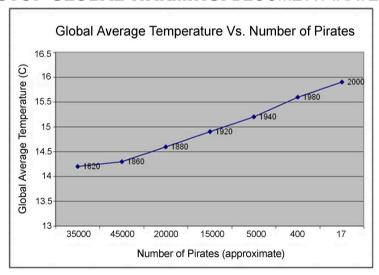
# Experimental Design and One-Way ANOVA

36-600

#### **Experiments and Causation**

- Throughout this course we have tried to uncover the association between a set of predictor variables  $\mathbf{x}$  and a response variable Y, if it exists
- However, even if a statistically significant association does exist...
  - ...association does not imply causation!

#### **STOP GLOBAL WARMING: BECOME A PIRATE**



WWW.VENGANZA.ORG

(image credit)

#### Case Study 1: Industrial Experiments

(The following case studies and the material on experimental design were provided by Professor Zach Branson)

- Apple wants to test the water durability of their laptops
  - they randomly sample 100 identical laptops for study, and pour water on half of them
  - 20 of 50 "treatment" (water-doused) keyboards continued to work, as opposed to 50 of 50 "control" keyboards
- Did the water *cause* the keyboards to break?
- Yes: the laptops were otherwise identical...the only difference was the treatment

### Case Study 2: Clinical Trials

- The Food and Drug Administration wanted to determine whether a new drug alleviated hypertension
  - they randomly picked 100 people with hypertension...
  - ...and placed 50 people each into the treatment and control groups
  - 30 of 50 treated people had alleviated hypertension, as opposed to 10 of 50 in the control group
  - a two-sample population proportions test yielded a *p*-value of 0.0001
- Did the new drug *cause* alleviated hypertension?
- We cannot be sure.
- Let's say it turns out that, totally by accident, the 50 people in the treatment group had health insurance, while the 50 people in the control group did not
  - so perhaps it was the drug, or the insurance (or the fact that those without insurance were poorer, or...)
  - o randomization leads to identical treatment and control groups, but only on average..."unlucky" randomization can happen

#### Case Study 3: Epidemiology

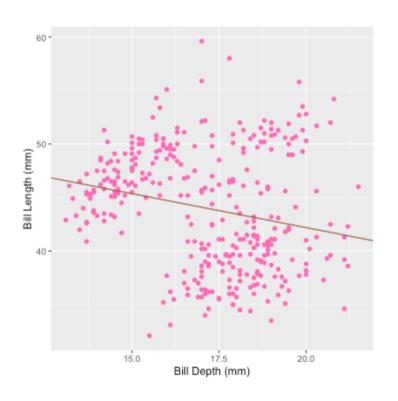
- We wish to study the effect of smoking on lung health
- Ideally, we would run an experiment in which we randomly place people into a smoking treatment group, and into a non-smoking control group
  - however, it is not ethical to force people to smoke
- So we randomly select 5000 smokers and 5000 non-smokers to study, and we find that the smokers have worse lung health
- Does smoking *cause* the observed deterioration of lung function?
- Again, we cannot be sure.
- For instance, smokers tend to be older, poorer, and to not have insurance
- Can we mitigate this issue?

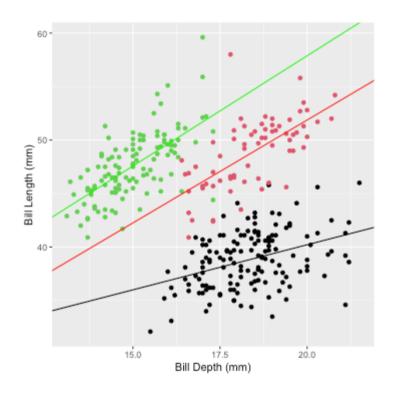
 $\rightarrow$  if we can identify subsets of treatment and control groups that have similar age, income, education, insurance, etc., and find similar results as before, we are in a position to argue more strenuously for causality

