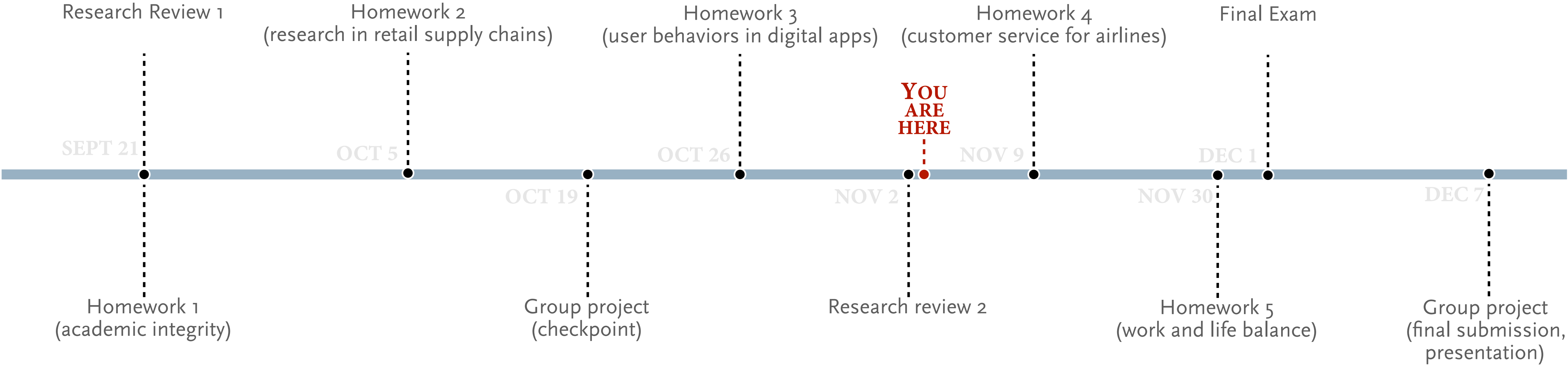


# Research Design, Fall 2021

**08: factorial experiments; analysis of variance; issues with multiple testing**



Initial questions?

factorial experiments

factorial experiments, investigating multiple explanatory *factors* or sets of treatments

factor ( $a, b, \dots$ )

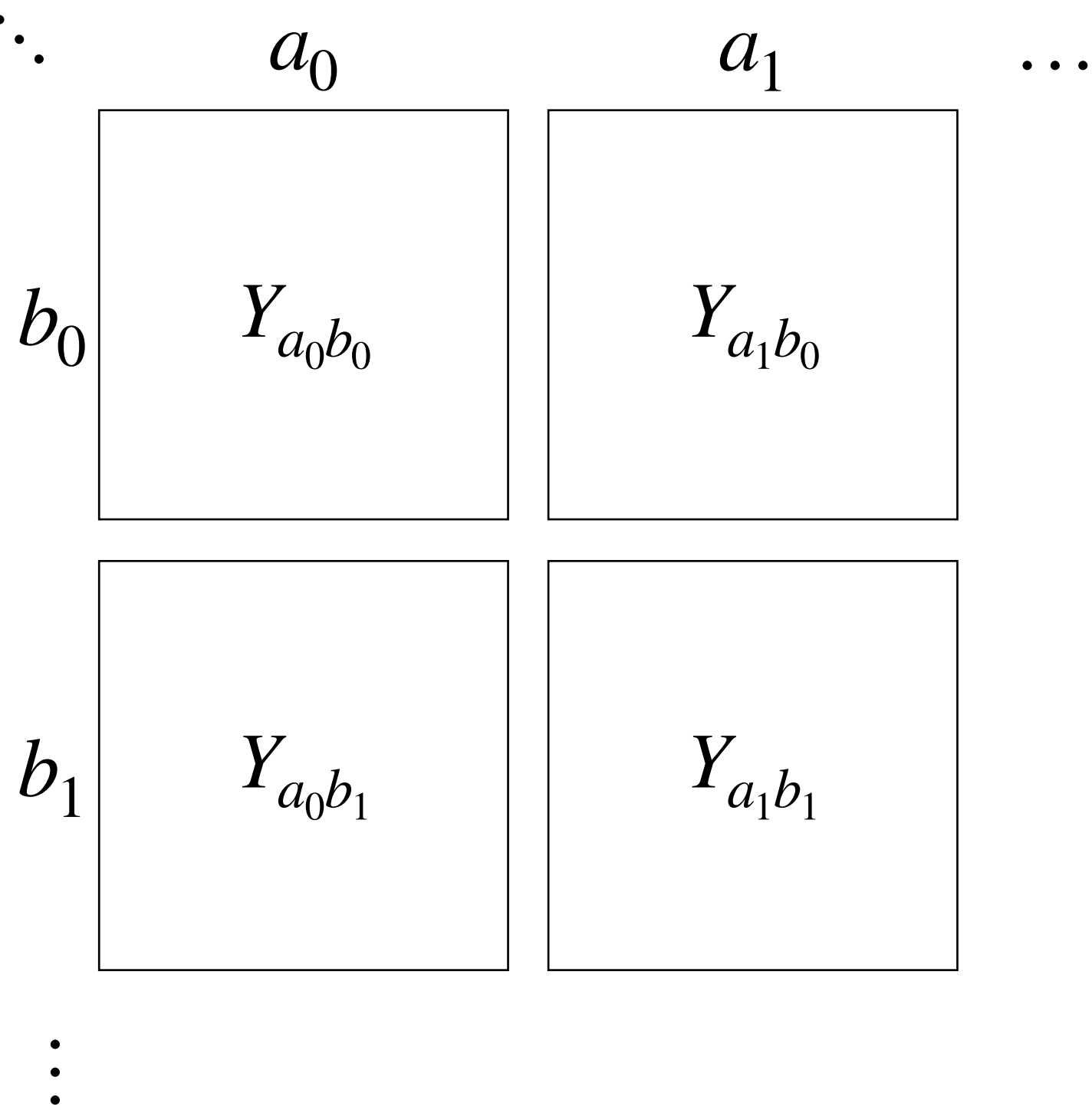
level ( $0, 1, \dots$ )

