

# Biostatistics

## Week XI – part I

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16 December 2021



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# Conflicting Results

- Researcher A conducts a study comparing the effects an intervention vs. placebo on reducing weight
  - 5 kg reduction among the intervention group ( $p = 0.01$ )
- Researcher B conducts a similar study comparing the effects an intervention vs. placebo on reducing weight
  - 5 kg reduction among the intervention group ( $p = 0.35$ )

# Statistical Power

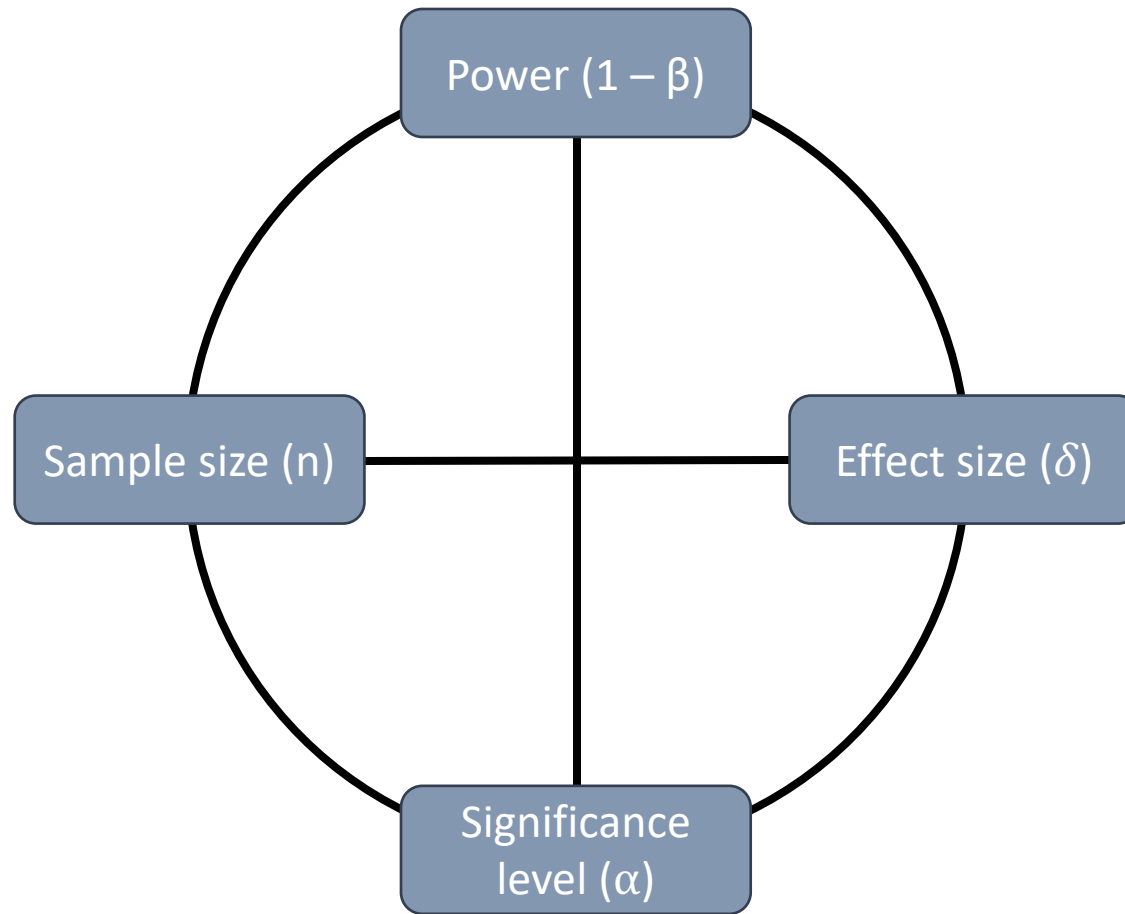
	Decision	
	Fail to reject	Reject
$H_0$		
True	Correct decision	<b>Type I Error</b> $\alpha$
False	<b>Type II Error</b> $\beta$	Correct decision

- **Statistical power** =  $1 - \beta$ 
  - $P(\text{reject } H_0 \mid H_0 \text{ is false})$

# Statistical Power

- Power is affected by:
  - Significance level ( $\alpha$ )
  - Effect size ( $\delta$ )
  - Sample size ( $n$ )

