four compaction methods

$$B \quad b = 4$$

Aggregate Type	Compaction Method			
	1 Static	2 Regular	3 Kneading Low	4 Very Low
Basalt	68	126	93	56
	63	128	101	59
	65	133	98	57
Silicious	71	107	63	40
	66	110	60	41
	66	116	59	44

A

$$a = 2$$

1

2

$$y_{231}$$

Source: A. M. Al-Marshed (1981), Compaction effects on asphaltic concrete durability. M.S. thesis, Civil Engineering, University of Arizona.

$$ab = 8$$

```
y <- c(68, 63, 65, 71, 66, 66, 126, 128, 133, 107, 110, 116,
      93, 101, 98, 63, 60, 59, 56, 59, 57, 40, 41, 44)
agg <- as.factor(rep(c(rep("B", 3), rep("S", 3)), 4))
comp <- as.factor(c(rep("st", 6), rep("r", 6),
                    rep("l", 6), rep("vl", 6)))
```

The two-way factorial experimental design

- ▶ Two factors of interest.
- ▶ Each factor comprehends a set of treatments, called its levels.
- ▶ EUs randomly assigned to treatments.
- ▶ Each treatment is a unique combinations of factor levels.
- ▶ If Factor A has a levels and Factor B has b levels.
- ▶ There are ab treatment groups.

We want to make inferences about:

1. The effects of each factor.
2. The interactions between the two factors.
3. Various differences in treatment group means.

Discuss: Give factors and their levels in the tensile strength experiment.

Main effects and interactions

A main effect is an effect of a factor which does not depend on the level of the other factor.

An interaction is any dependence on the effect of one factor on the level of the other factor.

Two-way treatment effects model

i : level of factor A
 j : level of factor B
 k : EU in the treatment combination

Suppose the responses arise as

$$Y_{ijk} = \mu + \tau_i + \gamma_j + (\tau\gamma)_{ij} + \varepsilon_{ijk}$$

Handwritten annotations:
 - μ : baseline mean
 - τ_i : tau
 - γ_j : gamma
 - ε_{ijk} : error term

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

n_{ij} : # EUs in treatment group ij .

- ▶ Y_{ijk} is the response for EU k under level i of A and level j of B.
- ▶ μ represents a baseline or overall mean
- ▶ The τ_i are the main effects for Factor A.
- ▶ The γ_j are the main effects for Factor B.
- ▶ The $(\tau\gamma)_{ij}$ are the interaction effects between A and B.
- ▶ The ε_{ijk} are $\text{Normal}(0, \sigma^2)$ error terms.

Assume for now

when you have same # EUs in each treatment combination.

1. a balanced design, i.e. $n_{ij} = n$ for all i, j
2. with replication, i.e. $n \geq 2$.

Parameter constraints in the treatment effects model

One-way: $Y_{ij} = \mu + \tau_i + \varepsilon_{ij}$, $i=1, \dots, a$, $j=1, \dots, n$
 We had $\mu, \tau_1, \dots, \tau_a$, so $a+1$ parameters describing a treatment means.
 So we set $\tau_1 = 0$. Just need to estimate $\mu, \tau_2, \dots, \tau_a$.

Treatment effects model has $a + b + ab + 1$ parameters for ab means...

To identify the parameters uniquely, we usually set

$$\tau_1 = 0, \quad \gamma_1 = 0, \quad \text{and} \quad (\tau\gamma)_{1j} = (\tau\gamma)_{i1} = 0 \text{ for all } i, j.$$

Then μ is the mean of the baseline treatment at level 1 of each factor.

$$Y_{ijk} = \mu + \tau_i + \delta_j + (\tau\delta)_{ij} + \varepsilon_{ijk}$$

\uparrow \uparrow \uparrow \uparrow \uparrow
 1 a b ab

$i=1, \dots, a$
 $j=1, \dots, b$
 $k=1, \dots, n$

A total of $1 + a + b + ab$ parameters.

$$Y_{ijk} = \underbrace{\mu + \tau_i + \delta_j + (\tau\delta)_{ij}}_{\mu_{ij}} + \varepsilon_{ijk}$$

$$= \mu_{ij} + \varepsilon_{ijk}$$

Exercise: Let $a = 2$ and $b = 3$. Can tabulate the response means as:

	1	2	3
1	$\mu + \tau_1 + \gamma_1 + (\tau\gamma)_{11}$	$\mu + \tau_1 + \gamma_2 + (\tau\gamma)_{12}$	$\mu + \tau_1 + \gamma_3 + (\tau\gamma)_{13}$
2	$\mu + \tau_2 + \gamma_1 + (\tau\gamma)_{21}$	$\mu + \tau_2 + \gamma_2 + (\tau\gamma)_{22}$	$\mu + \tau_2 + \gamma_3 + (\tau\gamma)_{23}$

$$\tau_i = 0, \delta_i = 0$$

Rewrite the table under the μ -as-baseline constraints

	1	2	3
1	μ	$\mu + \delta_2$	$\mu + \delta_3$
2	$\mu + \tau_2$	$\mu + \tau_2 + \delta_2 + (\tau\delta)_{22}$	$\mu + \tau_2 + \delta_3 + (\tau\delta)_{23}$

Cell means model representation

Assume

$$Y_{ijk} = \mu_{ij} + \varepsilon_{ijk}$$

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

- ▶ Y_{ijk} is the response for EU k under level i of A and level j of B.
- ▶ μ_{ij} is the response mean under level i of A and level j of B.
- ▶ The ε_{ijk} are $\text{Normal}(0, \sigma^2)$ error terms.

Define the marginal means

$$\bar{\mu}_{i.} = \frac{1}{b} \sum_{j=1}^b \mu_{ij}, \quad i = 1, \dots, a \quad \text{and} \quad \bar{\mu}_{.j} = \frac{1}{a} \sum_{i=1}^a \mu_{ij}, \quad j = 1, \dots, b.$$

and the overall mean $\bar{\mu}_{..} = \frac{1}{ab} \sum_{i=1}^a \sum_{j=1}^b \mu_{ij}$.

Hypotheses of interest in the two-way factorial experiment

1. H_0 : Factor A has no main effect.

$$H_0: \bar{\mu}_{1.} = \dots = \bar{\mu}_{a.}$$

2. H_0 : Factor B has no main effect.

$$H_0: \bar{\mu}_{.1} = \dots = \bar{\mu}_{.b}$$

3. H_0 : There is no interaction between Factor A and Factor B.

$$H_0: \mu_{ij} = \bar{\mu}_{i.} + \bar{\mu}_{.j} - \bar{\mu}_{..} \quad \text{for all } i, j.$$

(Factor A effects are same at all levels of Factor B and vice versa.)

$$\mu_{ij} = \bar{\mu}_{..} + (\bar{\mu}_{i.} - \bar{\mu}_{..}) + (\bar{\mu}_{.j} - \bar{\mu}_{..})$$

overall mean. From level i of Factor A From level j of factor B

Example: Let $a = 2$ and $b = 3$. Can tabulate the response means as:

	1	2	3
1	μ_{11}	μ_{12}	μ_{13}
2	μ_{21}	μ_{22}	μ_{23}

$$\bar{\mu}_{1.} = (\mu_{11} + \mu_{12} + \mu_{13}) / 3$$

$$\bar{\mu}_{2.} = (\mu_{21} + \mu_{22} + \mu_{23}) / 3$$

$$\bar{\mu}_{.1} = (\mu_{11} + \mu_{21}) / 2$$

$$\bar{\mu}_{.2} = (\mu_{12} + \mu_{22}) / 2$$

$$\bar{\mu}_{.3} = (\mu_{13} + \mu_{23}) / 2$$

$$\bar{\mu}_{..} = (\mu_{11} + \mu_{12} + \mu_{13} + \mu_{21} + \mu_{22} + \mu_{23}) / 6$$

Make sense of the hypotheses of no main effects and of no interaction.

$$\bar{\mu}_{.1} = (\mu_{11} + \mu_{21}) / 2$$

Goals in two-way factorial experiments

In the two-way treatment effects model

$$Y_{ijk} = \mu + \tau_i + \gamma_j + (\tau\gamma)_{ij} + \varepsilon_{ijk},$$

where $\varepsilon_{ijk} \stackrel{\text{ind}}{\sim} \text{Normal}(0, \sigma^2)$, we wish to

1. Visualize the data.
2. Estimate the parameters $\mu, \tau_i, \gamma_j, (\tau\gamma)_{ij}$.
3. Estimate the error term variance σ^2 .
4. Check whether the model assumptions are satisfied.
5. Decompose the variation in the Y_{ijk} into its sources.
6. Test whether there is *any* effect of the factors on the response.
7. Test for main effects and interaction effects.
8. Do multiple comparisons.

baseline mean

Effect of level i of Factor A

Effect of level j of Factor B

noise

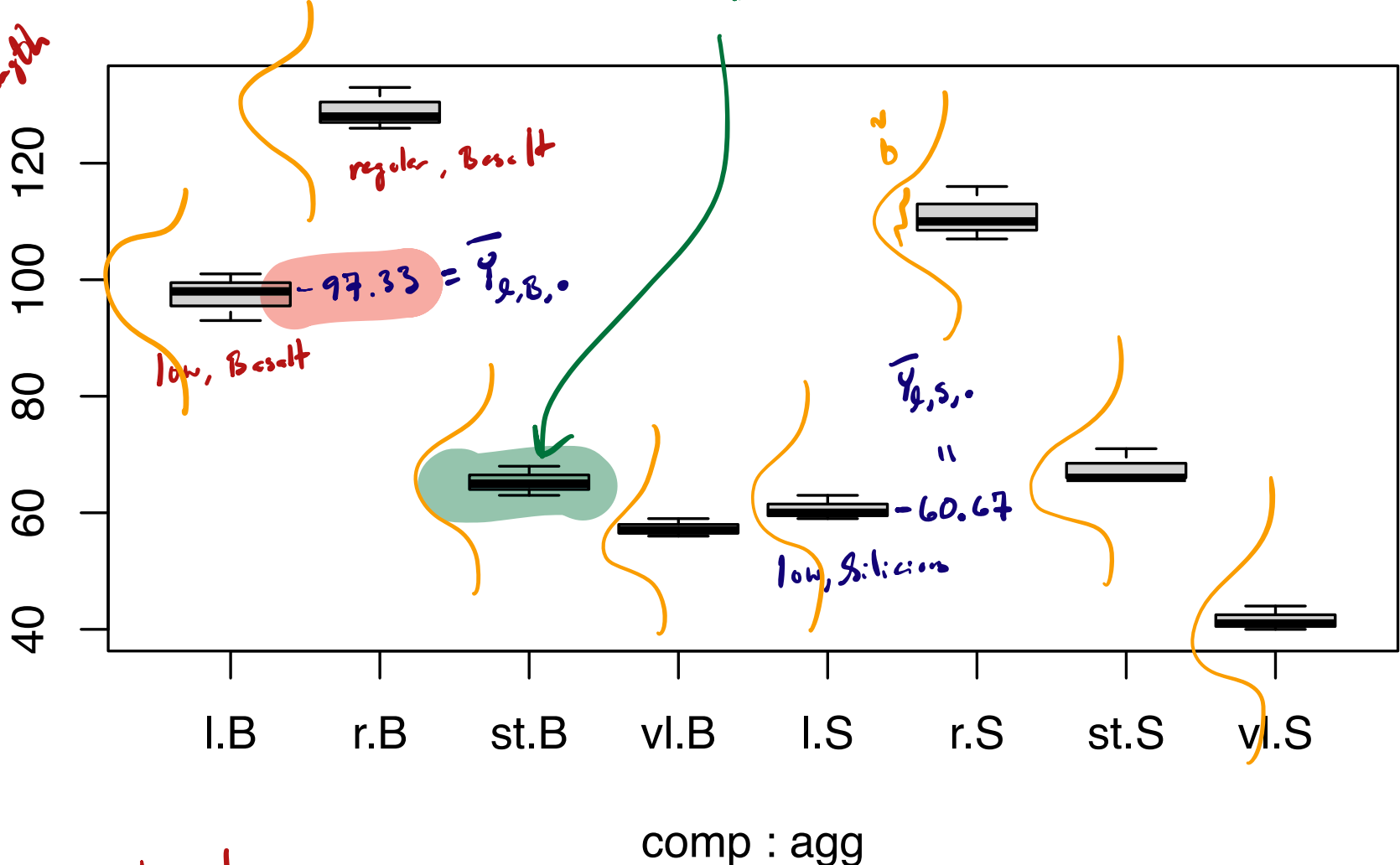
Interaction effect between levels i and j of Factors A and B

Tensile strength data (cont)

```
boxplot(y ~ comp:agg)
```

Suppose static comparison with Basalt is the baseline.

y = tensile strength



Comp: I r st vl
agg: B S

Tensile example (cont)

Use summary() on the lm() function output.

```
lm_out <- lm(y ~ agg + comp + agg:comp)
summary(lm_out)
```