complete factorial experiment

fractional factorial experiment

main effects

interaction effects



main effect  $a = \frac{(a_1 b_0 - a_0 b_0) + (a_1 b_1 - a_0 b_1)}{2}$

interaction effect  $ab = \frac{(a_1 b_1 - a_1 b_0) - (a_0 b_1 - a_0 b_0)}{2}$

**analysis of variance (ANOVA)**

ANOVA, generally — analyzing variation among three or more means

ANOVA | This analysis compares *variance ratios* to determine whether or not significant *differences exist among the means of several groups* of observations, where *each group follows a normal distribution*.

An analysis-of-variance extends the t-test, which is used to determine whether or not two means differ, to the case where there are *three or more means*.

⋮	$a_0$	$a_1$	...
$b_0$	$Y_{a_0b_0}$	$Y_{a_1b_0}$	
$b_1$	$Y_{a_0b_1}$	$Y_{a_1b_1}$	
⋮			

main effect  $a = \frac{(a_1b_0 - a_0b_0) + (a_1b_1 - a_0b_1)}{2}$

interaction effect  $ab = \frac{(a_1b_1 - a_1b_0) - (a_0b_1 - a_0b_0)}{2}$

# ANOVA, assumptions — i.i.d. + equal variance + residuals normally distributed

ANOVA | This analysis compares *variance ratios* to determine whether or not significant *differences exist among the means of several groups* of observations, where *each group follows a normal distribution*.

An analysis-of-variance extends the t-test, which is used to determine whether or not two means differ, to the case where there are *three or more means*.

ASSUMPTIONS | inferences assume,  
units are *independent, identically distributed*  
variance is equal (*homoscedastic*) within each group  
errors are *normally distributed*



# ANOVA, the test — do differences exist in the means of groups not likely explained by sampling and variation?

$$H_0 : \theta_1 = \theta_2 = \dots = \theta_k \quad , \quad H_a : \theta_i \neq \theta_j \text{ for some } i, j$$

NOTE | we don't always — *or usually* — believe there is zero effect, nor would we find that to be interesting.

ANOVA, one-way analysis of variance

Source of variation	Degrees of freedom	Sum of squares	Mean square	<i>F</i> statistic
Between treatment groups	$k - 1$	$SS_B = \sum n_i(\bar{y}_i - \bar{y})^2$	$MS_B = \frac{SS_B}{k - 1}$	$F = \frac{MS_B}{MS_W}$
Within treatment groups	$N - k$	$SS_W = \sum \sum (y_{ij} - \bar{y}_{i.})^2$	$MS_W = \frac{SS_W}{N - k}$	
Total	$N - 1$	$SS_T = \sum \sum (y_{ij} - \bar{y})^2$		