# Power Analysis/Sample Size Calculation



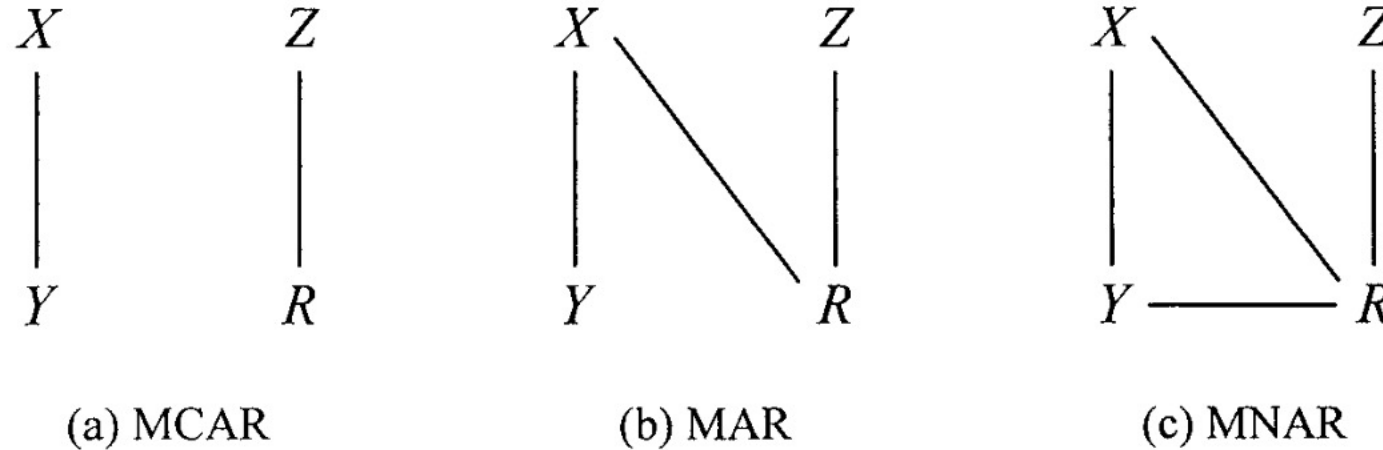
- Given any three, the fourth can be determined

# Default Values

- Power = usually **0.80**, 0.90
- Significance level = usually **0.05**, 0.01, 0.001
- Effect size
  - Literature review
  - Pilot study
  - Cohen's recommendations

# Missing Data

- Missing data, or missing values, occur when no data value is stored for the variable in an observation
- **complications** in handling and analyzing the data
- **bias** resulting from differences between missing and complete data



*Figure 2.* Graphical representations of (a) missing completely at random (MCAR), (b) missing at random (MAR), and (c) missing not at random (MNAR) in a univariate missing-data pattern.  $X$  represents variables that are completely observed,  $Y$  represents a variable that is partly missing,  $Z$  represents the component of the causes of missingness unrelated to  $X$  and  $Y$ , and  $R$  represents the missingness.



# Missing Data - Missing Completely at Random (MCAR)

- The missingness of the data is not associated with either any variable or outcome
- There is **nothing systematic** going on that makes some data more likely to be missing than others
- e.g., Questionnaire lost, blood tube for testing a blood level broken etc.

# Missing Data - Missing at Random (MAR)

- The missingness of the data is **associated with a variable**
- e.g., Supposing men are more likely to tell their weight than women, missingness in weight is MAR

# Missing Data - Missing Not at Random (MNAR)

- The missingness of the data is **related with the outcome**
- e.g., In a depression study, the depression score wasn't calculated for a participant because they committed suicide

# Missing Data

- There are several strategies to cope with missing data:
  - **Try to collect the missing data** (obvious best choice)
  - **Exclude** subjects with any missing data (may reduce the power of the study)
  - **Replace** the missing data with a conservative estimate (e.g., sample mean)
  - **Estimate** the missing data from other data on the same subject (imputation)

# Brief Summary

- Given any three of the following, the fourth can be determined:
  - Power
  - Significance level
  - Effect size
  - Sample size
- Determining sample size prior to starting a study is important
  - Too small of a sample size can under detect the effect of interest in your experiment
  - Too large of a sample size may lead to unnecessary wasting of resources
- There are 3 kinds of missing data:
  - MCAR: nothing systematic
  - MAR: missingness associated with a variable
  - MNAR: missingness related with the outcome

# Biostatistics

## Week XI – part II

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# Regression Analysis

- Regression analysis is used primarily to **model causality** and **provide prediction**
- Predict the values of a **dependent** (response) variable based on values of at least one **independent** (explanatory) variable
- Explain the **effect** of the independent variables on the dependent variable

# Regression Analysis

- Regression can be used to
  - Understand the relationship between variables
  - Predict the value of one (or more) variable(s) based on other variables
- Examples:
  - Quantifying the relative impacts of age, gender, and diet on BMI
  - Predicting whether the treatment will be successful or not based on certain variables



# Regression Analysis

- The variable to be predicted is called the **dependent variable**
  - Also called the **response variable**
- The value of this variable depends on the value of the **independent variable(s)**
  - Also called the **explanatory** or **predictor variable(s)**