#### **Experiment Design**

- Let's assume that we want to divide N people into two groups, a treatment group and a control group...(assume N is an even number)
- How might we do this?
- Via Bernoulli trials: assign people to groups effectively via coin flips
  - issue: the group sizes can end up being very different
- Via complete randomization: pick exactly N/2 people at random to be in the treatment group
  - o issue: while this resolves the group-size issue observed with Bernoulli trials, there is still the issue of covariates
  - for instance, perhaps we want to run a clinical trial that includes both smokers and non-smokers: it could turn out that one group ends up with many more smokers than the other
  - the "covariate issue" can be dealt with when analyzing the data...or during the experiment design stage
- Via *block randomization*: identify a potentially problematic covariate, divide people into groups on the basis of that covariate (e.g., smokers vs. non-smokers), and then perform complete randomization within each covariate group...this scheme can be extended to multiple covariates: e.g., male smokers, female non-smokers, etc.
  - o covariate examples: gender, socioeconomic status, geographic location, medical risk factors, education, etc.

## Digression: Why Is Identifying Covariates Important?





#### How to Analyze Designed Experiment Data

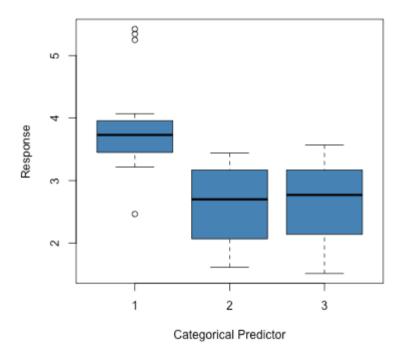
- Let's assume that our data consists of a predictor variable with k categories (or treatment groups) and a response variable.
- If k=2 and the response variable is normally distributed within each group, we can do a two-sample t test
- If k=2 and the response variable is not normally distributed, but the distributions are known or can be assumed, we can utilize other hypothesis tests like the population proportions test
  - o this is realm of so-called A/B testing...the test that is used depends on the distribution of the response values for groups A and B
- If k>2 and the response variable is normally distributed within each group (with equal variances), we can do one-way analysis of variance (or ANOVA)
  - o if there are two categorical predictors, we can do a two-way ANOVA
- If k > 2 and the response variable is normally distributed within each group (with equal variances), and there is another continuous predictor that can be treated as a covariate (e.g., age), we can do (one-way) analysis of covariance (or ANCOVA)
- Etc.
- **NOTE:** if your work potentially involves the design of experiments, consider taking 36-749, *Experimental Design for Behavioral and Social Sciences* (offered every fall)

### The One-Way ANOVA Setting

- Recall that a statistical model is a description of the data-generating process
- ullet In the simple linear regression setting, our data consists of a continuously valued predictor variable old x that we attempt to relate to the values of response variable old Y
  - however, what if instead the values of the predictor variable are discretely valued...and specifically, what if they represent groups (or categories)?
- If there are k>2 groups we would utilize one-way *analysis of variance* (or one-way *ANOVA*)
  - o "one-way" simply indicates that (in our chosen setting) there is only one (categorical) predictor variable

## Why Not Use Simple Linear Regression?

- Because categories/groups may have no natural numerical order
- If we were to apply linear regression alone, then switch the placement of groups 1 and 2 and apply linear regression again, the slope would change!



### Why Not Use Simple Linear Regression?

- Let's define the predictor variable x as a *factor variable*
- This causes R to change the definition of the model...
  - $\circ$  a factor variable with k levels is split into k-1 so-called *dummy variables*; the other level becomes the so-called *reference level*
- The linear regression model becomes

$$Y_i = eta_0 + \mathcal{I}_{x_i=2}eta_2 + \mathcal{I}_{x_i=3}eta_3 + \epsilon_i$$