Let,

- $N$  be the number of observations
- $k$  be the number of groups
- $i$  be a particular group
- $j$  be a particular observation in a group

# ANOVA, test — where does the $F$ -statistic fall in the $F$ -distribution?

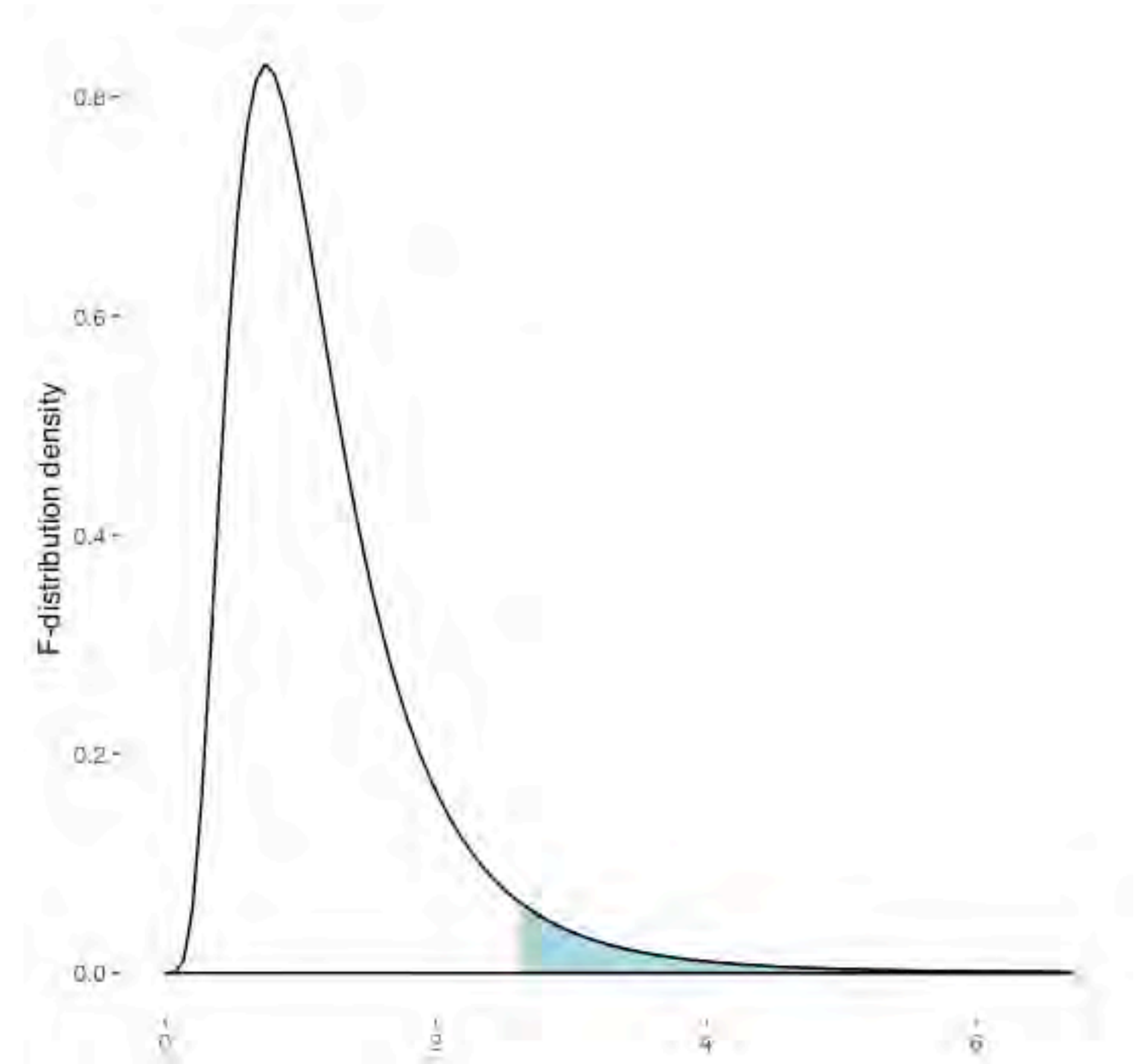
$$H_0 : \theta_1 = \theta_2 = \dots = \theta_k \quad , \quad H_a : \theta_i \neq \theta_j \text{ for some } i, j$$

```
library(ggplot2)
library(ggthemes)

df1 <- 15
df2 <- 12
alpha <- 0.05
xmax <- qf(0.001, df1, df2, lower.tail = FALSE)

ggplot() +
  theme_tufte(base_family = "sans") +
  stat_function(fun = df,
               args = list(df1 = df1, df2 = df2),
               geom = "density",
               fill = "white",
               xlim = c(0, xmax) ) +
  stat_function(fun = df,
               args = list(df1 = df1, df2 = df2),
               geom = "density",
               fill = "lightblue",
               xlim = c(qf(alpha, df1, df2, lower.tail = FALSE), xmax) ) +
  labs(y = "F-distribution Density")
```

```
F <- ____
P <- pf(F, df1, df2)
```