Factor A: $\overset{1}{B}$ $\overset{2}{S}$

Factor B: $\overset{1}{L}$ $\overset{2}{r}$ $\overset{3}{st}$ $\overset{4}{vl}$

```
Call:
lm(formula = y ~ agg + comp + agg:comp)
```

```
Residuals:
    Min       1Q   Median       3Q      Max
-4.3333 -1.6667 -0.6667  2.3333  5.0000
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	97.333	1.780	54.697	< 2e-16 ***
aggS	-36.667	2.517	-14.570	1.18e-10 ***
compr	31.667	2.517	12.583	1.03e-09 ***
compst	-32.000	2.517	-12.716	8.85e-10 ***
compvl	-40.000	2.517	-15.894	3.20e-11 ***
aggS:compr	18.667	3.559	5.245	8.01e-05 ***
aggS:compst	39.000	3.559	10.958	7.58e-09 ***
aggS:compvl	21.000	3.559	5.900	2.24e-05 ***

Sets $\tau_1 = 0$
 $\delta_1 = 0 \dots$

$\hat{\tau}_s$

$\hat{(\tau\delta)}_{23}$

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

```
Residual standard error: 3.082 on 16 degrees of freedom
Multiple R-squared:  0.9921, Adjusted R-squared:  0.9887
F-statistic: 287.6 on 7 and 16 DF, p-value: 1.305e-15
```

Reject H_0 : All treatment means are the same.

$$F_{stat} = \frac{MST_{Treat}}{MSE_{Error}}$$

$$ab - 1 = 2 \cdot 4 - 1 = 7$$

$$ab(n-1) = 2 \cdot 4 \cdot (3-1) = 16$$

$$\sqrt{\hat{\sigma}^2}, \quad \hat{\sigma}^2 = 9.5$$

$$Y_{ijk} = \mu + \tau_i + \tau_j + (\tau\tau)_{ij} + \epsilon_{ijk}$$

$\underbrace{\hspace{10em}}_{\mu_{ij}}$

Exercise: See how the parameter estimates build the treatment means.

```
aggregate(y, by = list(agg, comp), FUN = mean)
```

	Group.1	Group.2	x
1	B	l	97.33333
2	S	l	60.66667
3	B	r	129.00000
4	S	r	111.00000
5	B	st	65.33333
6	S	st	67.66667
7	B	vl	57.33333
8	S	vl	41.66667

$$\hat{\mu} + \hat{\tau}_B + \hat{\tau}_l + (\hat{\tau\tau})_{Bl}$$

$$= 97.33 + (-36.667) = 60.667$$

$$\hat{\mu} + \hat{\tau}_B + \hat{\tau}_r + (\hat{\tau\tau})_{Br}$$

$$= 97.33 + 0 + 31.667 = 129.00 \dots$$

$$\hat{\mu} + \hat{\tau}_S + \hat{\tau}_{vl} + (\hat{\tau\tau})_{Svl}$$

$$= 97.33 + (-36.67) + (-40) + 21.0 = 41.667$$

The fitted values

Define the treatment group means

$$\bar{Y}_{ij.} = \frac{1}{n} \sum_{k=1}^n Y_{ijk} \quad \text{for } i = 1, \dots, a, \quad j = 1, \dots, b.$$

*# EUs of each
Factor level combination*

Then the

- ▶ fitted values are the treatment group means, i.e. $\hat{Y}_{ijk} = \bar{Y}_{ij.}$
- ▶ residuals are the deviations from the group means $\varepsilon_{ijk} = Y_{ijk} - \bar{Y}_{ij.}$

In the cell means model, we estimate μ_{ij} with $\hat{\mu}_{ij} = \bar{Y}_{ij.} \quad \forall ij.$

Estimating the error term variance

An unbiased estimator of the error term variance σ^2 is given by

$$\hat{\sigma}^2 = \frac{1}{ab(n-1)} \sum_{i=1}^a \sum_{j=1}^b \sum_{k=1}^n \overbrace{(Y_{ijk} - \bar{Y}_{ij.})^2}^{\hat{\sigma}_{ijk}^2}, \quad \left(= 9.5 \text{ on concrete data} \right)$$

degrees of freedom

which is the sum of the squared residuals divided by $ab(n-1)$.

We have $a \times b \times n$ total observations

$\underline{abn} - \underline{ab}$ — We had first estimate the $a \times b$ group means in order to get the residuals.

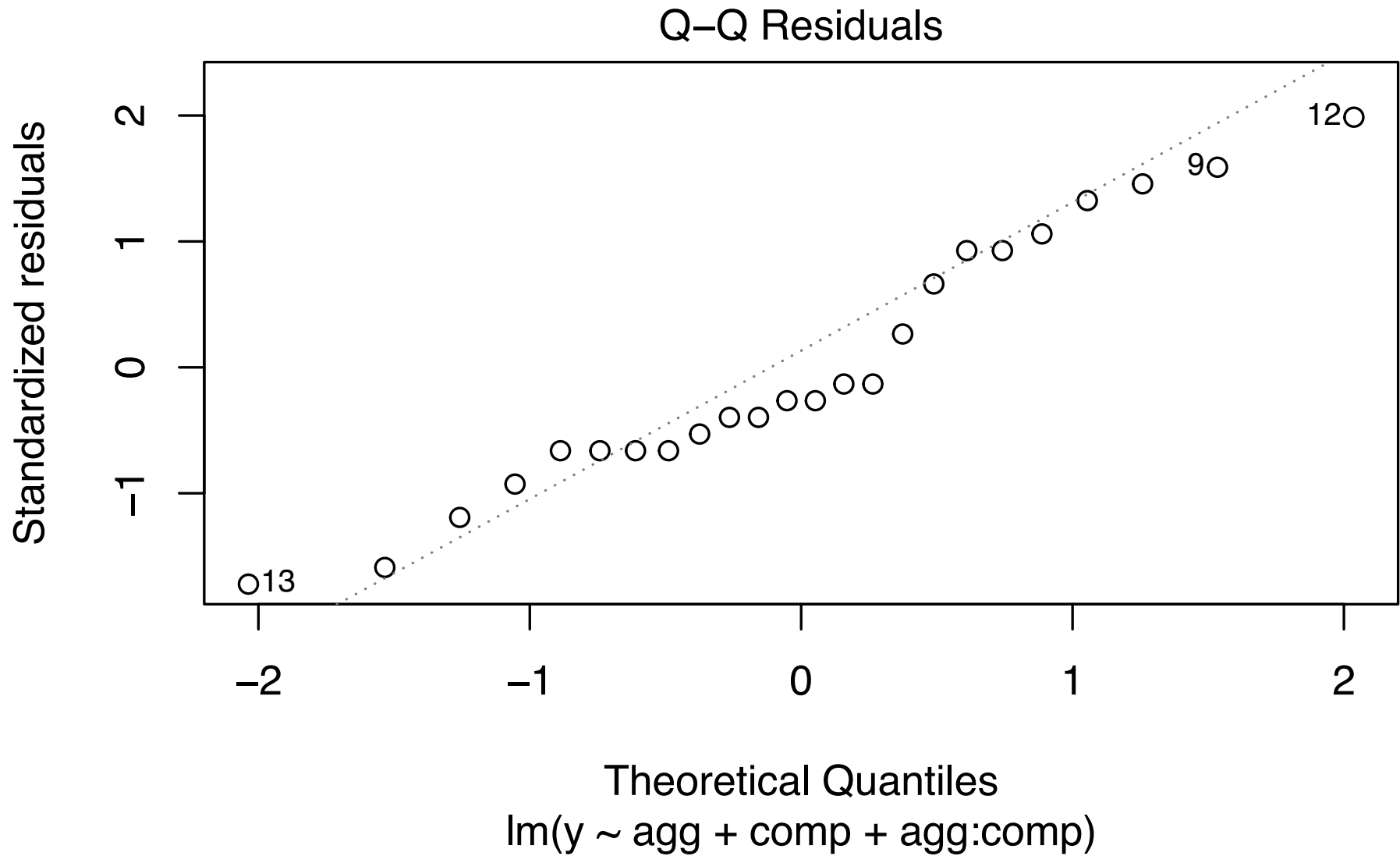
Checking model assumptions

Validity of the methods in these slides depends on these assumptions:

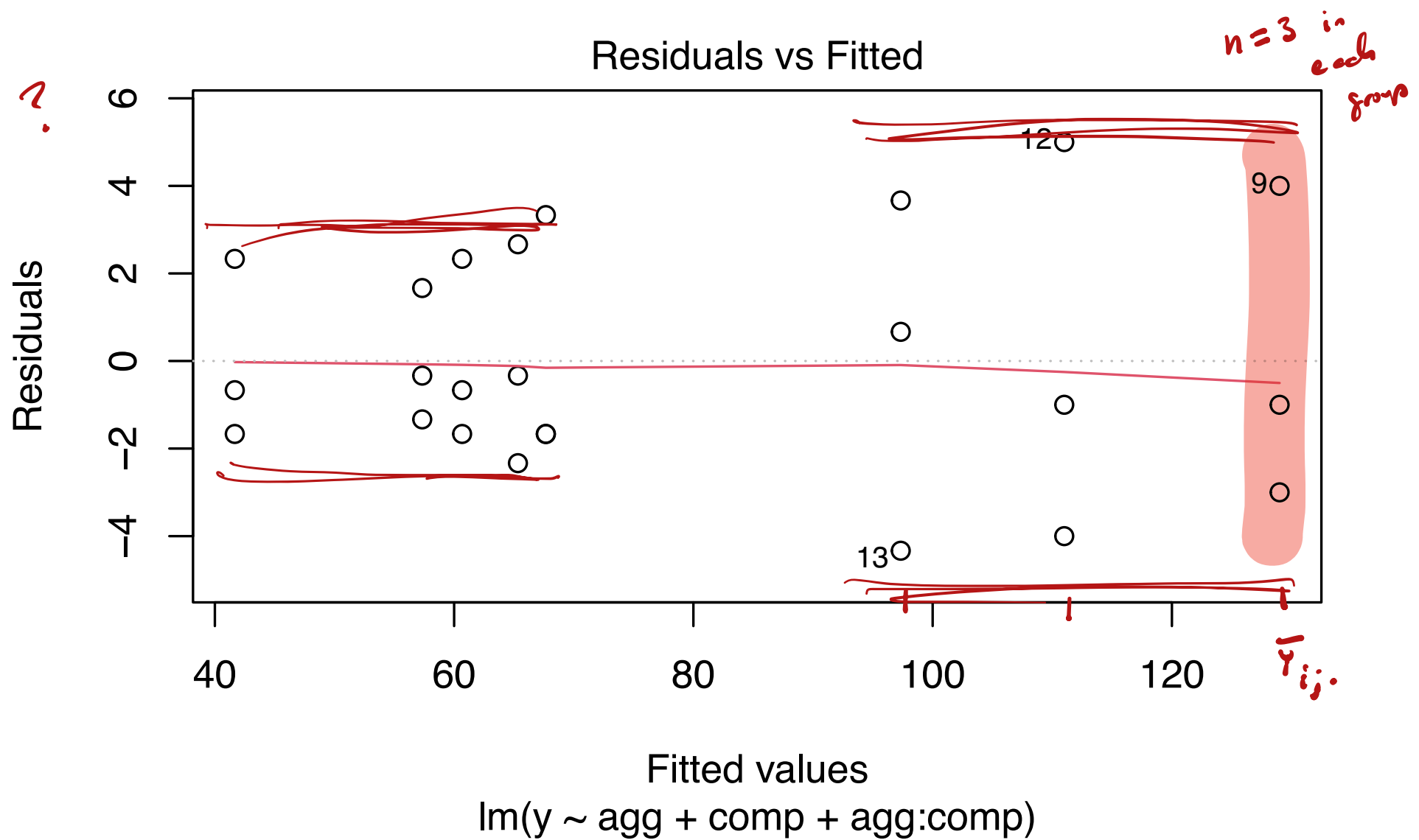
1. The responses are normally distributed around the treatment means (Check QQ plot of residuals).
2. The response has the same variance in all treatment groups (Check residuals vs fitted values plot).
3. The response values are independent of each other (No way to check; must trust experimental design).

Tensile example (cont)

