Regression

ES/STT 7140: Statistical Modeling for Environmental Data

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- Unexcelled accuracy
- Capable of handling large data sets
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- What is the shape of the data (i.e., how does it cluster?)
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-Leo Breiman

Statistical models

• In the one-sample t-test, we are interested in learning about the mean of a normal distribution/population

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ight), \quad i=1,2,\dots,n$$

- For example, y might represent the shell length of a randomly selected zebra mussel from a stream or lake in Michigan
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- ∘ data = mean + remainder
- The above two expressions are mathematically equivalent
- The remainder is the difference between the observed values and the mean, often refered to as the residuals

• In the two-sample *t*-test problem, we are interested in the difference between the means of two populations (or groups):

$$egin{aligned} y_{1i} &\sim N\left(\mu_1, \sigma^2
ight), \quad i=1,2,\ldots,n_1 \ \ y_{2j} &\sim N\left(\mu_2, \sigma^2
ight), \quad j=1,2,\ldots,n_2 \ \ \delta &= \mu_2 - \mu_1 \end{aligned}$$

• As a linear model, we could use

$$y_k = \mu_1 + \delta g_k + \epsilon_k, \quad \sim N\left(0, \sigma^2
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- Illustration: clams.R
- A similar approach can be used for ANOVA procedures as well

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 - nonlinear regression (NLR) models
- In the next chapter, we will look at a more general class of regression models called *generalized linear models* (GzLMs)
 - GzLMs include both *logistic regression* and *Poisson regression* models

Simple linear regression

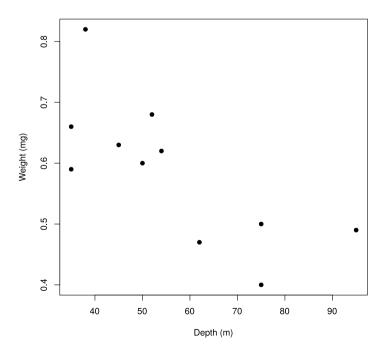
- A regression model is a formal means of expressing the two essential ingredients of a statistical model:
 - 1. A tenancy of the **response variable**, y, with a **predictor variable**, x, in some systematic fashion
 - 2. A scattering of points around the hypothesized curve of statistical relationship
- These two characteristics are embodied in a regression model by postulating that:
 - 1. There is a **probability distribution** of y for each level of x
 - 2. The means of these probability distributions vary in some systematic fashion with x

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 - For instance, we may want to determine the effect of increasing levels of DDT on eggshell thicknesses. How does increasing levels of DDT effect eggshell thickness?
- Another common use of regression models is to predict a response
 - For instance, if water is contaminated with a certain level of toxin, can we predict the amount of accumulation of this toxin in a fish that lives in the water?

- Many statistical applications deal with modeling how a single response variable, denoted y, depends on a single predictor, denoted x
- Illustration: shrimp.R



- From the previous *scatterplot*, one can see a fairly strong relationship between between the weight of the *Diporeia* and the depth of water where the Diporeia are found
- The scatterplot suggests that a straight line relationship between weight of Diporeia (\$y\$) and water depth (\$x\$) may be a reasonable way to model the data:

$$weight = eta_0 + eta_1 depth$$

- Here, β_0 is the *y*-intercept of the regression line and β_1 is the slope (i.e., rate of change)
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$$weight_i = \beta_0 + \beta_1 depth_i + \epsilon_i, \quad i = 1, \dots, 11$$

Assumptions of the SLR model

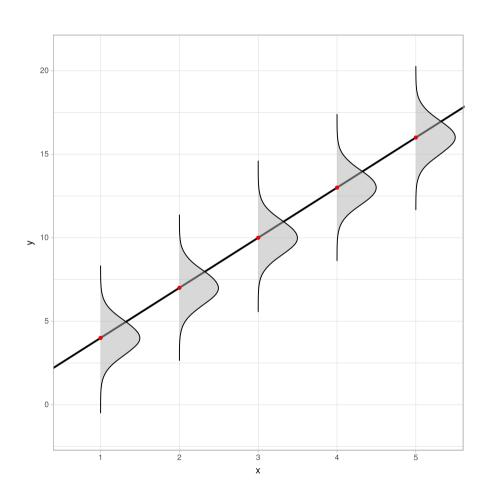
• The SLR model

$$y_i = eta_0 + eta_1 x_i + \epsilon_i, \quad i = 1, \dots, n$$

assumes that

- 1. Independent observations (i.e., the random errors are independent)
- 2. The errors have constant variance (i.e., *homoscedasticity*)
- 3. The errors are normally distributed (for statistical inference)
- If these assumptions are not met, then alternative methods need to be applied (e.g., weighted least squares or mixed-effects models)

Assumptions of the SLR model



- How do we estimate the model coefficients β_0 and β_1 ?
- There are an infinite number of lines passing through the data points $\{x_i, y_i\}_{i=1}^n$
- The *least squares* (LS) solution seeks to find β_0 and β_1 that minimize the *sum of squares*:

$$SS\left(eta_{0},eta_{1}
ight)=\sum_{i=1}^{n}\left(y_{i}-eta_{0}-eta_{1}x_{i}
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- So, how do we minimize $SS(\beta_0, \beta_1)$?
 - CALCULUS!!