# ANOVA, example from pre-lecture notes (refactored for another approach)

```
# example in pre-lecture notes
library(dplyr)
dat <- read.csv("quiz video and text data.csv", header = TRUE)

# F-statistic
k <- nlevels( factor(dat$video) )
N <- nrow( dat )

SS_B <- dat %>%
  mutate(bar_quiz = mean(quiz)) %>%
  group_by(video) %>%
  summarise(sb = n() * (mean(quiz) - first(bar_quiz)) ^ 2) %>%
  ungroup() %>%
  summarise(SS_B = sum(sb)) %>% .$SS_B

SS_W <- dat %>%
  group_by(video) %>%
  mutate(sw = (quiz - mean(quiz)) ^ 2 ) %>%
  ungroup() %>%
  summarise(SS_W = sum(sw) ) %>%
  .$SS_W

Fstat <- ( SS_B / (k - 1) ) / ( SS_W / (N - k) )

# probability of this or greater variation in means from F-distribution
p <- pf(Fstat, df1 = k - 1, df2 = N - k, lower.tail = F)
```

Compare with R function, which relies on a linear regression model:

```
summary( aov(quiz ~ video, dat) )
```

Source of variation	Degrees of freedom	Sum of squares	Mean square	<i>F</i> statistic
Between treatment groups	$k - 1$	$SS_B = \sum n_i(\bar{y}_i - \bar{y})^2$	$MS_B = \frac{SS_B}{k - 1}$	$F = \frac{MS_B}{MS_W}$
Within treatment groups	$N - k$	$SS_W = \sum \sum (y_{ij} - \bar{y}_i)^2$	$MS_W = \frac{SS_W}{N - k}$	
Total	$N - 1$	$SS_T = \sum \sum (y_{ij} - \bar{y})^2$		

# ANOVA, example from pre-lecture notes (refactored for another approach)

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library(dplyr)
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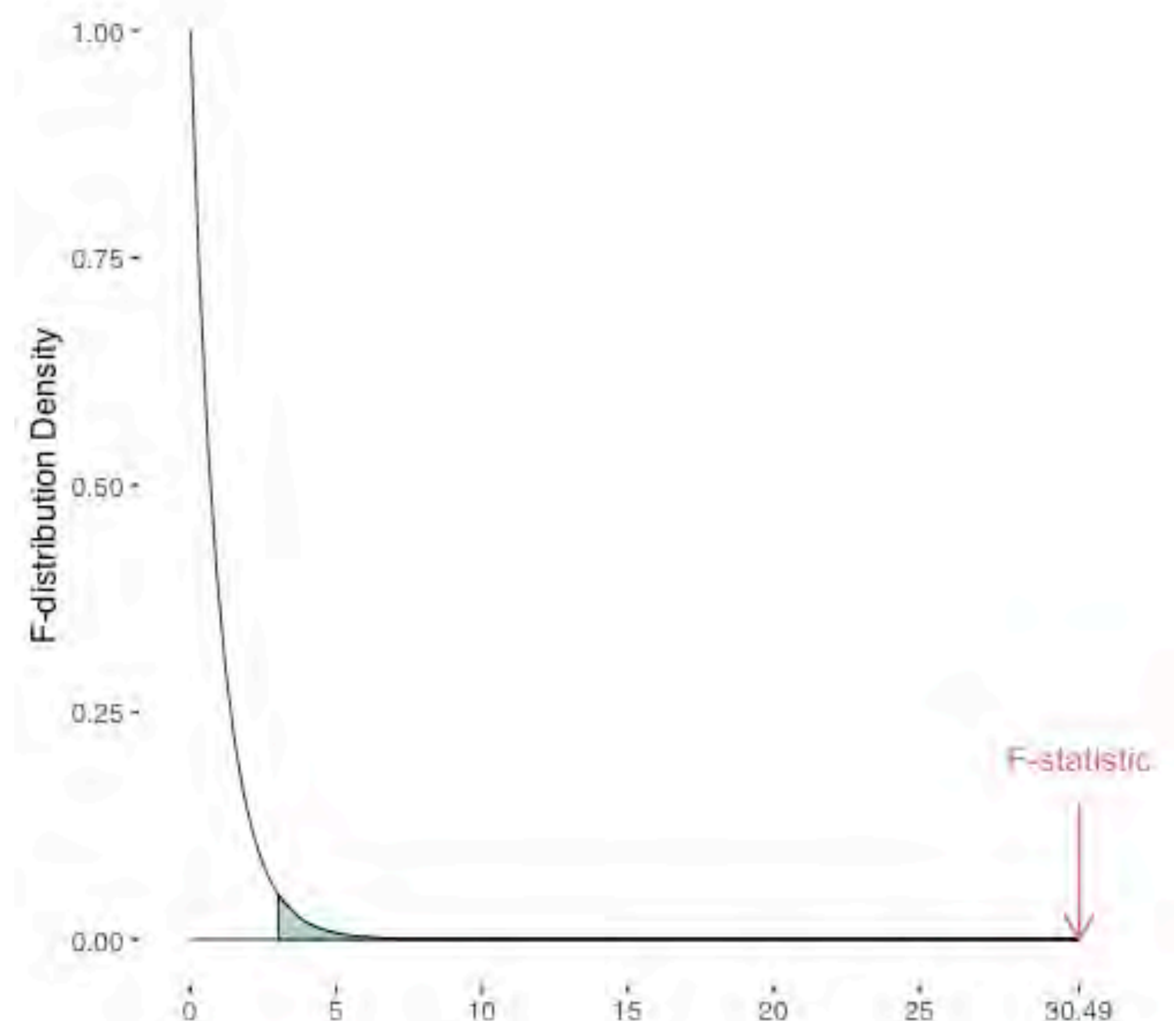
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  summarise(SS_W = sum(sw) ) %>%
  .$SS_W