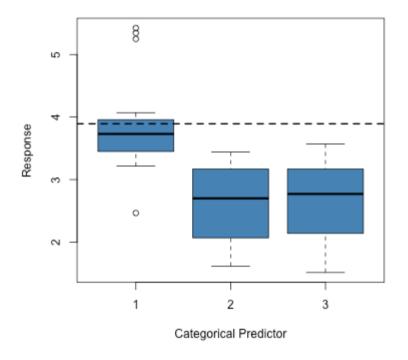
- $\mathcal{I}$  is the *indicator function*, and it takes on value 1 if the condition is true and 0 otherwise
  - $\circ~$  for instance,  $\mathcal{I}_{x_i=4}$  is 0 if  $x_i=3$  or 5 and 1 if  $x_i=4$
- We can rewrite the model in a form that might be more intuitive:

$$Y_i = \left\{egin{array}{ll} eta_0 & x_i = 1 \ eta_0 + eta_2 & x_i = 2 \ eta_0 + eta_3 & x_i = 3 \end{array}
ight.$$

• If we change the ordering of the groups, the  $\beta_i$ 's might change, but only because we perhaps define a new reference level

## Why Not Use Simple Linear Regression?

- The dashed line is  $eta_0$  and represents the predicted response for group 1
  - the negative coefficients for x2 and x3 in the lm() output indicate that the model is predicting that the means in groups 2 and 3 are smaller than the mean in group 1
- So...why not use simple linear regression? It turns out, we do use it...but the model definition changes since x is categorical and not quantitative
- So what then is ANOVA?
  - it is simply a mechanism for running a hypothesis test on linear regression output



### One-Way Analysis of Variance

• For one-way ANOVA, the statistical model is

$$Y_{ij} = \mu + au_i + \epsilon_{ij}$$

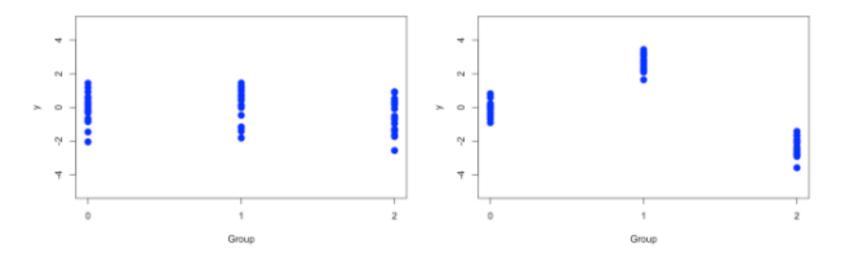
- $\circ \; i$  denotes the treatment group (there are k groups overall)
- $\circ \; j$  denotes an observed datum within group i
- $\circ \mu$  is the overall mean response
- $\circ \;\; au_i$  is the  $\mathit{deterministic}$  (i.e., not random) effect of treatment in group i
- $\circ~\epsilon_{ij}$  are the error terms, assumed to be independent, normally distributed, and of constant variance  $\sigma^2$ ...thus  $Y_{ij}\sim\mathcal{N}(\mu+ au_i,\sigma^2)$

#### ANOVA: Goal

• The goal of ANOVA is to perform the hypothesis test

$$H_o: \tau_1 = \cdots = \tau_k = 0$$
 vs. at least one value differs from zero

 $\circ~$  to reject the null, the value of one or more of the  $au_i$ 's has to be large with respect to  $\sigma$ 



• The left figure represents a situation in which we would fail to reject the null hypothesis, while the right figure represents a situation in which we would reject the null

#### ANOVA Example

anova(lm(Y~x))

- the first column shows k-1 and n-k (so we can determine that k=3 and n=32)
- the second column shows the SST (top) and SSE (bottom) (see appendix)
- the third columns shows the MST (= SST /( k-1); top) and MSE (= SSE /( n-k); bottom) (again, see appendix)
- the fourth column shows F=MST/MSE
- the fifth and last column shows the *p*-value
- We observe that  $p \ll \alpha = 0.05$ , so we reject the null hypothesis and conclude that at least one of the means is different from the others

