### ECOLOGICAL DATA ANALYSIS IN R



### 11.1 Simple (bivariate) Linear Regression

Luis Malpica Cruz lmalpica@uabc.edu.mx - @luismalpicaC







### A case study







✓ PEER-REVIEWE

Sooty tern (*Onychoprion fuscatus*) survival, oil spills, shrimp fisheries, and hurricanes

Research article Conservation Biology Ecology Marine Biology Veterinary Medicine Zoology

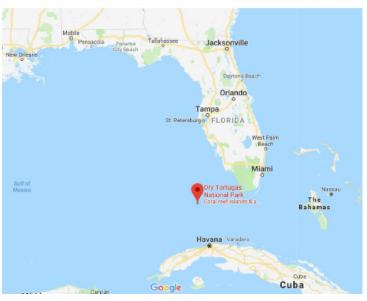
Ryan M. Huang <sup>1</sup>, Oron L. Bass Jr<sup>2</sup>, Stuart L. Pimm<sup>2</sup> <sup>1</sup>

Published May 10, 2017 PubMed 28503374

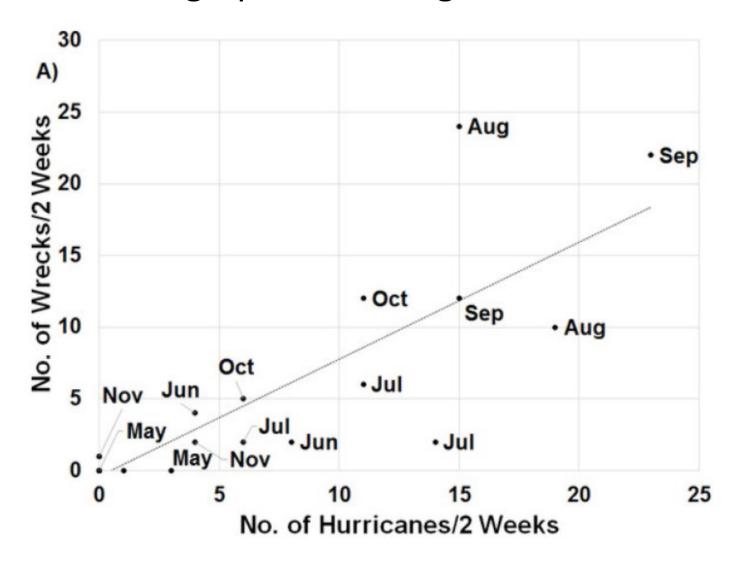
#### https://peerj.com/articles/3287/

We combined telemetry data on *Onychoprion* fuscatus (sooty terns) with a long-term capture-mark-recapture dataset from the Dry Tortugas National Park to map the movements at sea for this species, calculate estimates of mortality, and investigate the impact of hurricanes on a migratory seabird. ... Indices of hurricane strength and occurrence are positively correlated with annual mortality and indices of numbers of wrecked birds.





### Which Q is this graph answering?

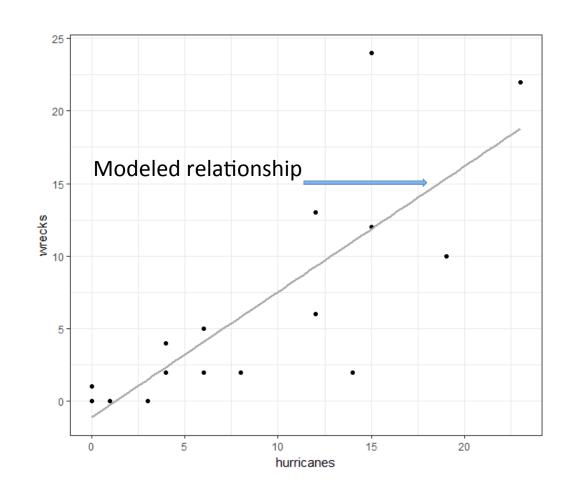


What is the relationship between hurricanes and wrecked birds?