```
plot(lm_out, which = 2)
```



```
plot(lm_out, which = 1)
```



Sums of squares in the two-way factorial experiment

$$\bar{y}_{...} = \frac{1}{abn} \sum_{i=1}^a \sum_{j=1}^b \sum_{k=1}^n y_{ijk}$$

Handwritten: Variability owing to all the different treatment combinations

SS	Symbol	Formula
Total	SS_{Tot}	$\sum_{i=1}^a \sum_{j=1}^b \sum_{k=1}^n (Y_{ijk} - \bar{Y}_{...})^2$
Treatments	SS_{Trt}	$\sum_{i=1}^a \sum_{j=1}^b n(\bar{Y}_{ij.} - \bar{Y}_{...})^2$
Error	SS_{Error}	$\sum_{i=1}^a \sum_{j=1}^b \sum_{k=1}^n (Y_{ijk} - \bar{Y}_{ij.})^2$

Handwritten: \sum_{ijk}^2

Handwritten: noise

$$\hat{\sigma}^2 = \frac{SS_{Error}}{ab(n-1)}$$

- ▶ We have $SS_{Tot} = SS_{Trt} + SS_{Error}$
- ▶ We again define $R^2 = \frac{SS_{Trt}}{SS_{Tot}}$. Always have $0 \leq R^2 \leq 1$
- ▶ Can further decompose SS_{Trt} into SS from main effects and interactions.

ANOVA table for overall F test

Source	Df	SS	MS	F value	p value
Treatments	$ab - 1$	SS_{Trt}	MS_{Trt}	F_{stat}	$P(F > F_{\text{stat}})$
Error	$ab(n - 1)$	SS_{Error}	MS_{Error}		
Total	$abn - 1$	SS_{Tot}			

$abn - ab$
 ↑ first have to compute \bar{Y}_{ij} for all i, j , so ab quantities

total # obs

must first compute $\bar{Y} \dots$

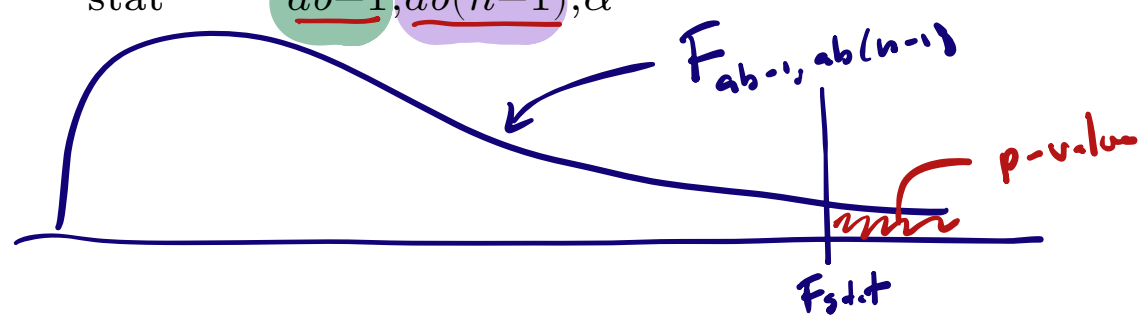
$$Df_{\text{Trt}} + Df_{\text{Error}} = Df_{\text{Tot}}$$

$$ab - 1 + ab(n - 1) = abn - 1$$

► The F statistic $F_{\text{stat}} = \frac{MS_{\text{Trt}}}{MS_{\text{Error}}}$ tests

H_0 : All the μ_{ij} are the same. ("None of the treatment combinations has any effect")
 H_1 : The μ_{ij} are not all the same.

► Reject H_0 if $F_{\text{stat}} > F_{ab-1, ab(n-1), \alpha}$



Exercise: Make ANOVA table for the tensile strength data. Interpret!

```
a <- 2
b <- 4
n <- 3
yhat <- predict(lm_out)
ehat <- y - yhat
SSE <- sum(ehat^2)
MSE <- SSE / (a*b*(n-1))
SSR <- sum((yhat - mean(y))^2)
MSR <- SSR / (a*b - 1)
SST <- sum((y - mean(y))^2)
Fstat <- MSR / MSE
pval <- 1 - pf(Fstat, a*b-1, a*b*(n-1))
```

Source	Df	SS	MS	F value	p value
Treatments	7	19122.50	2731.79	287.5564	0.0000
Error	16	152.00	9.50		
Total	23	19274.50			

Further decomposition of treatments sum of squares

$$SS_{\text{TPT}} = \sum_{i=1}^a \sum_{j=1}^b n (\bar{Y}_{ij.} - \bar{Y}_{...})^2$$

Define main effect and interaction sums of squares SS_A , SS_B , and SS_{AB} :

▶ $SS_A = bn \sum_{i=1}^a (\bar{Y}_{i..} - \bar{Y}_{...})^2$

$$\bar{Y}_{i..} = \frac{1}{bn} \sum_{j=1}^b \sum_{k=1}^n Y_{ijk}$$

▶ $SS_B = an \sum_{j=1}^b (\bar{Y}_{.j.} - \bar{Y}_{...})^2$

▶ $SS_{AB} = n \sum_{i=1}^a \sum_{j=1}^b (\bar{Y}_{ij.} - (\bar{Y}_{i..} + \bar{Y}_{.j.} - \bar{Y}_{...}))^2$
 $((\bar{Y}_{ij.} - \bar{Y}_{..}) - (\bar{Y}_{i..} - \bar{Y}_{...}) - (\bar{Y}_{.j.} - \bar{Y}_{...}))^2$

In the balanced design we have

$$SS_{\text{Trt}} = SS_A + SS_B + SS_{AB}.$$

$$SS_{AB} = SS_{\text{TPT}} - (SS_A + SS_B)$$

Aggregate Type	Compaction Method			
	1 Static	2 Regular	3 Kneading Low	4 Very Low
1 Basalt	68	126	93	56
	63	128	101	59
	65	133	98	57
2 Silicious	71	107	63	40
	66	110	60	41
	66	116	59	44

Source: A. M. Al-Marshed (1981), Compaction effects on asphaltic concrete durability. M.S. thesis, Civil Engineering, University of Arizona.

$\bar{y}_{.1.}$

"

66.5

$\bar{y}_{.2.}$

"

120.0

$\bar{y}_{.3.}$

"

79.0

$\bar{y}_{.4.}$

"

49.5

$\bar{y}_{1..} = 87.25$

$\bar{y}_{2..} = 70.25$

a b

$\bar{y}_{...} = 78.75$

↑
mean of all observations

$$SS_A = 4 \times 3 \left[(87.25 - 78.75)^2 + (70.25 - 78.75)^2 \right]$$

$$SS_B = 2 \times 3 \left[(66.5 - 78.75)^2 + \dots \right]$$

$$SS_{AB} = SS_{TOT} - (SS_A + SS_B)$$

Full sums of squares in two-way factorial experiment

$$SS_{TFT} = SS_A + SS_B + SS_{AB}$$

SS	Symbol	Formula
Total	SS_{Tot}	$\sum_{i=1}^a \sum_{j=1}^b \sum_{k=1}^n (Y_{ijk} - \bar{Y}_{...})^2$
A	SS_A	$bn \sum_{i=1}^a (\bar{Y}_{i..} - \bar{Y}_{...})^2$
B	SS_B	$an \sum_{j=1}^b (\bar{Y}_{.j.} - \bar{Y}_{...})^2$
AB	SS_{AB}	$n \sum_{i=1}^a \sum_{j=1}^b (Y_{ij.} - (\bar{Y}_{i..} + \bar{Y}_{.j.} - \bar{Y}_{...}))^2$
Error	SS_{Error}	$\sum_{i=1}^a \sum_{j=1}^b \sum_{k=1}^n (Y_{ijk} - \bar{Y}_{ij.})^2$

► We have $SS_{Tot} = \underbrace{SS_A + SS_B + SS_{AB}}_{SS_{TFT}} + SS_{Error}$.

Full ANOVA table for balanced two-way factorial design

$$Df_{TOT} = ab - 1 = Df_A + Df_B + Df_{AB}$$

$$= a - 1 + b - 1 + (a - 1)(b - 1)$$

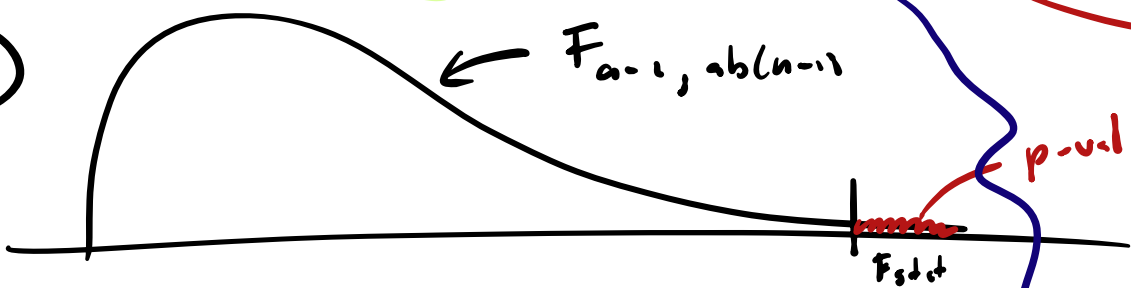
SS_{TOT}

Source	Df	SS	MS	F value
A	$a - 1$	SS_A	MS_A	$F_A = MS_A / MS_{Error}$
B	$b - 1$	SS_B	MS_B	$F_B = MS_B / MS_{Error}$
AB	$(a - 1)(b - 1)$	SS_{AB}	MS_{AB}	$F_{AB} = MS_{AB} / MS_{Error}$
Error	$ab(n - 1)$	SS_{Error}	MS_{Error}	
Total	$abn - 1$	SS_{Tot}		

$\bar{\mu}_{1.} = \dots = \bar{\mu}_{a.}$ (Ignoring Factor B, Factor A has no effect)

1. Reject H_0 : no Factor A main effect if $F_A > F_{a-1, ab(n-1), \alpha}$.
2. Reject H_0 : no Factor B main effect if $F_B > F_{b-1, ab(n-1), \alpha}$.
3. Reject H_0 : no A and B interaction if $F_{AB} > F_{(a-1)(b-1), ab(n-1), \alpha}$.

①



(Ignoring Factor A, Factor B has no effect)

Tensile strength data (cont)

"Effects of one factor are same no matter the level of the other factor."

Obtain ANOVA table with `anova()` function on `lm()` output.