#### But Which Mean is Different?

• To try to determine *which* of the means is different from the others, we can use a "post-hoc" test such as the Tukey HSD (honest significant difference) test

```
TukeyHSD(aov(Y~x)) # one quirk: it won't work with anova() output, just aov() output
## Tukey multiple comparisons of means
## 95% family-wise confidence level
```

- - the first column of output shows the groups being compared (2 vs. 1, etc.)
  - the second column gives the observed mean difference
  - the third and fourth columns provide confidence intervals on the *true* mean difference...if the interval does not contain zero, we conclude the means are different
  - the last column reinforces the confidence interval by providing a *p*-value (where the null is that the true difference is zero)

##

#### But Which Mean is Different?

```
TukeyHSD(aov(Y~x)) # one quirk: it won't work with anova() output, just aov() output
```

```
Tukey multiple comparisons of means
##
##
       95% family-wise confidence level
##
## Fit: aov(formula = Y ~ x)
##
## $x
##
              diff
                          lwr
                                     upr
                                             p adj
## 2-1 -1.27593333 -2.0039372 -0.5479295 0.0004657
## 3-1 -1.26000000 -2.2306718 -0.2893282 0.0088999
## 3-2 0.01593333 -0.9846122 1.0164789 0.9991476
```

- What do we conclude here?
  - o group 1 has a different mean from group 2, and from group 3...while the means in groups 2 and 3 are not significantly different

#### But Which Mean is Different?

- A final point to make is that the Tukey HSD test attempts to correct for *multiple comparisons*, i.e., for running many separate hypothesis tests
- If you run many tests, then you are more likely, by chance, to see p-values that are less than  $\alpha$  even if the null is always true
- The Tukey HSD test attempts to control the "family-wise error rate" such that if all the nulls are correct, the probability of seeing  $p < \alpha$  occur *once* is  $\alpha$ 
  - *however*, the algorithm for controlling the rate of false positives tends to be overly conservative
  - we will simply point out here that alternative test schemes are available, e.g., Dunnett's test, that one might want to explore using, particularly when the number of groups is large

### Appendix: Sum of Squares of Errors and of Treatment Groups

ullet We break the total sum of squared differences between each datum  $Y_{ij}$  and the overall mean  $ar{Y}$  into two pieces

$$\sum_{i=1}^k \sum_{j=1}^{n_i} (Y_{ij} - ar{Y})^2 = \sum_{i=1}^k \sum_{j=1}^{n_i} (Y_{ij} - ar{Y}_{iullet})^2 + \sum_{i=1}^k n_i (ar{Y}_{iullet} - ar{Y})^2 = SSE + SST$$

- $\circ \ \ ar{Y}_{iullet}$  is the sample mean of the data of group i
- $\circ n_i$  is the sample size in group i
- $\circ$  SSE is the sum of squares of the errors (where the "error" is how far each datum is from its group mean)
- $\circ$  SST is the sum of squares for each treatment group (how far each group mean is from the overall mean)

## Appendix: Hypothesis Testing

• We can form two statistics:

$$\frac{SSE}{\sigma^2}$$
 and  $\frac{SST}{\sigma^2}$ 

• Under the null hypothesis that  $au_1 = \dots = au_k = 0$ , we can write that

$$rac{SSE}{\sigma^2} \sim \chi^2_{n-k} ~~ ext{and}~~ rac{SST}{\sigma^2} \sim \chi^2_{k-1}$$

- $\circ \chi^2_{\nu}$  is a chi-square distribution for  $\nu$  degrees of freedom
- The following ratio defines a random variable that is sampled from an F distribution:

$$rac{SST/(k-1)}{SSE/(n-k)} = rac{MST}{MSE} = F \sim F_{k-1,n-k}$$

- $\circ k-1$  and n-k are the number of *numerator* and *denominator* degrees of freedom, respectively
- We reject the null hypothesis if the value of F is (far) larger than its mean value, (n-k)/(n-k-2) (which is pprox 1 if  $n\gg k$ )