### How is this made?

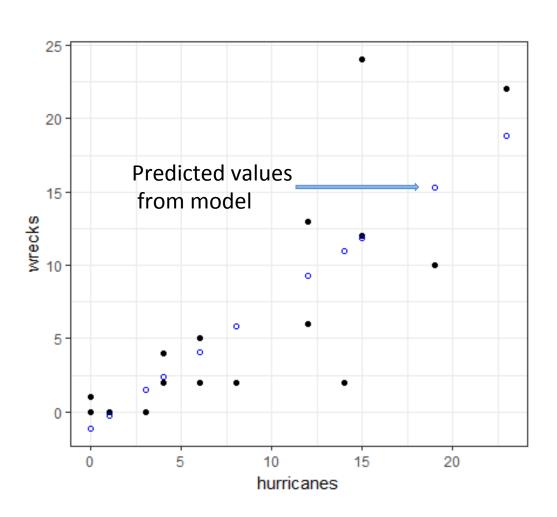
A line ("of best fit") is drawn through the points (data) minimizing the distance between the line and each point

This produces a modelled relationship between X and Y



### How is this made?

A 'predicted Y value' at each X value is calculated and connected with a line

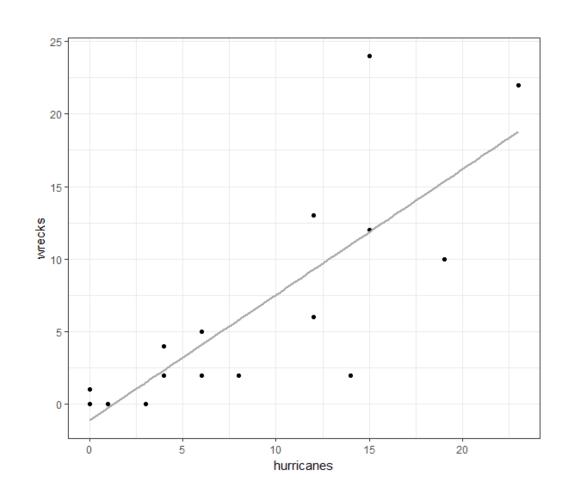


### How is this called?

This is a model, a mathematical description of a real-life relationship

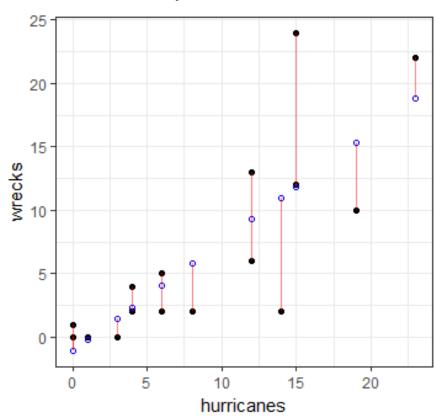
"All models are wrong, but some are useful"

- George Box, Statistician



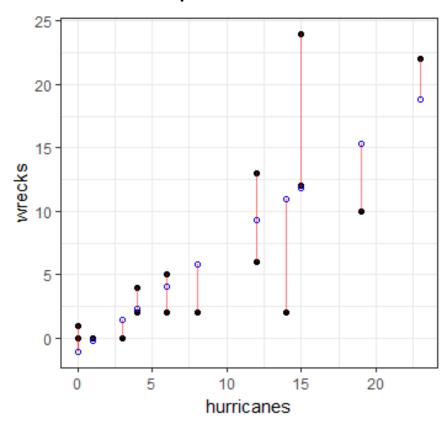
### Important terms: Residuals

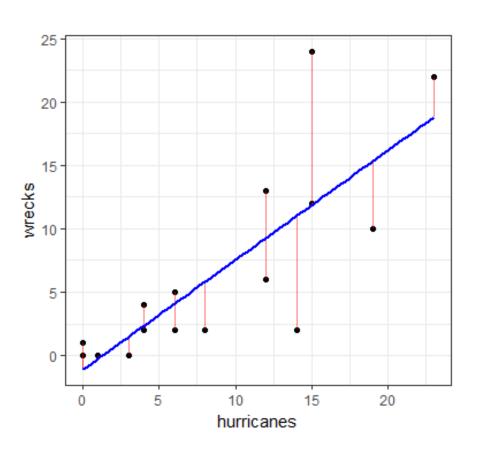
# Distance between the observed values and predicted values