```
anova(lm(y ~ agg + comp + agg:comp))
```

Analysis of Variance Table

Response: y

Source	Df	Sum Sq	Mean Sq	F value	Pr(>F)
A agg	1	1734	1734.0	182.526	3.628e-10 ***
B comp	3	16244	5414.5	569.947	< 2.2e-16 ***
AB agg:comp	3	1145	381.7	40.175	1.124e-07 ***
Error Residuals	16	152	9.5		

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

$a-1 = 2-1 = 1$

$b-1 = 4-1 = 3$

small

$(a-1)(b-1) = (2-1)(4-1) = 3$

$ab(n-1) = 2 \cdot 4 (3-1) = 16$

Reject null hypothesis of no interaction. I should expect to see evidence of interaction in interaction plots.

Important: `anova()` function only appropriate for a balanced design.

Interaction is significant. Now what?

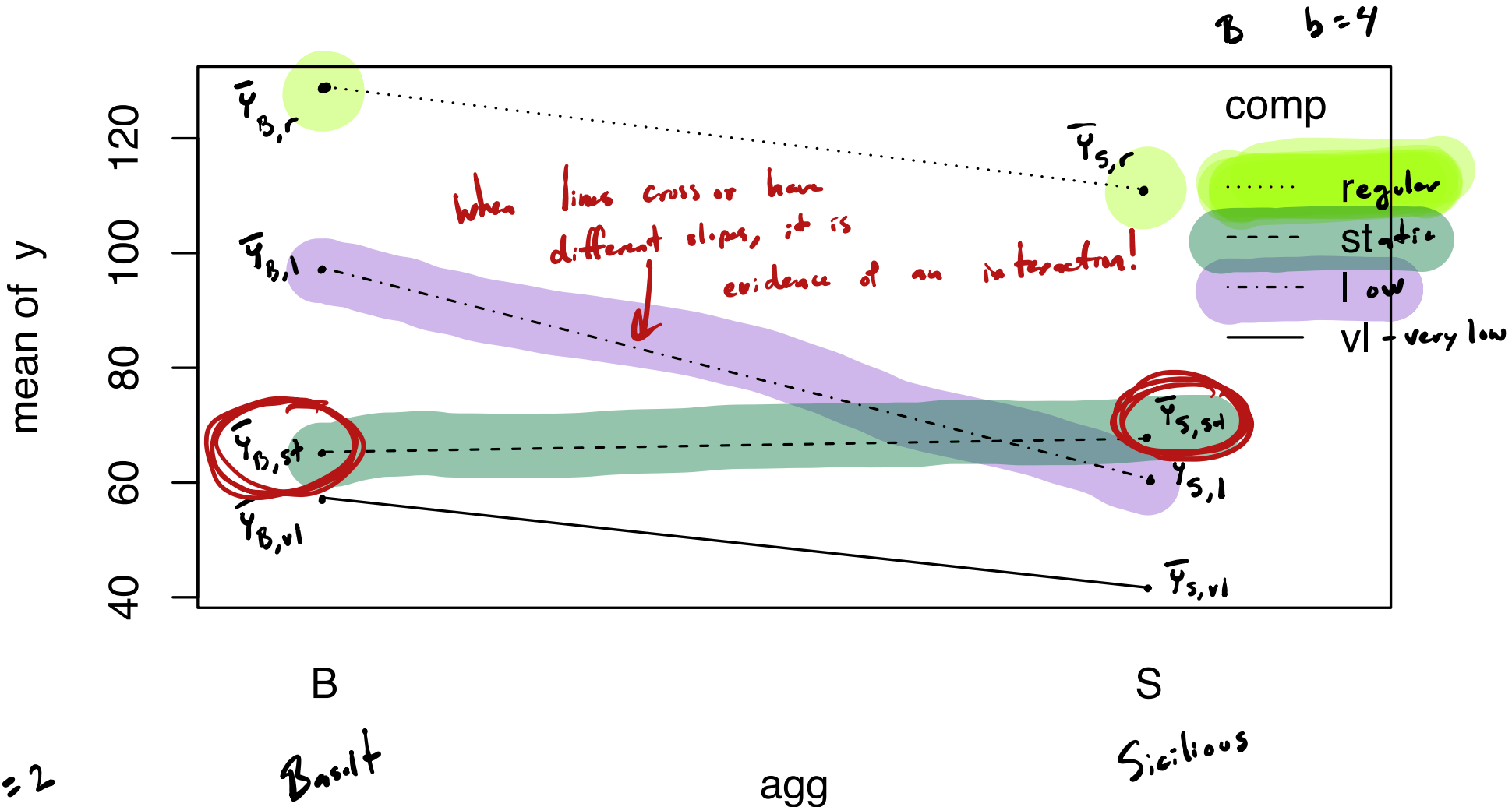
When you find a significant interaction:

1. Make interaction plots (next slides).
2. Be very cautious about interpreting main effects, even when these are statistically significant.

Tensile strength data (cont)

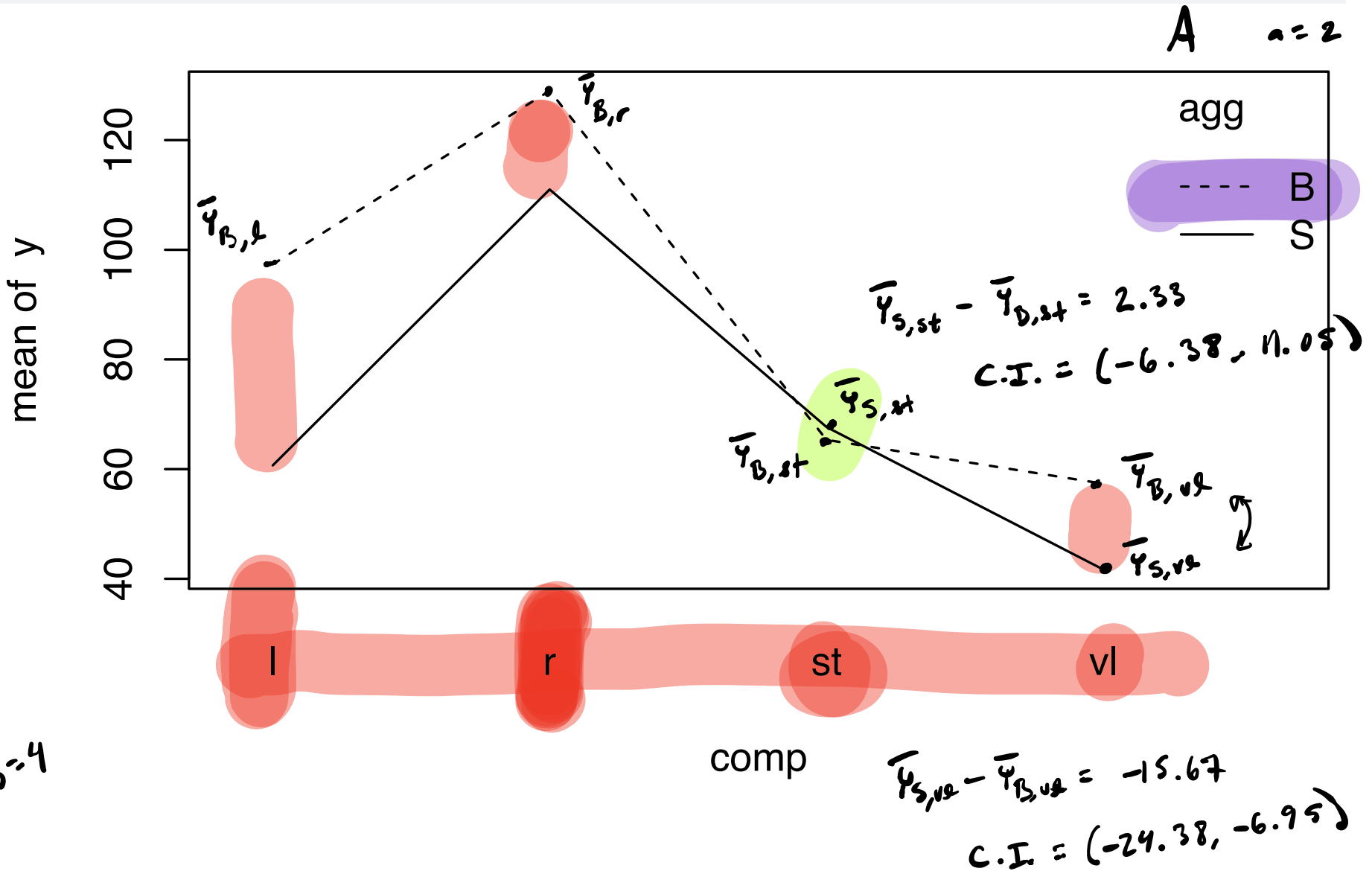
Use the `interaction.plot()` function to visualize an interaction:

```
interaction.plot(agg, comp, y)
```



Interactions appear as crossing lines or differing slopes.

```
interaction.plot(comp, agg, y)
```



Estimates of cell and marginal means in the balanced case

The estimators of the cell and marginal means are given by

- ▶ $\hat{\mu}_{ij} = \bar{Y}_{ij.}, i = 1, \dots, a, j = 1, \dots, b$
- ▶ $\hat{\mu}_{i.} = \bar{Y}_{i..}, i = 1, \dots, a.$
- ▶ $\hat{\mu}_{.j} = \bar{Y}_{.j.}, j = 1, \dots, b.$

We estimate $\hat{\mu}_{i.}$ with $\bar{Y}_{i..}$ (and $\hat{\mu}_{.j}$ with $\bar{Y}_{.j.}$) only when $n_{ij} = n \forall ij.$

Tensile data

	1	2	3	4	
1	$\bar{Y}_{11.}$	$\bar{Y}_{12.}$	$\bar{Y}_{13.}$	$\bar{Y}_{14.}$	$\bar{Y}_{1..}$
2	$\bar{Y}_{21.}$	$\bar{Y}_{22.}$	$\bar{Y}_{23.}$	$\bar{Y}_{24.}$	$\bar{Y}_{2..}$
	$\bar{Y}_{.1.}$	$\bar{Y}_{.2.}$	$\bar{Y}_{.3.}$	$\bar{Y}_{.4.}$	$\bar{Y}_{...}$

Factor A marginal means

Factor B marginal means

$$\hat{\mu}_{11} = \bar{y}_{11}$$

μ_{11}	μ_{12}	μ_{13}	μ_{14}	$\bar{\mu}_{1.}$
μ_{21}	μ_{22}	μ_{23}	μ_{24}	$\bar{\mu}_{2.}$
$\bar{\mu}_{.1}$	$\bar{\mu}_{.2}$	$\bar{\mu}_{.3}$	$\bar{\mu}_{.4}$	$\bar{\mu}_{..}$

Some CI formulas (without familywise adjustment)

These CI formulas are for the balanced design $n_{ij} = n \forall ij$.

Target (1 - α)100% confidence interval

μ_{ij}	$\bar{Y}_{ij.} \pm t_{ab(n-1), \alpha/2} \hat{\sigma} \sqrt{\frac{1}{n}}$
$\mu_{ij} - \mu_{i'j'}$	$\bar{Y}_{ij.} - \bar{Y}_{i'j'}. \pm t_{ab(n-1), \alpha/2} \hat{\sigma} \sqrt{\frac{2}{n}}$
$\bar{\mu}_{i.}$	$\bar{Y}_{i..} \pm t_{ab(n-1), \alpha/2} \hat{\sigma} \sqrt{\frac{1}{bn}}$
$\bar{\mu}_{.j}$	$\bar{Y}_{.j.} \pm t_{ab(n-1), \alpha/2} \hat{\sigma} \sqrt{\frac{1}{an}}$
$\bar{\mu}_{i.} - \bar{\mu}_{i'..}$	$\bar{Y}_{i..} - \bar{Y}_{i'..} \pm t_{ab(n-1), \alpha/2} \hat{\sigma} \sqrt{\frac{2}{bn}}$
$\bar{\mu}_{.j} - \bar{\mu}_{.j'}$	$\bar{Y}_{.j.} - \bar{Y}_{.j'}. \pm t_{ab(n-1), \alpha/2} \hat{\sigma} \sqrt{\frac{2}{an}}$

In the above $\hat{\sigma} = \sqrt{\text{MS}_{\text{Error}}}$.

Comparing means at all factor level combinations

Tensile

$a=2$ $b=4$

μ_{11}	μ_{12}	μ_{13}	μ_{14}
μ_{21}	μ_{22}	μ_{23}	μ_{24}

$ab = 2(4) = 8$ total means
 $\binom{8}{2} = \frac{8 \cdot 7}{2} = 28$ comparisons

- Tukey's for comparing all pairs among μ_{ij} , $i = 1, \dots, a$, $j = 1, \dots, b$:

$$\bar{Y}_{ij.} - \bar{Y}_{i'j'}. \pm q_{ab, ab(n-1), \alpha} \hat{\sigma} \frac{1}{\sqrt{n}}, \quad (i, j) \neq (i', j').$$

Handwritten notes:
 q_{ab} is circled in red, with a red box around $ab(n-1)$.
 $\hat{\sigma}$ is circled in red.
 $\sqrt{MS_{Error}}$ is written above the fraction, with an arrow pointing to $\hat{\sigma}$.
 Df_{Error} is written above the box, with an arrow pointing to $ab(n-1)$.
 $\# TPT$ combinations is written below the box.

- Dunnnett's for comparing all means μ_{ij} to a baseline μ_{ab} :

$$\bar{Y}_{ij.} - \bar{Y}_{ab.} \pm d_{ab, ab(n-1), \alpha} \hat{\sigma} \sqrt{\frac{2}{n}}, \quad (i, j) \neq (a, b).$$

Handwritten notes:
 $\bar{Y}_{ab.}$ is circled in green and labeled "Baseline".
 d_{ab} is circled in green, with a green box around $ab(n-1)$.
 $\sqrt{\frac{2}{n}}$ has a green arrow pointing to the 2.
 $\# TPT$ combinations is written below the box.
 Df_{Error} is written below the box.

Use $\hat{\sigma} = \sqrt{MS_{Error}}$.

These may make more comparisons than are of interest...

Tensile strength data (cont)

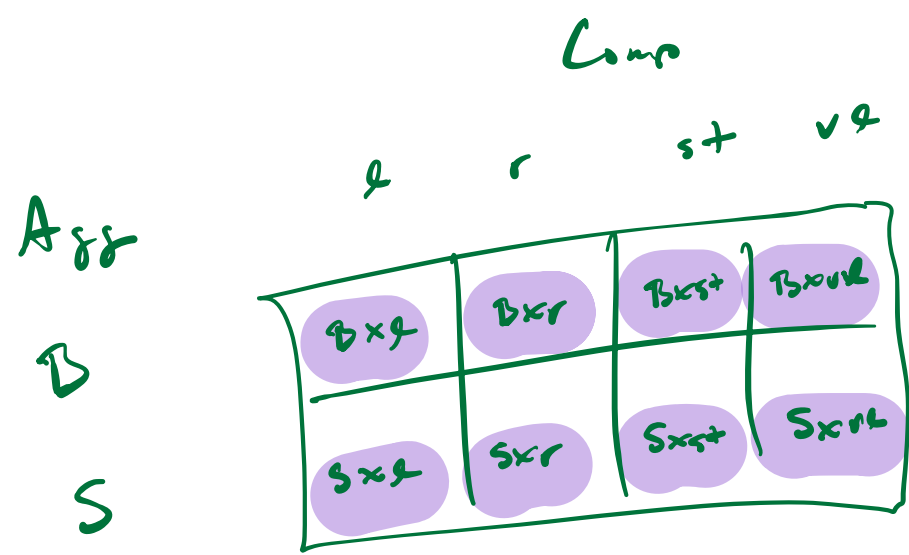
```
TukeyHSD(aov(lm(y ~ agg:comp)))
```

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