• The values of β_0 and β_1 that minimize $SS(\beta_0, \beta_1)$ are given by

$$\circ$$
 $\widehat{eta}_1 = rac{\sum_{i=1}^n (x_i - ar{x})(y_i - ar{y})}{\sum_{i=1}^n \left(x_i - ar{x}
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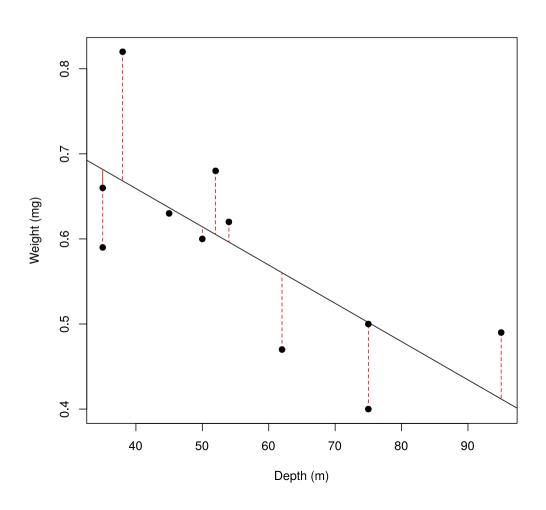
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- These are called the LS estimators of β_0 and β_1
- Under the usual assumptions for the SLR model (normality not required), the LS estimators:
 - Are **unbiased** estimators of β_0 and β_1
 - Have **minimum variance** among all *linear* unbiased estimators of β_0 and β_1 !
- How do we interpret $\widehat{\beta}_0$ and $\widehat{\beta}_1$ for a fitted SLR model?



- The SLR model is belongs to a broad class of models called *linear models* (LMs)
 - In an LM, the response is a **linear function of the coefficients**
 - Later on we'll see how to deal with nonlinear models where the response is not linearly related to the model parameters
- The classic two-sample *t*-test and ANOVA are linear models where the predictors are indicators for the levels of the factors involved
- In the R software, the lm() function can be used to fit regression models
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- For the shrimp example, we have $\hat{y} = 0.839135 0.004498x$, where \hat{y} is the predicted value of y

- Typically, the parameter of primary interest in SLR is the slope, β_1
- The slope measures the average rate of change in y relative to x
- Occasionally, interest also lies in the y-intercept, β_0 , but usually only in cases where x values are collected **near the origin**
- Otherwise, the *y*-intercept may not have any practical meaning
- For the shrimp example, the estimate slope is $\widehat{\beta}_1 = -0.0045$ (how do we interpret this number?)
- In SLR, it is natural to ask whether or not the **slope differs significantly from zero**.
 - If the slope equals zero and the model is correctly specified, then y will not depend on x (i.e., a horizontal regression line)
 - If the relation is quadratic, then one could fit a straight line and get an estimated slope near zero which could be very misleading (always plot your data)

- If $\epsilon \sim N\left(0,\sigma^2\right)$, then $\widehat{eta}_1 \sim N\left(eta,\sigma_{eta_1}^2\right)$ (why?)
- The formulas for $\widehat{SE}\left(\widehat{\beta_0}\right)$ and $\widehat{SE}\left(\widehat{\beta_1}\right)$ are messy, but are provided my most statistical software
 - In R, these are located in the column labeled Std. Error after applying the summary() function (e.g., summary(slr))

```
summary(slr)
##
## Call:
## lm(formula = weight ~ depth)
##
## Residuals:
## Min
                  10 Median
                                     30
                                              Max
## -0.101819 -0.056004 -0.006746 0.049235 0.151772
##
## Coefficients:
##
            Estimate Std. Error t value Pr(>|t|)
## (Intercept) 0.839135 0.081394 10.310 2.77e-06 ***
## depth -0.004498 0.001382 -3.254 0.00993 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.08363 on 9 degrees of freedom
## Multiple R-squared: 0.5406, Adjusted R-squared: 0.4896
## F-statistic: 10.59 on 1 and 9 DF, p-value: 0.009926
```

• In SLR, one can test the following hypothesis:

$$H_0:eta_1=eta_{10}\quad vs.\quad H_1:eta_1
eqeta_{10}$$

- Typically, $\beta_{10} = 0$
- If the null hypothesis is true, then

$$t_{obs} = rac{\widehat{eta}_1 - eta_{10}}{\widehat{SE}\left(\widehat{eta_1}
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will have a t-distribution with n-2 degrees of freedom

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- Similar results hold for the y-intercept β_0

The coefficient of determination

- One of the most important statistics in regression analysis is the coefficient of determination, better known as the \mathbb{R}^2 ("R-squared")
- In the SLR model, \mathbb{R}^2 is just the square of the (Pearson) correlation between x and y
- In MLR (i.e., when we have more than one predictor), a more general definition of \mathbb{R}^2 is required
- Illustration: shrip.R
- The value of \mathbb{R}^2 is always between zero and one and represents the proportion of variability in the response that is explained by the regression model
- **Question:** Interpret R^2 from the SLR model fit to the shrimp data

Assessing the SLR fit

• TBD.