### Important terms: Residuals

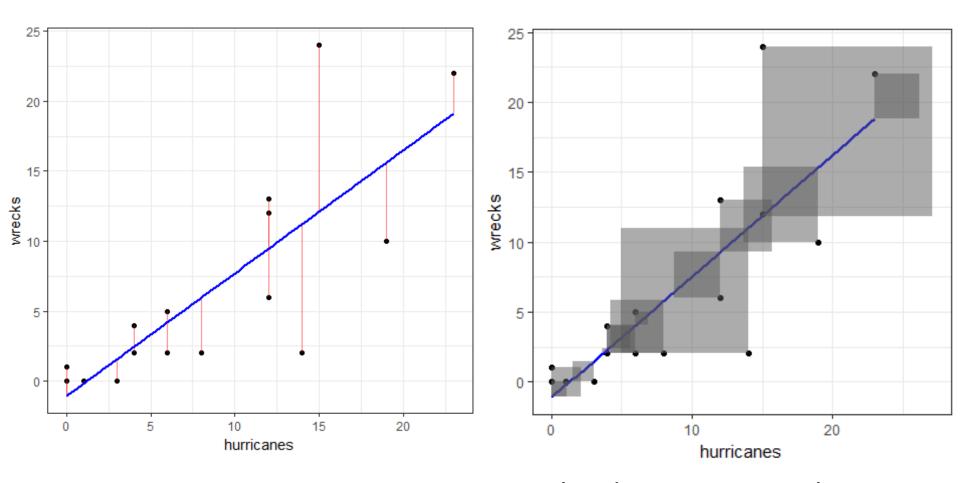
# Distance between the observed values and predicted values





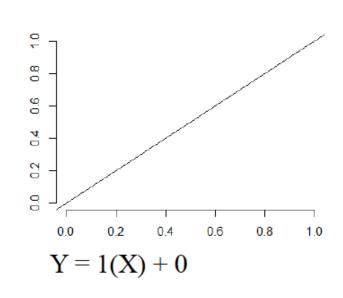
The line is fit so as to minimize residuals

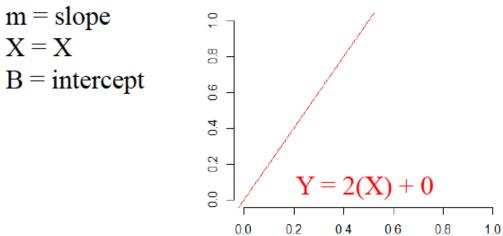
### Technically, to minimize the *Residual Sum of Squares* (RSS or SSR)



This demonstrates why outliers are so impactful!

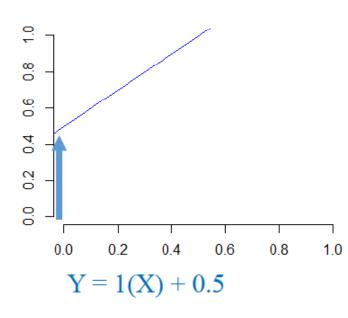
### Recall, how to graph a line?