```
Fit: aov(formula = lm(y ~ agg:comp))
```

```
$`agg:comp`
```

	diff	lwr	upr	p adj
S:l-B:l	-36.666667	-45.379555	-27.9537785	0.0000000
B:r-B:l	31.666667	22.953778	40.3795548	0.0000000
S:r-B:l	13.666667	4.953778	22.3795548	0.0011210
B:st-B:l	-32.000000	-40.712888	-23.2871118	0.0000000
S:st-B:l	-29.666667	-38.379555	-20.9537785	0.0000001
B:vl-B:l	-40.000000	-48.712888	-31.2871118	0.0000000
S:vl-B:l	-55.666667	-64.379555	-46.9537785	0.0000000
B:r-S:l	68.333333	59.620445	77.0462215	0.0000000
S:r-S:l	50.333333	41.620445	59.0462215	0.0000000
B:st-S:l	4.666667	-4.046222	13.3795548	0.5964603
S:st-S:l	7.000000	-1.712888	15.7128882	0.1678762
B:vl-S:l	-3.333333	-12.046222	5.3795548	0.8765993
S:vl-S:l	-19.000000	-27.712888	-10.2871118	0.0000257
S:r-B:r	-18.000000	-26.712888	-9.2871118	0.0000501
B:st-B:r	-63.666667	-72.379555	-54.9537785	0.0000000
S:st-B:r	-61.333333	-70.046222	-52.6204452	0.0000000
B:vl-B:r	-71.666667	-80.379555	-62.9537785	0.0000000
S:vl-B:r	-87.333333	-96.046222	-78.6204452	0.0000000
B:st-S:r	-45.666667	-54.379555	-36.9537785	0.0000000
S:st-S:r	-43.333333	-52.046222	-34.6204452	0.0000000
B:vl-S:r	-53.666667	-62.379555	-44.9537785	0.0000000
S:vl-S:r	-69.333333	-78.046222	-60.6204452	0.0000000
S:st-B:st	2.333333	-6.379555	11.0462215	0.9785200
B:vl-B:st	-8.000000	-16.712888	0.7128882	0.0842128
S:vl-B:st	-23.666667	-32.379555	-14.9537785	0.0000015
B:vl-S:st	-10.333333	-19.046222	-1.6204452	0.0145554
S:vl-S:st	-26.000000	-34.712888	-17.2871118	0.0000004
S:vl-B:vl	-15.666667	-24.379555	-6.9537785	0.0002561



To use Dunnett's

Convert to
One-way ANOVA

↑ P-val
Compare to
Dunnett's output!!

Pretend we have 8 treatments
of one single factor

Easiest way to do Dunnett's is to convert the design to a one-way:

```
agg_comp <- as.factor(paste(agg, comp, sep="_"))
levels(agg_comp)
```

```
[1] "B_l" "B_r" "B_st" "B_vl" "S_l" "S_r" "S_st" "S_vl"
```

```
library(DescTools)
DunnettTest(y ~ agg_comp, control = "B_st", conf.level = 0.95)
```

Dunnett's test for comparing several treatments with a control :
95% family-wise confidence level

Basalt and static

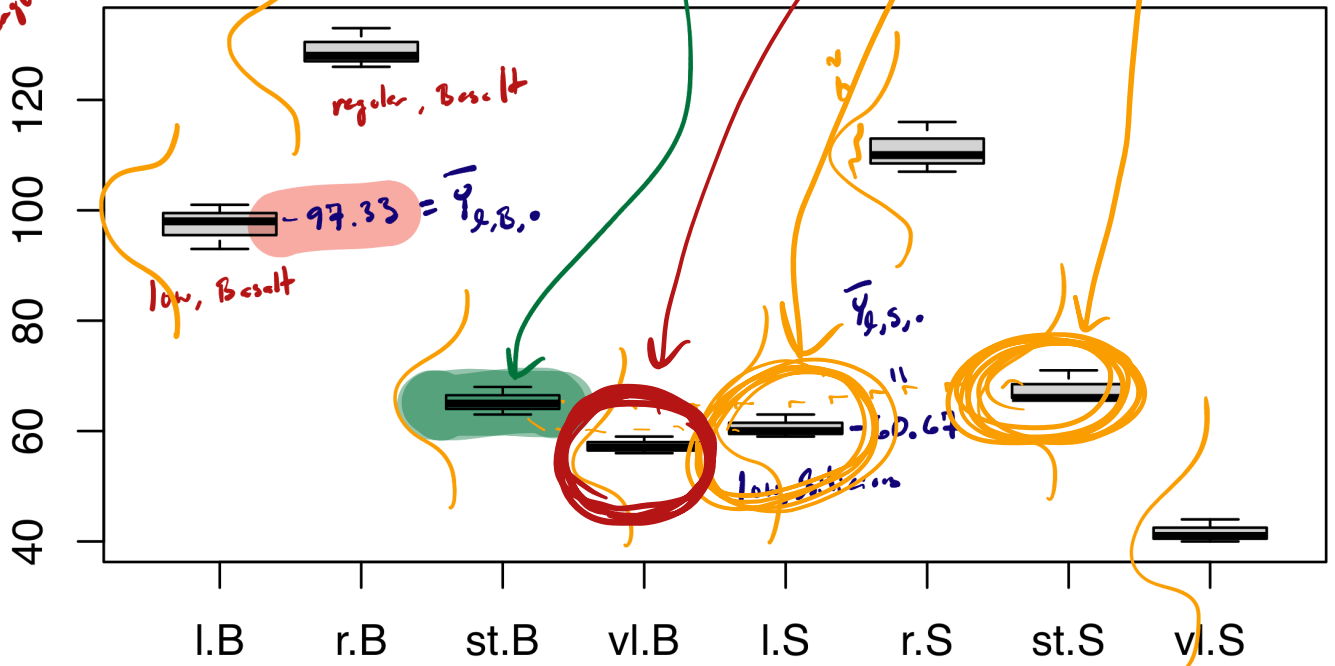
\$B_st	diff	lwr.ci	upr.ci	pval	
B_l-B_st	32.000000	24.643829	39.3561715	3.8e-10	***
B_r-B_st	63.666667	56.310495	71.0228381	< 2e-16	***
B_vl-B_st	-8.000000	-15.356171	-0.6438285	0.0304	*
S_l-B_st	-4.666667	-12.022838	2.6895048	0.3264	
S_r-B_st	45.666667	38.310495	53.0228381	2.8e-15	***
S_st-B_st	2.333333	-5.022838	9.6895048	0.8882	
S_vl-B_st	-23.666667	-31.022838	-16.3104952	1.1e-07	***

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

```
boxplot(y ~ comp:agg)
```

Suppose static comparison with Basalt is the baseline.

$y = \text{tensile strength}$



Comparing factor level means at fixed level of other factor

E.g. With Besolt aggregate, compare all pairs of comparison methods.

Fix Factor A at level i and compare means across factor B:

- ▶ Tukey's for comparing all pairs among $\bar{\mu}_{i1}, \dots, \bar{\mu}_{ib}$:

$$\bar{Y}_{ij.} - \bar{Y}_{ij'.} \pm q_{b, ab(n-1), \alpha} \hat{\sigma} \frac{1}{\sqrt{n}}, \quad 1 \leq j < j' \leq b.$$

Tukey table (pointing to $q_{b, ab(n-1), \alpha}$) *# levels of factor B* (pointing to b)

Same level of factor A, different levels of factor B. (pointing to $\bar{Y}_{ij.} - \bar{Y}_{ij'.}$) *Df Error* (pointing to $ab(n-1)$)

- ▶ Dunnett's for comparing $\bar{\mu}_{ij}, \dots, \bar{\mu}_{i,b-1}$ to baseline μ_{ib} :

$$\bar{Y}_{ij.} - \bar{Y}_{ib.} \pm d_{b, ab(n-1), \alpha} \hat{\sigma} \sqrt{\frac{2}{n}}, \quad j = 1, \dots, b - 1.$$

To do this at *all* levels $i = 1, \dots, a$, divide α by a (à la Bonferroni)!

Likewise if comparing means of factor B at fixed levels of factor A.

Tensile strength data (cont)

For Basalt and then for Silicious

For each aggregate type, compare all pairs of compaction method means.

Ensure that the familywise coverage probability is at least 0.95.

Use $q_{4,16,0.05/2} = \text{qtukey}(1-0.05/2, 4, 16) = 4.54763$.

Ex. $\bar{y}_{B,t} - \bar{y}_{B,st} \pm t_{b, ab(n-1), \alpha} \frac{\hat{\sigma}}{a} \sqrt{\frac{1}{n}}$

$a = 2$
 $b = 4$
 $n = 3$

```
y11. <- mean(y[agg == "B" & comp == "l"])
y12. <- mean(y[agg == "B" & comp == "r"])
y13. <- mean(y[agg == "B" & comp == "st"])
y14. <- mean(y[agg == "B" & comp == "v1"])
```

```
y21. <- mean(y[agg == "S" & comp == "l"])
y22. <- mean(y[agg == "S" & comp == "r"])
y23. <- mean(y[agg == "S" & comp == "st"])
y24. <- mean(y[agg == "S" & comp == "v1"])
```

```
alpha <- 0.05
```

```
me <- qtkey(1-alpha/a, b, a*b*(n-1)) * sqrt(MSE) / sqrt(n)
```

$t_{4, 16, 0.05}$
 3.082

$2.4(3-1) = 16$