Fstat <- ( SS_B / (k - 1) ) / ( SS_W / (N - k) )

# probability of this or greater variation in means from F-distribution
p <- pf(Fstat, df1 = k - 1, df2 = N - k, lower.tail = F)
```

Compare with R function, which  
relies on a linear regression model:

```
summary( aov(quiz ~ video, dat) )
```



# ANOVA, continuing example using two-way ANOVA with interaction

```
summary( aov(quiz ~ video + text + video:text, dat) )
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
video	2	2093	1046.5	39.61	< 2e-16	***
text	1	1530	1529.8	57.90	1.80e-13	***
video:text	2	1923	961.7	36.40	2.57e-15	***
Residuals	424	11202	26.4			

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

While ANOVA tests the probability of variation across all combinations of means if from a *F*-distribution ...

... it provides no information on what is usually *more important: effect sizes!* Use regression for those.

multiple hypothesis testing

In any test, because we're using a significance level,  $\alpha$ , we end up with *false positives* about that often.

As the tests multiply, so does the chance of getting false positives — **dramatically**.



For  $m$  number of tests, can  
adjust  $\alpha$  significance level:

$$\alpha_{\text{adj}} = \frac{\alpha}{m}$$

but with many tests, Bonferroni's  $\alpha_{\text{adj}}$  can also lead to *inflated false negatives*. Other methods are available, including Tukey's *Honest Significant Difference* test:

```
TukeyHSD( aov(quiz ~ video + text + video:text, data = dat), conf.level = 0.95 )
```

# References

**Abelson**, Robert P. *Statistics as Principled Argument*. Psychology Press, 1995.

**Casella**, George, and Roger L. Berger. “Analysis of Variance and Regression, Chp. 11.” In *Statistical Inference*. 2nd ed. Australia ; Pacific Grove, CA: Thomson Learning, 2002.

**Cox**, D. R., and N. Reid. “Factorial designs: basic ideas, Chp. 5.” In *The Theory of the Design of Experiments*. Monographs on Statistics and Applied Probability 86. Boca Raton: Chapman & Hall/CRC, 2000.

**Gelman**, Andrew, Tue Tjur, Peter McCullagh, Joop Hox, Herbert Hoijtink, and Alan M. Zaslavsky. “*Analysis of Variance? Why It Is More Important than Ever. With Discussion and Rejoinder.*” The Annals of Statistics 33, no. 1 (February 2005): 1–53.

**Gelman**, Andrew, and Eric Loken. “*The Statistical Crisis in Science.*” American Scientist 102 (November 2014): 1–6.