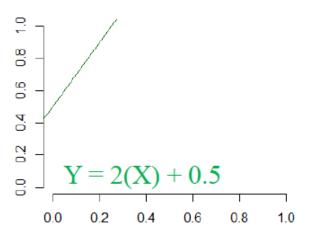




Y = mX + B

Y = Y





### Simple Linear regression model

AKA "Bivariate linear regression"

$$Y_i = \beta_o + \beta X_i + \epsilon_i$$

$$\beta_0$$
= Intercept (sometimes denoted  $\alpha$ )

$$\varepsilon$$
= residual error –information not

explained by model

$$\varepsilon_i \sim N(0, \sigma_i^2)$$

Error is normally distributed with mean of zero, variance  $\sigma_i^2$ 

Same form as:

Y = mX + B

## Example of a simple linear regression model

(Continuous or predictive (X) variable)

Spoken language: tern wrecks = intercept + hurricanes + error

Model (math) language: wrecks<sub>i</sub>=  $\beta_0$ +  $\beta_1$ hurricanes<sub>i</sub> + error<sub>i</sub>

R language: Im(wrecks ~ hurricanes, data = terns)



## Example of a simple linear regression model

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Spoken language: tern wrecks = intercept + hurricanes + error

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R language: Im(wrecks ~ hurricanes, data = terns)



Let's run a simple linear model in R...



### Model output interpretation

```
fit <- lm(wrecks ~ hurricanes, data=terns)
 call:
 lm(formula = wrecks ~ hurricanes, data = terns)
 Residuals:
   Min
           10 Median 30
                             Max
 -8.999 -2.372 0.195 1.773 12.135
 Coefficients:
            Estimate Std. Error t value Pr(>|t|)
 (Intercept) -1.1205 1.9699 -0.569 0.578500
 hurricanes 0.8657 0.1761 4.916 0.000228 ***
 Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
 Residual standard error: 4.796 on 14 degrees of freedom
 Multiple R-squared: 0.6332, Adjusted R-squared: 0.607
 F-statistic: 24.16 on 1 and 14 DF, p-value: 0.0002275
```

### Model fit, & coefficient magnitude Vs. significance

#### Coefficients:

Assess magnitude of effect on Y

Assess significance

### Assess model fit

```
Residual standard error: 4.796 on 14 degrees of freedom Multiple R-squared: 0.6332, Adjusted R-squared: 0.607 F-statistic: 24.16 on 1 and 14 DF, p-value: 0.0002275
```

### Model coefficient magnitude

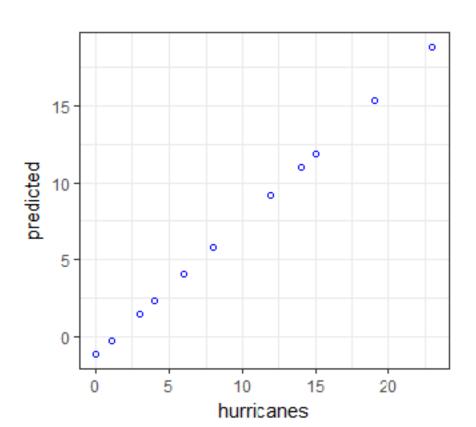
#### Coefficients:

Estimate Std. Error (Intercept) -1.1205 1.9699 hurricanes 0.8657 0.1761

$$Y = \beta_0 + \beta_1 X_1 + error$$

wrecks = -1.12 + 0.87 \* hurricanes

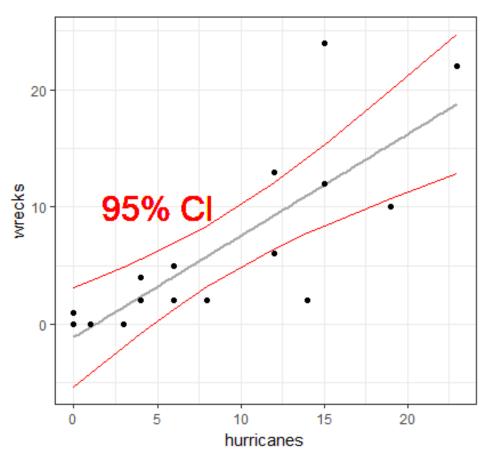
"For every additional hurricane, there were 0.87 more wrecks"



### 95% Confidence Interval of coefficient

If we repeated this study an infinite number of times, our interval would encapsulate the population mean at that X value 95% of the time

NOT: "95% of values fall within these bands"



