

# Multiple Linear Regression: Global tests and Multiple Testing

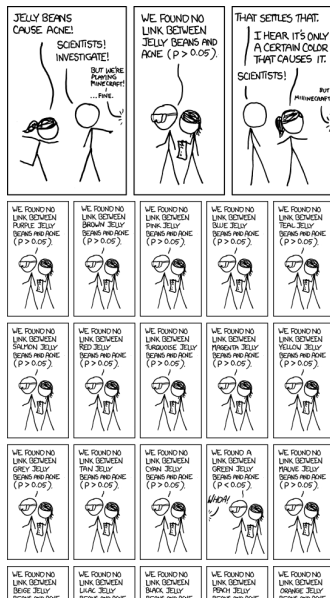
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# Today's Lecture

- Multiple testing - preserving your Type I error rate.



# Inference about multiple coefficients

Our model contains multiple parameters; often we want to perform multiple tests:

$$H_{01} : \beta_1 = 0$$

$$H_{02} : \beta_2 = 0$$

$$\vdots = \vdots$$

$$H_{0k} : \beta_k = 0$$

where each test has a size of  $\alpha$

- For any individual test,  $P(\text{reject } H_{0i} | H_{0i}) = \alpha$

## Inference about multiple coefficients

For any individual test,  $P(\text{reject } H_{0i} | H_{0i}) = \alpha$ .

But this DOES NOT MEAN that

$$P(\text{reject at least one } H_{0i} | \text{all } H_{0i} \text{ are true}) = \alpha$$

. This is called the Family-wise error rate (FWER). Ignore it at your own peril!

# Family-wise error rate

To calculate the FWER

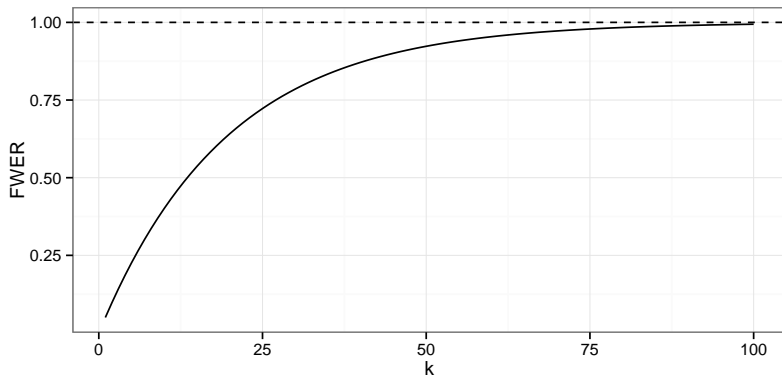
- First note  $P(\text{no rejections} | \text{all } H_{0i} \text{ are true}) = (1 - \alpha)^k$
- It follows that

$$\begin{aligned}\text{FWER} &= P(\text{at least one rejection} | \text{all } H_{0i} \text{ are true}) \\ &= 1 - (1 - \alpha)^k\end{aligned}$$

# Family-wise error rate

$$\text{FWER} = 1 - (1 - \alpha)^k$$

```
alpha <- .05  
k <- 1:100  
FWER <- 1-(1-alpha)^k  
qplot(k, FWER, geom="line") + geom_hline(yintercept = 1, lty=2)
```



# Addressing multiple comparisons

## Three general approaches

- Do nothing in a reasonable way
  - ▶ Don't trust scientifically implausible results
  - ▶ Don't over-emphasize isolated findings
- Correct for multiple comparisons
  - ▶ Often, use the Bonferroni correction and use  $\alpha_i = \alpha/k$  for each test
  - ▶ Thanks to the Bonferroni inequality, this gives an overall  $FWER \leq \alpha$
- Use a global test

# Global tests

Compare a smaller “null” model to a larger “alternative” model

- Smaller model must be nested in the larger model
- That is, the smaller model must be a special case of the larger model
- For both models, the  $RSS$  gives a general idea about how well the model is fitting
- In particular, something like

$$\frac{RSS_S - RSS_L}{RSS_L}$$

compares the relative  $RSS$  of the models



# Nested models

- These models are nested:

Smaller = Regression of  $Y$  on  $X_1$

Larger = Regression of  $Y$  on  $X_1, X_2, X_3, X_4$

- These models are not:

Smaller = Regression of  $Y$  on  $X_2$

Larger = Regression of  $Y$  on  $X_1, X_3$

## Global $F$ tests

- Compute the test statistic

$$F_{obs} = \frac{(RSS_S - RSS_L)/(df_S - df_L)}{RSS_L/df_L}$$

- If  $H_0$  (the null model) is true, then  $F_{obs} \sim F_{df_S - df_L, df_L}$
- Note  $df_S = n - p_S - 1$  and  $df_L = n - p_L - 1$
- We reject the null hypothesis if the p-value is above  $\alpha$ , where

$$\text{p-value} = P(F_{df_S - df_L, df_L} > F_{obs})$$

# Global $F$ tests

There are a couple of important special cases for the  $F$  test

- The null model contains the intercept only
  - ▶ When people say ANOVA, this is often what they mean (although all  $F$  tests are based on an analysis of variance)
- The null model and the alternative model differ only by one term
  - ▶ Gives a way of testing for a single coefficient
  - ▶ Turns out to be equivalent to a two-sided  $t$ -test:  $t_{df_L}^2 \sim F_{1,df_L}$

# Lung data: multiple coefficients simultaneously

You can test multiple coefficients simultaneously using the  $F$  test

```
mlr_null <- lm(disease ~ nutrition, data=dat)
mlr1 <- lm(disease ~ nutrition+ airqual + crowding + smoking, data=dat)
anova(mlr_null, mlr1)

## Analysis of Variance Table
##
## Model 1: disease ~ nutrition
## Model 2: disease ~ nutrition + airqual + crowding + smoking
##   Res.Df    RSS Df Sum of Sq    F    Pr(>F)
## 1       97 9192.7
## 2       94 1248.0  3    7944.7 199.47 < 2.2e-16 ***
## ---
## Signif. codes:
## 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

## Lung data: single coefficient test

The  $F$  test is equivalent to the  $t$  test when there's only one parameter of interest

```
mlr_null <- lm(disease ~ nutrition, data=dat)
mlr1 <- lm(disease ~ nutrition + airqual, data=dat)
anova(mlr_null, mlr1)

## Analysis of Variance Table
##
## Model 1: disease ~ nutrition
## Model 2: disease ~ nutrition + airqual
##   Res.Df    RSS Df Sum of Sq    F    Pr(>F)
## 1      97 9192.7
## 2      96 5969.5  1    3223.1 51.833 1.347e-10 ***
## ---
## Signif. codes:
## 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

summary(mlr1)$coef

##              Estimate Std. Error  t value    Pr(>|t|)
## (Intercept) 37.62538251  2.43946243  15.423637 9.946294e-28
## nutrition   -0.03469855  0.01692446  -2.050202 4.307101e-02
## airqual      0.36114435  0.05016218   7.199535 1.346935e-10
```

# Today's Big Ideas

$F$  tests can control for multiple comparisons!

- hands-on example