```

ttab <- rbind(c(y11. - y12. - me, y11. - y12. + me),
             c(y11. - y13. - me, y11. - y13. + me),
             c(y11. - y14. - me, y11. - y14. + me),
             c(y12. - y13. - me, y12. - y13. + me),
             c(y12. - y14. - me, y12. - y14. + me),
             c(y13. - y14. - me, y13. - y14. + me),
             c(y21. - y22. - me, y21. - y22. + me),
             c(y21. - y23. - me, y21. - y23. + me),
             c(y21. - y24. - me, y21. - y24. + me),
             c(y22. - y23. - me, y22. - y23. + me),
             c(y22. - y24. - me, y22. - y24. + me),
             c(y23. - y24. - me, y23. - y24. + me))
rownames(ttab) <- c("B:l-r", "B:l-st", "B:l-vl", "B:r-st", "B:r-vl", "B:st-vl",
                  "S:l-r", "S:l-st", "S:l-vl", "S:r-st", "S:r-vl", "S:st-vl")
colnames(ttab) <- c("lower", "upper")

```

Tukey/Bonferroni-adjusted confidence intervals comparing all pairs of compaction methods for each aggregate type:

ttab

$$b = 4 \quad \binom{4}{2} = \frac{4 \cdot 3}{2} = 6$$

Compare all pairs of compaction methods under each aggregate

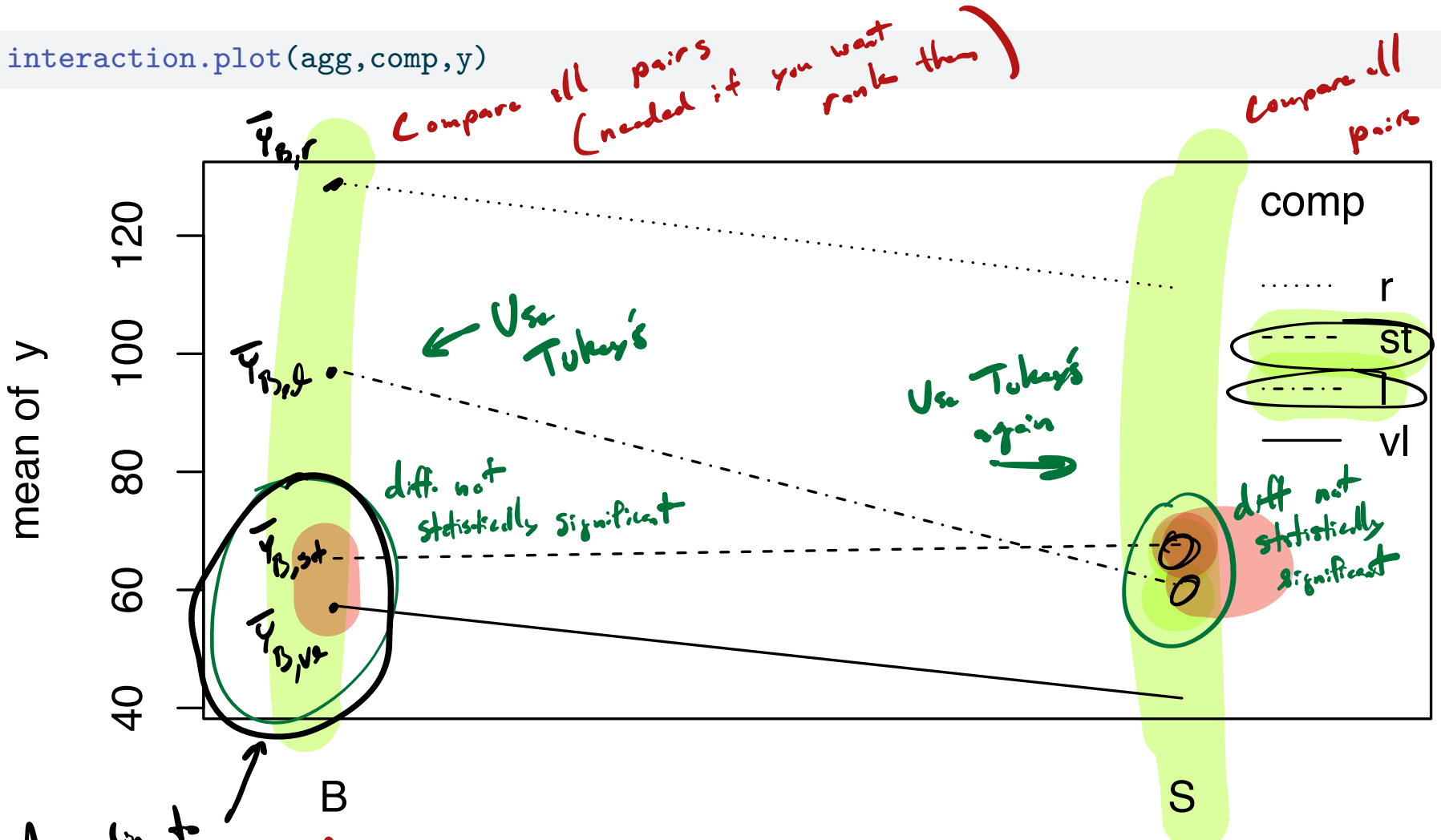
" " " "

	lower	upper
B:l-r	-39.75923299	-23.574100
B:l-st	23.90743367	40.092566
B:l-vl	31.90743367	48.092566
B:r-st	55.57410034	71.759233
B:r-vl	63.57410034	79.759233
B:st-vl	-0.09256633	16.092566
S:l-r	-58.42589966	-42.240767
S:l-st	-15.09256633	1.092566
S:l-vl	10.90743367	27.092566
S:r-st	35.24076701	51.425900
S:r-vl	61.24076701	77.425900
S:st-vl	17.90743367	34.092566

We applied Tukey's method twice, so we should divide α by two (Bonferroni correction).

Identify which differences are *not* significant in the interaction plot:

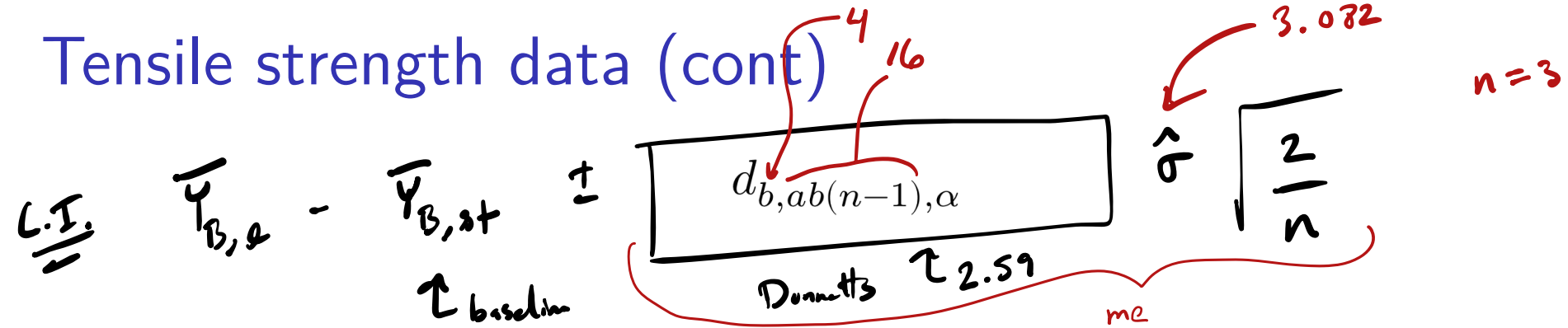
```
interaction.plot(agg, comp, y)
```



According to Dunnett's, this diff. is significant.

Using Tukey's two times, so divide α by 2.

Tensile strength data (cont)



For only the basalt aggregate type, compare all compaction method means to that of the static method.

Use $d_{4,16,0.05} = 2.59$ from the table on the next slide.

```
me <- 2.59 * sqrt(MSE) * sqrt(2/n)
dtab <- rbind(c(y11. - y13. - me, y11. - y13. + me),
             c(y12. - y13. - me, y12. - y13. + me),
             c(y14. - y13. - me, y14. - y13. + me))
rownames(dtab) <- c("B:l-st", "B:r-st", "B:vl-st")
colnames(dtab) <- c("lower", "upper")
```

Table A.5 Critical Values for Dunnett's Two-Sided Test of Treatments versus Control.

"Baseline"

Error df	Two-sided α	$T = \text{Number of Groups Counting Both Treatments and Control}$						
		2	3	4	5	6	7	8
5	0.05	2.57	3.03	3.29	3.48	3.62	3.73	3.82
5	0.01	4.03	4.63	4.97	5.22	5.41	5.56	5.68
6	0.05	2.45	2.86	3.10	3.26	3.39	3.49	3.57
6	0.01	3.71	4.21	4.51	4.71	4.87	5.00	5.10
7	0.05	2.36	2.75	2.97	3.12	3.24	3.33	3.41
7	0.01	3.50	3.95	4.21	4.39	4.53	4.64	4.74
8	0.05	2.31	2.67	2.88	3.02	3.13	3.22	3.29
8	0.01	3.36	3.77	4.00	4.17	4.29	4.40	4.48
9	0.05	2.26	2.61	2.81	2.95	3.05	3.14	3.20
9	0.01	3.25	3.63	3.85	4.01	4.12	4.22	4.30
10	0.05	2.23	2.57	2.76	2.89	2.99	3.07	3.14
10	0.01	3.17	3.53	3.74	3.88	3.99	4.08	4.16
11	0.05	2.20	2.53	2.72	2.84	2.94	3.02	3.08
11	0.01	3.11	3.45	3.65	3.79	3.89	3.98	4.05
12	0.05	2.18	2.50	2.68	2.81	2.90	2.98	3.04
12	0.01	3.05	3.39	3.58	3.71	3.81	3.89	3.96
13	0.05	2.16	2.48	2.65	2.78	2.87	2.94	3.00
13	0.01	3.01	3.33	3.52	3.65	3.74	3.82	3.89
14	0.05	2.14	2.46	2.63	2.75	2.84	2.91	2.97
14	0.01	2.98	3.29	3.47	3.59	3.69	3.76	3.83
15	0.05	2.13	2.44	2.61	2.73	2.82	2.89	2.95
15	0.01	2.95	3.25	3.43	3.55	3.64	3.71	3.78
16	0.05	2.12	2.42	2.59	2.71	2.80	2.87	2.92
16	0.01	2.92	3.22	3.39	3.51	3.60	3.67	3.73
17	0.05	2.11	2.41	2.58	2.69	2.78	2.85	2.90
17	0.01	2.90	3.19	3.36	3.47	3.56	3.63	3.69
18	0.05	2.10	2.40	2.56	2.68	2.76	2.83	2.89
18	0.01	2.88	3.17	3.33	3.44	3.53	3.60	3.66
19	0.05	2.09	2.39	2.55	2.66	2.75	2.81	2.87
19	0.01	2.86	3.15	3.31	3.42	3.50	3.57	3.63
20	0.05	2.09	2.38	2.54	2.65	2.73	2.80	2.86
20	0.01	2.85	3.13	3.29	3.40	3.48	3.55	3.60
25	0.05	2.06	2.34	2.50	2.61	2.69	2.75	2.81
25	0.01	2.79	3.06	3.21	3.31	3.39	3.45	3.51
30	0.05	2.04	2.32	2.47	2.58	2.66	2.72	2.77
30	0.01	2.75	3.01	3.15	3.25	3.33	3.39	3.44
40	0.05	2.02	2.29	2.44	2.54	2.62	2.68	2.73
40	0.01	2.70	2.95	3.09	3.19	3.26	3.32	3.37
60	0.05	2.00	2.27	2.41	2.51	2.58	2.64	2.69
60	0.01	2.66	2.90	3.03	3.12	3.19	3.25	3.29

2.59

This table produced from the SAS System using function PROBMC('DUNNETT2',...1 - α ,df,k), where $k = T - 1$.

Figure 1: Table A.5 from Mohr, Wilson, and Freund (2021)

Dunnett's comparison of compaction method means to the static method when the aggregate type is basalt:

dtab

	lower	upper
B:l-st	25.48198	38.518024
B:r-st	57.14864	70.184690
B:vl-st	-14.51802	-1.481976

None contain zero,
means all are different
from the baseline.

Interaction *not* significant. Then what?

If the interaction is not significant:

1. We can interpret main effects.
2. We can make meaningful comparisons among marginal means.

Comparing marginal means in the absence of interaction

For making comparisons among the marginal means of Factor A:

- ▶ Tukey's for comparing all pairs among $\bar{\mu}_1, \dots, \bar{\mu}_a$:

$$\bar{Y}_{i..} - \bar{Y}_{i'..} \pm q_{a, ab(n-1), \alpha} \hat{\sigma} \frac{1}{\sqrt{bn}}, \quad 1 \leq i < i' \leq a.$$

(When $a=2$, this comes from a t -distribution)

- ▶ Dunnett's for comparing $\bar{\mu}_1, \dots, \bar{\mu}_{a-1}$ to a control mean $\bar{\mu}_a$:

$$\bar{Y}_{i..} - \bar{Y}_{a..} \pm d_{a, ab(n-1), \alpha} \hat{\sigma} \sqrt{\frac{2}{bn}}, \quad i = 1, \dots, a - 1.$$

Still use $\hat{\sigma} = \sqrt{MS_{\text{Error}}}$.

Do likewise for making comparisons among $\bar{\mu}_{.1}, \dots, \bar{\mu}_{.b}$ of Factor B.

Method	Glucose level			Marginal means for Method
	1	2	3	
1	μ_{11}	μ_{12}	μ_{13}	$\bar{\mu}_{1\cdot}$
2	μ_{21}	μ_{22}	μ_{23}	$\bar{\mu}_{2\cdot}$
	$\bar{\mu}_{\cdot 1}$	$\bar{\mu}_{\cdot 2}$	$\bar{\mu}_{\cdot 3}$	$\bar{\mu}_{\cdot\cdot}$ (Overall)

Marginal means for Glucose

Build a C.I. for the diff $\bar{\mu}_{1\cdot} - \bar{\mu}_{2\cdot}$.

$$\bar{Y}_{i\cdot} - \bar{Y}_{i'\cdot} \pm q_{a, ab(n-1), \alpha} \hat{\sigma} \frac{1}{\sqrt{bn}}$$

$a=2$ $b=3$
 $ab(n-1) = 12$

$$\bar{\mu}_{1\cdot} - \bar{\mu}_{2\cdot} \pm t_{2, 12, 0.05} \hat{\sigma} \frac{1}{\sqrt{9}} = [0.042, 0.070]$$

95% C.I.
 t-key (0.95, 2, 12)
 " 3.081
 0.6135

Serum glucose example from Kuehl (2000)

Two methods for measuring serum glucose level at three glucose levels.

Glucose Level	Method 1			Method 2		
	1	2	3	1	2	3
	42.5	138.4	180.9	39.8	132.4	176.8
	43.3	144.4	180.5	40.3	132.4	173.6
	42.9	142.7	183.0	41.2	130.3	174.9

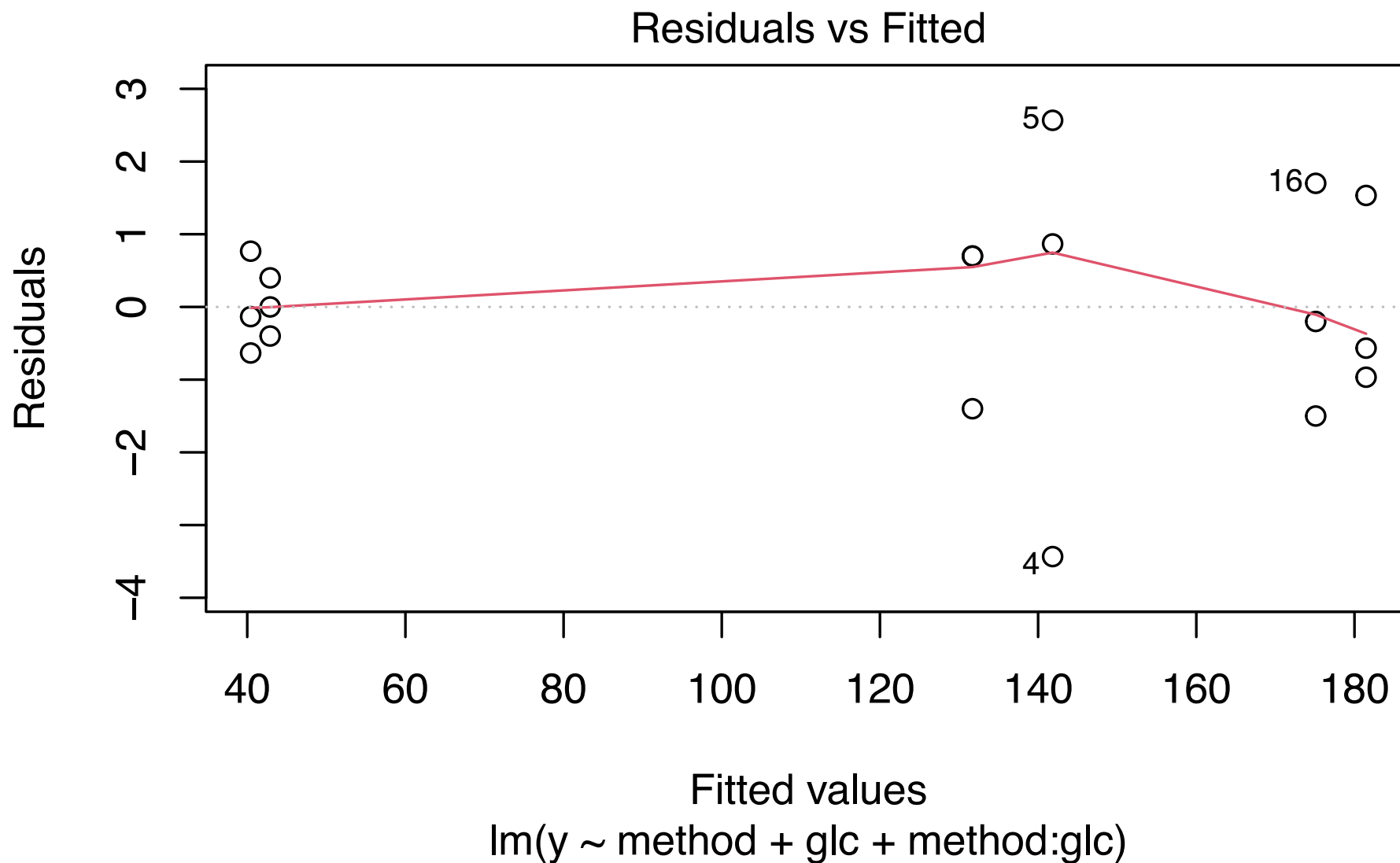
Source: Dr. J. Anderson, Beckman Instruments Inc.

Handwritten notes: $\bar{y}_{M_1..}$ (pointing to Method 1 data) and $\bar{y}_{M_2..}$ (pointing to Method 2 data)