### 95% CI of coefficient interpretation

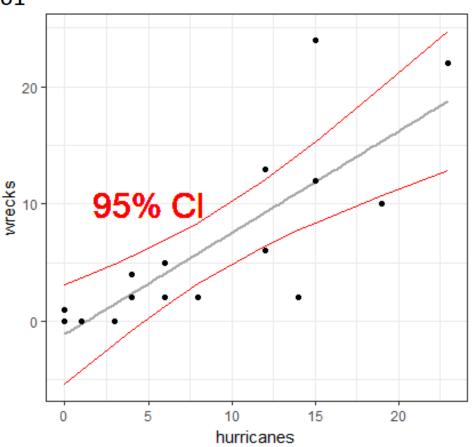
#### Coefficients:

Estimate Std. Error (Intercept) -1.1205 1.9699 hurricanes 0.8657 0.1761

Here, 95% C.I. of  $\beta_1$  = 0.8657  $\pm$  1.96 \* 0.1761

= 0.52 to 1.21

"For every additional hurricane, there were between 0.52 and 1.21 additional wrecked birds (95% C.I. = 0.87)"



### What if the 95% C.I. of a coefficient ( $\beta$ ) spans zero?

Y= 
$$\beta_o$$
+  $\beta_1$ \*X + error  
Y=  $\beta_o$ + 0\*X + error  
Y=  $\beta_o$ + error

Definition of 95% CI: If you did this experiment an infinite number of times, the population Beta would be encapsulated by the interval 95% of the time

In other words, we can't rule out that the 'true' beta value is zero

In OTHER words...

This variable has no statistically significant effect on Y

### Model coefficient significance

```
What's this?

Coefficients:

Estimate Std. Error t value Pr(>|t|)

(Intercept) -1.1205 1.9699 -0.569 0.578500

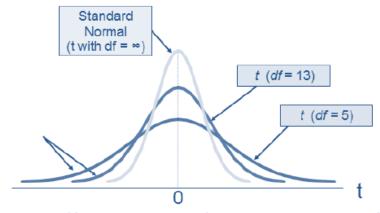
hurricanes 0.8657 0.1761 4.916 0.000228 **
```

"How many standard deviations is our coefficient away from zero?"

This is compared against the T-distribution

Larger absolute value of T ->
Further to the tails on the distribution ->
Lower P value

P < 0.05 = "Reject hypothesis that this parameter's value is zero"



https://financetrain.com/students-t-distribution/

### Model fit, residual standard error

```
Residual standard error: 4.796 on 14 degrees of freedom
Multiple R-squared: 0.6332, Adjusted R-squared: 0.607
F-statistic: 24.16 on 1 and 14 DF, p-value: 0.0002275
```

Average deviation of observed values from regression line

- When R fits a linear regression model, the sum of all residuals adds to zero

(intuitively: because there should be as many points "above" line as below, at roughly same overall distance)

- Therefore, calculate 'spread' of residuals by the following formula:

```
> sqrt(sum(residuals(fit)^2) / df.residual(fit))
[1] 4.796254
```

14 degrees of freedom (df) because... 16 data points 2 coefficients (intercept and slope) So... 16 – 2 = 14

### Model fit, R-squared (r<sup>2</sup>)

```
Residual standard error: 4.796 on 14 degrees of freedom
Multiple R-squared: 0.6332, Adjusted R-squared: 0.607
F-statistic: 24.16 on 1 and 14 DF, p-value: 0.0002275

% of variance in Y explainable by X
```

1 = perfect explanatory power (never happens)

0 = no explanatory power

As above, but penalizes model for having additional parameters

### Model fit, F-stat & p-value

```
Residual standard error: 4.796 on 14 degrees of freedom
Multiple R-squared: 0.6332, Adjusted R-squared: 0.607
F-statistic: 24.16 on 1 and 14 DF, p-value: 0.0002275
```

Variance explained by model

\_\_\_\_\_

Unexplained variance

Bigger F-stat means stronger evidence to reject null hypothesis

```
> anova(fit)
Analysis of Variance Table
```

```
Response: wrecks

Df Sum Sq Mean Sq F value Pr(>F)
hurricanes 1 555.88 555.88 24.165 0.0002275 ***
Residuals 14 322.06 23.00
```

Note: If you have large sample sizes, even an F-ratio of just over 1 may be significant

### Simple Linear regression model assumptions

$$Y_i = \beta_o + \beta X_i + \epsilon_i$$

$$\varepsilon_i \sim N(0, \sigma_i^2)$$

Y<sub>i</sub>= Response

X<sub>i</sub>= Explanatory variable

 $\beta_o$ = Intercept

β= Population slope

 $\varepsilon$ = residual error –information

not explained by model

#### **Assumptions:**

Assume error (i.e. residuals) is/are normally distributed with mean of zero, variance  $\sigma_i^2$ 

Assume  $\sigma_i^2$  is equal across the entire range of data

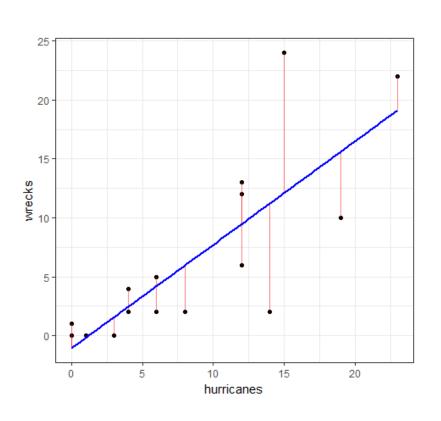
Assume replicates are truly independent.

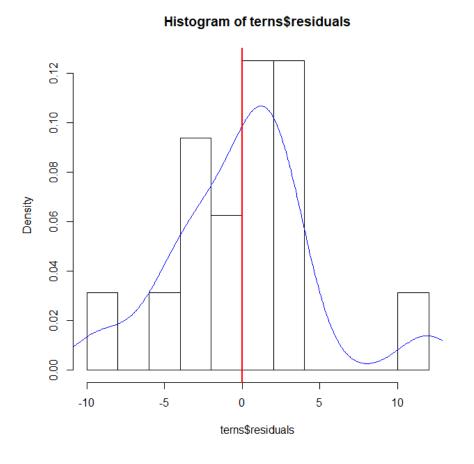
Y values at a given X should not influence Y values at other X positions

Assume fixed X

### Checking model assumptions, normality

 $\varepsilon_i \sim N(0, \sigma_i^2) \rightarrow$  Do our data meet this?





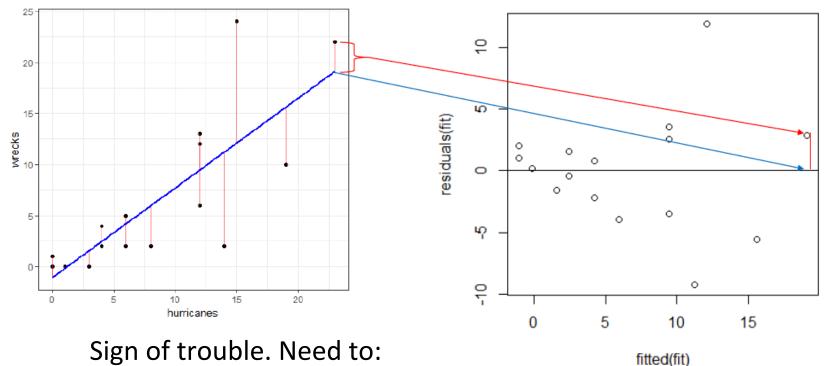
Yes-ish

### Checking model assumptions, homoscedasticity

Q: Is  $\sigma_i^2$  equal across the entire range of data?

Plot residuals vs. fitted values

A: No, we see bigger residuals at higher values



- Transform
  - Allow for different variance in Y across X (GLS)
  - Allow for different underlying distribution (GLM) ← later

### Checking model assumptions, independence

Assume replicates are truly independent.