```
y <- c(42.5, 43.3, 42.9, 138.4, 144.4, 142.7, 180.9, 180.5, 183.0,
       39.8, 40.3, 41.2, 132.4, 132.4, 130.3, 176.8, 173.6, 174.9)
method <- as.factor(c(1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 2, 2))
glc <- as.factor(c(1, 1, 1, 2, 2, 2, 3, 3, 3, 1, 1, 1, 2, 2, 2, 3, 3, 3))
```

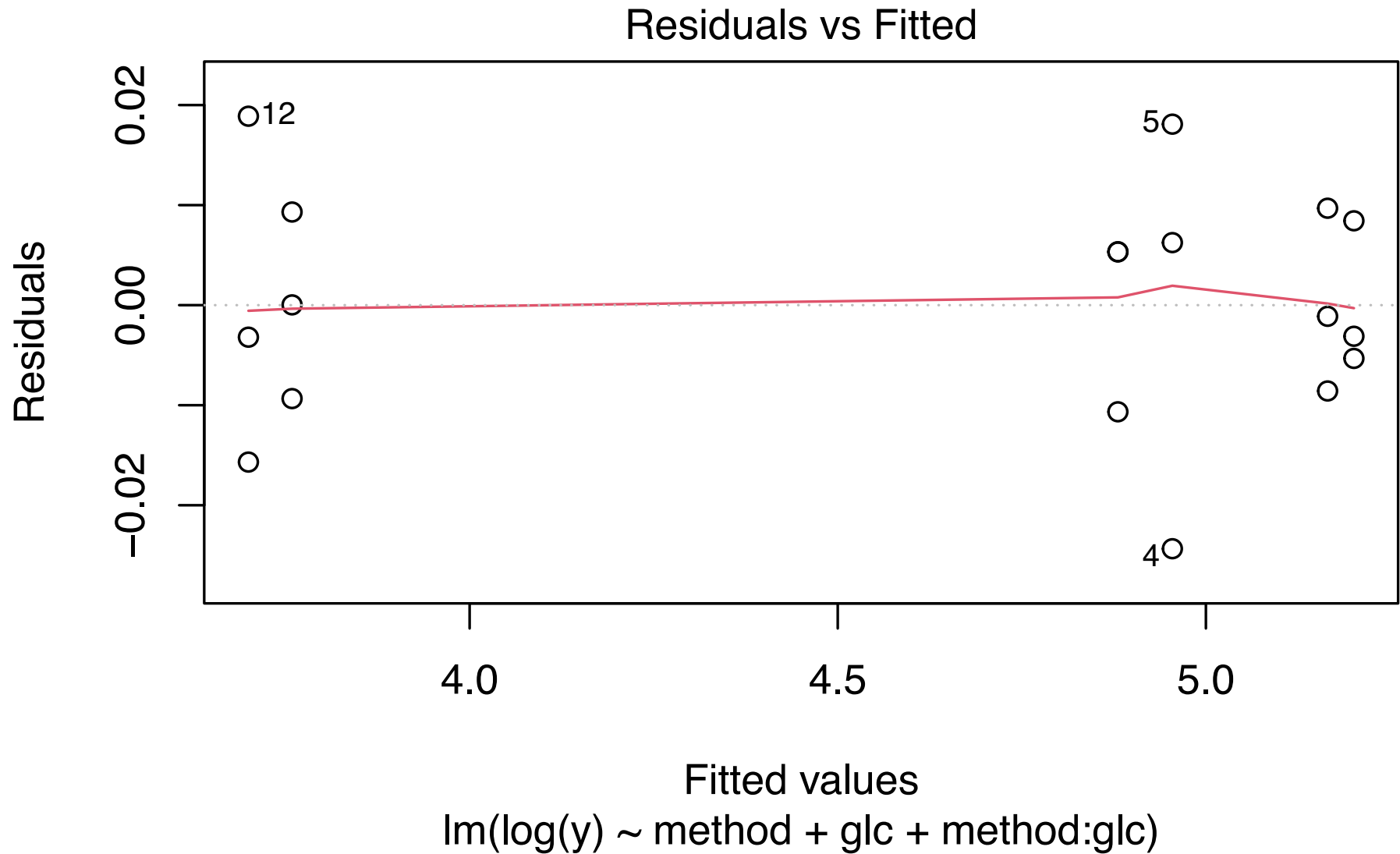
- ▶ Is there an interaction between the method and the glucose level?
- ▶ If not, can we describe the main effect of the method?

```
lm_glc <- lm(y ~ method + glc + method:glc)
plot(lm_glc, which = 1)
```



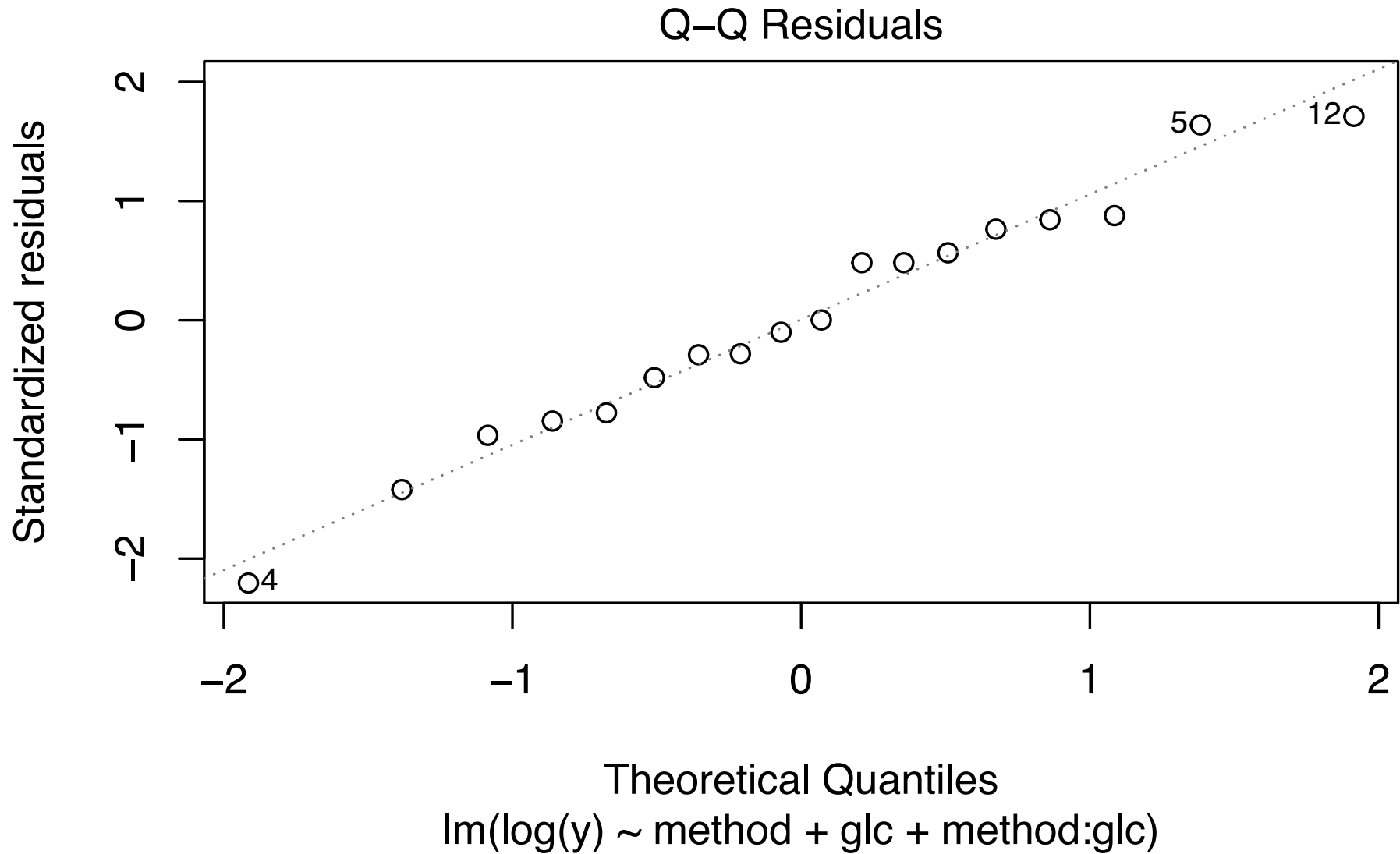
Variance appears smaller at lower glucose level. Try using $\log(Y_{ijk})$.

```
lm2_glc <- lm(log(y) ~ method + glc + method:glc)
plot(lm2_glc, which = 1)
```



This looks better.