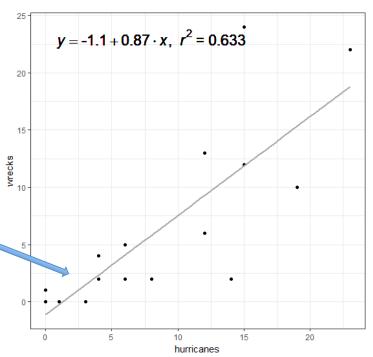
Y values at a given X should not influence Y values at other X positions

Dependence can be due to study design

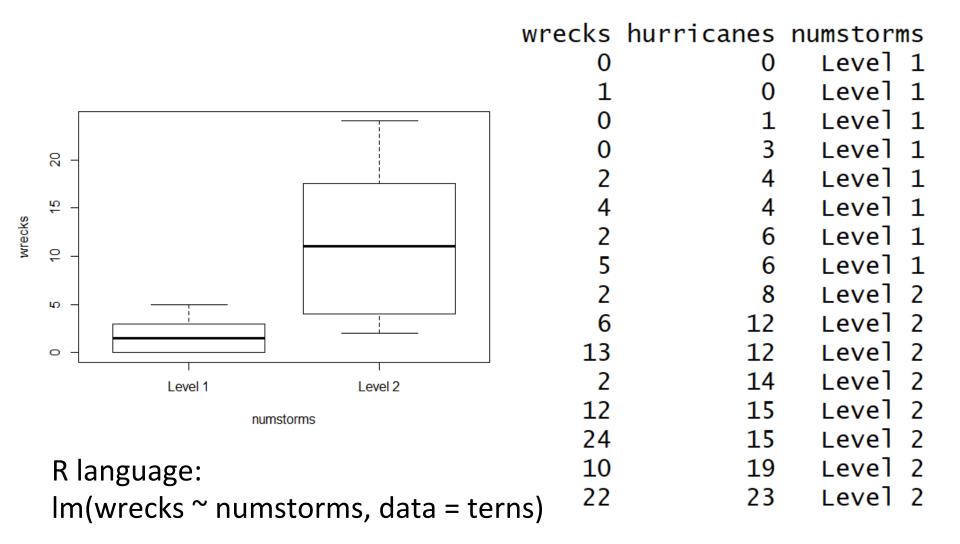
Dependence can also be due to poor model fit

At low X values, Y's are more similar than they are at high X values

Dependence due to model misfit



### Categorical predictive (X) variable (2 levels)



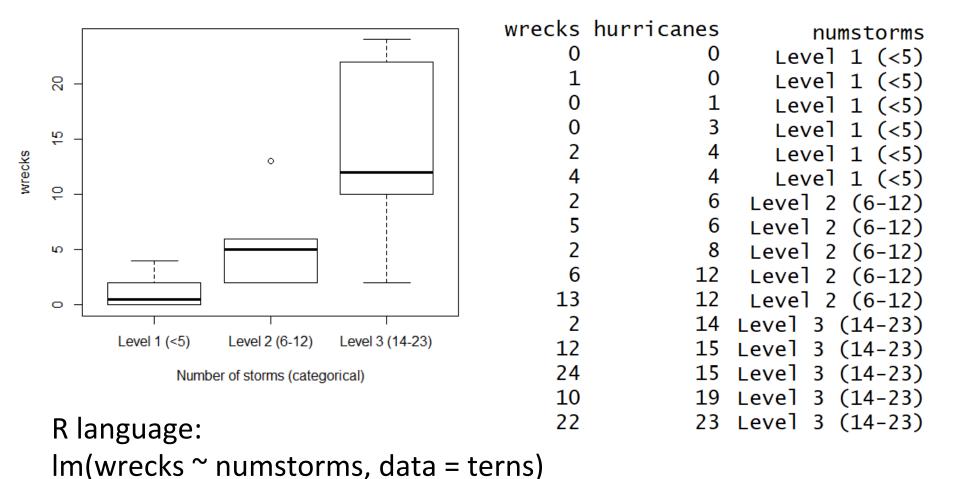
### If two levels in X, model output is like t-test