```
plot(lm2_glc, which = 2)
```



Normality check looks okay.

Factor A = Method

Factor B = Glucose level

a = 2

b = 3

n = 3

```
anova(lm2_glc)
```

Glucose Level	Method 1			Method 2		
	1	2	3	1	2	3
	42.5	138.4	180.9	39.8	132.4	176.8
	43.3	144.4	180.5	40.3	132.4	173.6
	42.9	142.7	183.0	41.2	130.3	174.9

Source: Dr. J. Anderson, Beckman Instruments Inc.

Analysis of Variance Table

Response: log(y)

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
method	1	0.0143	0.0143	78.1091	1.337e-06 ***
glc	2	7.1935	3.5967	19670.4837	< 2.2e-16 ***
method:glc	2	0.0011	0.0006	3.0574	0.0845 .
Residuals	12	0.0022	0.0002		

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

$q-1=2-1$

$b-1=3-1$

$(a-1)(b-1) = (2-1)(3-1) = 2$

$ab(n-1) = 2 \cdot 3 \cdot (3-1) = 12$

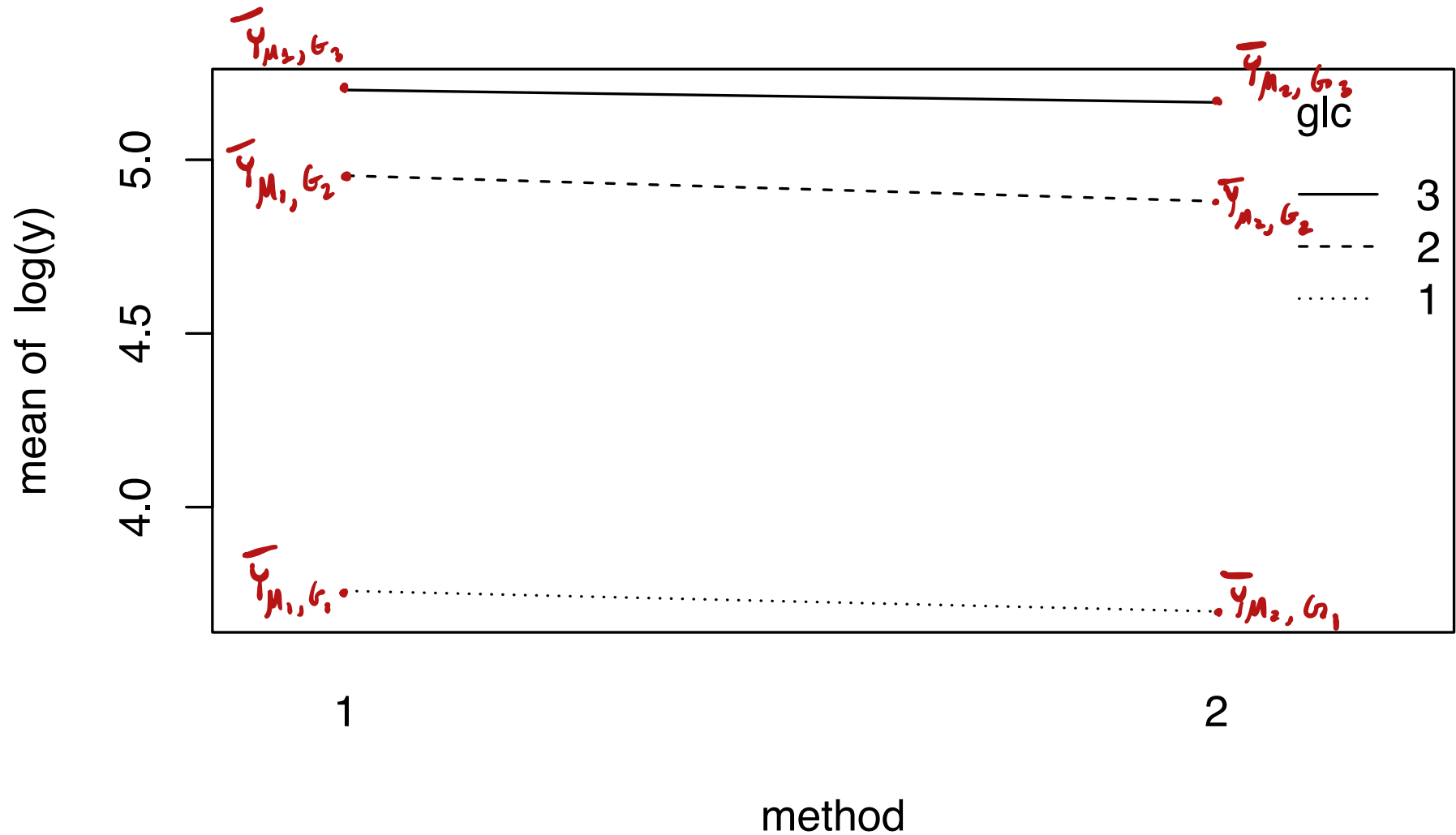
for testing H_0 : no interaction
 \Rightarrow Proceed as though interaction is not significant.

There is only weak evidence of interaction. Check interaction plot.

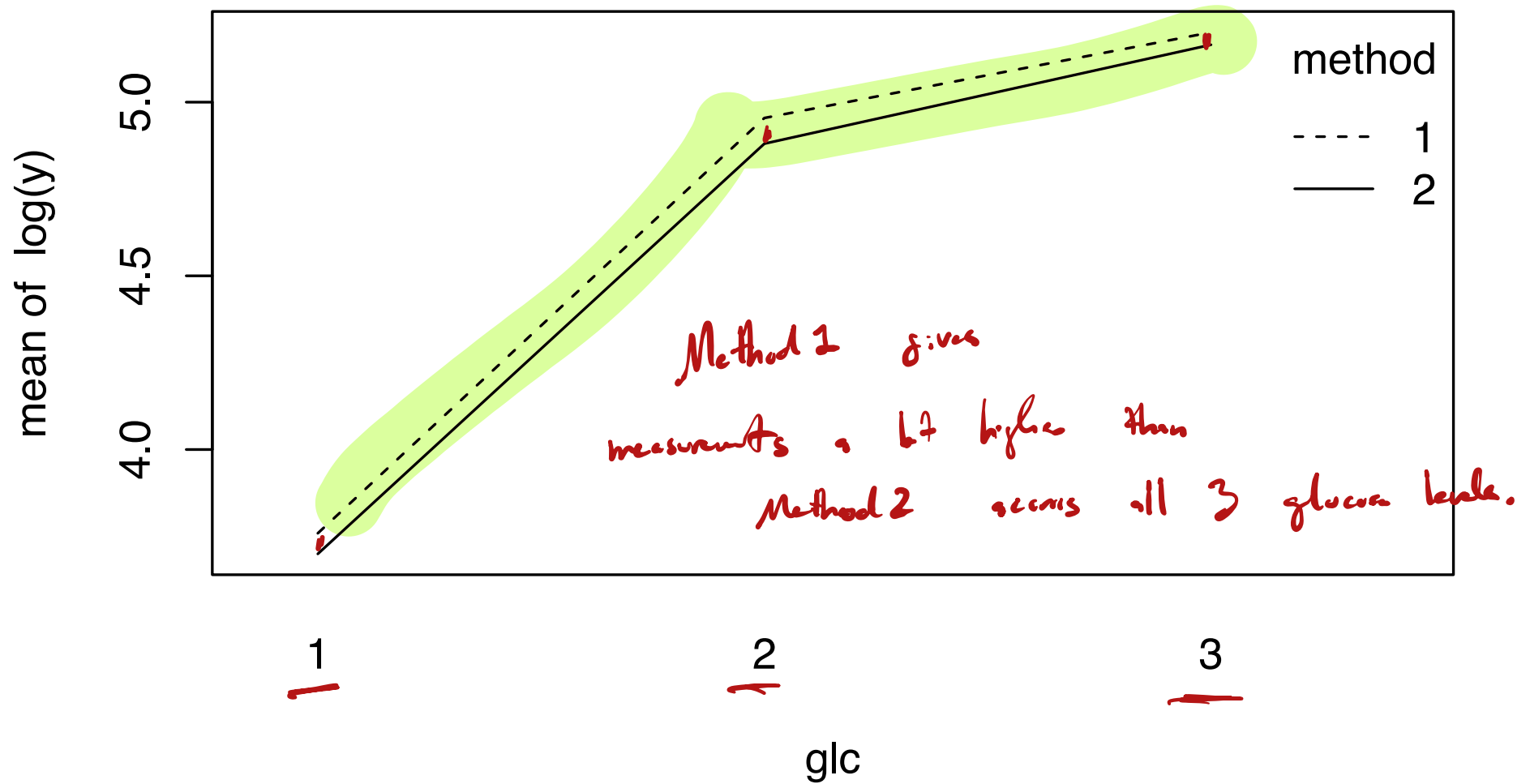
SS_{TOT}

Source	Df	SS	MS	F value
A	a - 1	SS _A	MS _A	$F_A = MS_A / MS_{Error}$
B	b - 1	SS _B	MS _B	$F_B = MS_B / MS_{Error}$
AB	(a - 1)(b - 1)	SS _{AB}	MS _{AB}	$F_{AB} = MS_{AB} / MS_{Error}$
Error	$ab(n - 1)$	SS _{Error}	MS _{Error}	
Total	abn - 1	SS _{Tot}		

```
interaction.plot(method, glc, log(y))
```



```
interaction.plot(glc,method,log(y))
```



It appears safe to ignore the interaction and report on main effects.

Call the method Factor A; build a CI for $\bar{\mu}_{1.} - \bar{\mu}_{2.}$ (just one comparison).

Since $a = 2$, $b = 3$, and $n = 3$, use $\bar{Y}_{1..} - \bar{Y}_{2..} \pm q_{2,2 \cdot 3(3-1),0.05} \hat{\sigma} \frac{1}{\sqrt{3 \cdot 3}}$.

```
a <- 2
b <- 3
n <- 3
alpha <- 0.05
y1.. <- mean(log(y[method == 1])) # remember we are using log(y)
y2.. <- mean(log(y[method == 2]))
MSE <- sum(lm2_glc$residuals^2) / (a*b*(n-1))
me <- qtukey(1-alpha,a,a*b*(n-1)) * sqrt(MSE) / sqrt(n*b)
lo <- y1.. - y2.. - me
up <- y1.. - y2.. + me
c(lo,up)
```

```
[1] 0.04244809 0.07022545
```

Since $a = 2$, $q_{a,ab(n-1),\alpha} = \sqrt{2} \cdot t_{ab(n-1),\alpha/2}$, so it is just a t -interval.

Possible workflow for factorial experiments

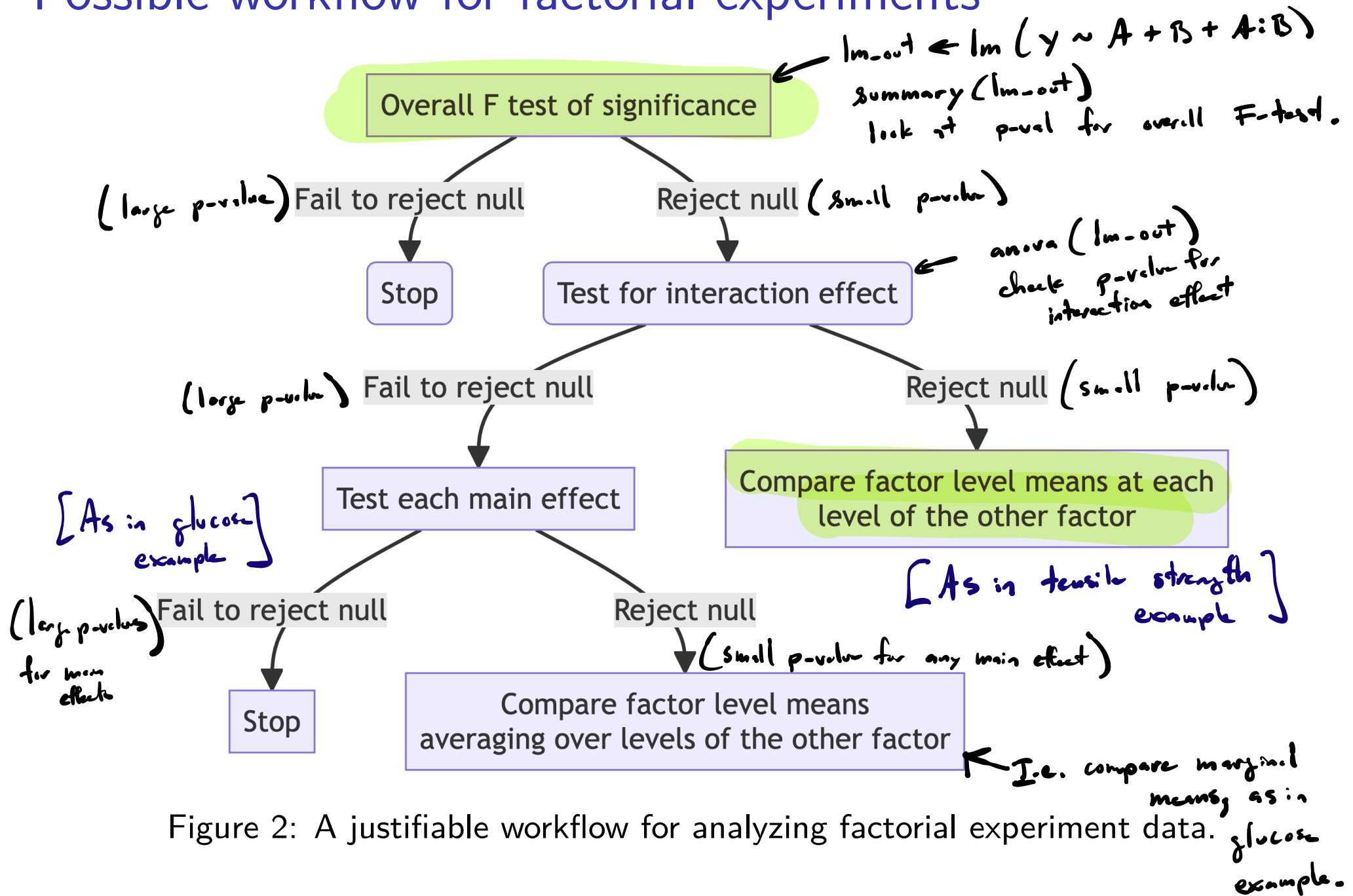


Figure 2: A justifiable workflow for analyzing factorial experiment data.

References

- Kuehl, R. O. 2000. *Design of Experiments: Statistical Principles of Research Design and Analysis*. Duxbury/Thomson Learning.
- Mohr, Donna L, William J Wilson, and Rudolf J Freund. 2021. *Statistical Methods*. Academic Press.