> categorical\_lm1 <- lm(wrecks ~ numstorms, data=terns2)</pre>

```
> summary(categorical_lm1)
call:
lm(formula = wrecks ~ numstorms, data = terns2)
Residuals:
   Min
            1Q Median
                                  Max
-9.375 -1.750 -0.250 1.781 12.625
                                                                           > t.test(terns2$wrecks ~ terns2$numstorms)
Coefficients:
                                                                                  Welch Two Sample t-test
                   Estimate Std. Error t value Pr(>|t|)
                                                                                 ters2$wrecks by terns2$numstorms
(Intercept)
                      1.750
                                  2.128
                                                                              -3.1976 df = 7.7388, p-value = 0.01322
numstormsLevel
                      9.625
                                  3.010
                                                   0.00645 **
                                                                            ternative hypothesis: true difference in means is not equal to 0
                                                                           95 percent confidence interval:
                   '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Signif. codes:
                                                                            -16.60713 -2.64287
                                                                           sample estimates.
                                                                           mean in group Level 1 mean in group Level 2
Residual standard error: 6.02 on 14 degrees of freedom
                                                                                                           11.375
                                                                                         1.750
Multiple R-squared: 0.4221,
                                   Adjusted R-squared: 0.3808
F-statistic: 10.22 on 1 and 14 DF, p-value: 0.006451
```

1.75 + 9.625 = 11.375

### Categorical predictive (X) variable (>2 levels)



### If >2 levels in X, model output is like ANOVA

```
> categorical_lm <- lm(wrecks ~ numstorms, data=terns3)</pre>
> summary(categorical_lm)
call:
lm(formula = wrecks ~ numstorms, data = terns3)
Residuals:
    Min
              10 Median
                                        Max
-12.0000 -2.4000 -0.8833 1.3333 10.0000
Coefficients:
                        Estimate Std. Error t value Pr(>|t|)
(Intercept)
                           1.167
                                     2.326 0.502 0.62436
numstormsLevel 2 (6-12)
                           4.433
                                     3.450
                                             1.285 0.22123
numstormsLevel 3 (14-23)
                                     3.450
                                            3.720 0.00257 **
                          12.833
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 5.698 on 13 degrees of freedom
Multiple R-squared: 0.3193, Aujusted K-squared: 0.4453
F-statistic 7.022 on 2 and 13 DF, p-value: 0.008555
> anova_version <- aov(wrecks ~ numstorms, data=terns3)</pre>
> summary(anova_version)
             Df Sum Sq Mean Sq F value Pr(>F)
            2 455.9 227.9
                               7.022 0.00856 **
numstorms
Residuals
             13 422.0
                         32.46
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

### If >2 levels in X, model output is like ANOVA

```
> categorical_lm <- lm(wrecks ~ numstorms, data=terns3)</pre>
> summary(categorical_lm)
call:
lm(formula = wrecks ~ numstorms, data = terns3)
Residuals:
    Min
              10
                  Median
                                        Max
-12.0000 -2.4000 -0.8833 1.3333 10.0000
Coefficients:
                        Estimate Std. Error t value Pr(>|t|)
(Intercept)
                           1.167
                                      2.326
                                             0.502 0.62436
numstormsLevel 2 (6-12)
                           4.433
                                      3.450
                                              1.285 0.22123
                                      3.450
numstormsLevel 3 (14-23)
                          12.833
                                              3.720 0.00257 **
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
Residual standard error: 5.698 on 13 degrees of freedom
Multiple R-squared: 0.5155, Adjusted K-squared:
F-statistic 7.022 on 2 and 13 DF, p-value: 0.008555
```

Interpreting  $\beta$ 's between levels:

As you go from Level 1 to Level 2, model predicts 4.43 more wrecks

1.167 + 4.43 = 5.6 wrecks at X = 2

As you go from Level 1 to Level 3, model predicts 12.83 more wrecks

 $1.167 + 12.83 = ^14.0$  wrecks at X = 3