# Simple Linear Regression

*E.g., quantifying the impact of age on BMI*

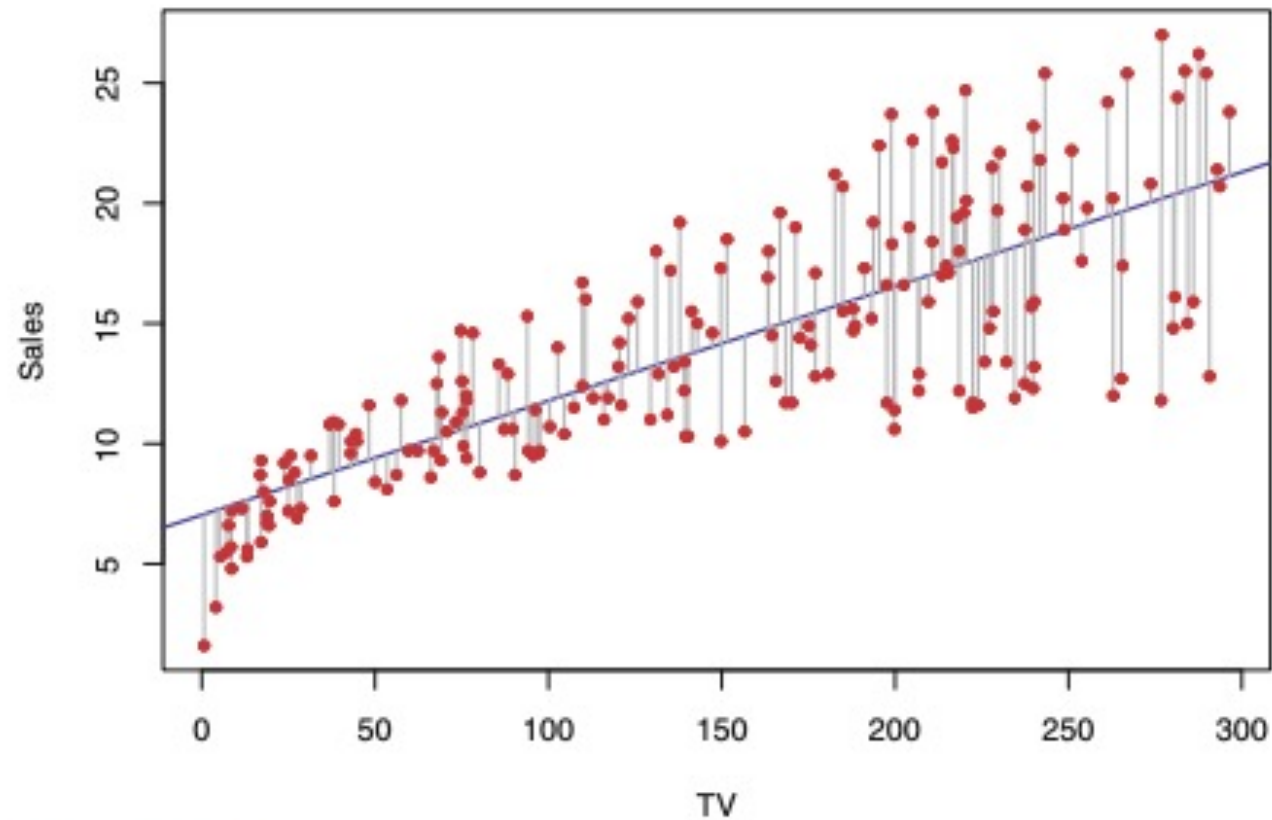
- Linear regression is a method for estimating the **linear relationship** between the dependent and independent variables
- Relationship between variables is described by a linear function

$$Y_i = \beta_0 + \beta_1 X_i + \varepsilon_i \quad \varepsilon \sim N(0, \sigma^2)$$

Diagram illustrating the components of the linear regression equation:

- $Y_i$ : Dependent variable
- $\beta_0$ : Intercept
- $\beta_1$ : slope
- $X_i$ : Independent variable
- $\varepsilon_i$ : residual

- The coefficients are estimated by minimizing the sum of the squared errors/residuals (Least squares)



**FIGURE 3.1.** For the Advertising data, the least squares fit for the regression of sales onto TV is shown. The fit is found by minimizing the sum of squared errors. Each grey line segment represents an error, and the fit makes a compromise by averaging their squares. In this case a linear fit captures the essence of the relationship, although it is somewhat deficient in the left of the plot.

$$\text{Error/residual} = \text{Actual value} - \text{Predicted value}$$

# Estimation of Coefficients

- $\widehat{\beta}_0$  and  $\widehat{\beta}_1$  are estimates by **least squares**
  - minimizing the sum of the squares of the residuals

$$\textit{Residual sum of squares}(RSS) = \sum_{i=1}^n (y_i - \hat{y}_i)^2 = \sum_{i=1}^n (y_i - \widehat{\beta}_0 - \widehat{\beta}_1 x_i)^2$$

$$\widehat{\beta}_0 = \bar{y} - \widehat{\beta}_1 \bar{x}$$

$$\widehat{\beta}_1 = \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\sum_{i=1}^n (x_i - \bar{x})^2}$$

# Estimate of $\sigma$

- Known as the **residual standard error (RSE)**

$$RSE = \sqrt{\frac{RSS}{n - 2}}$$

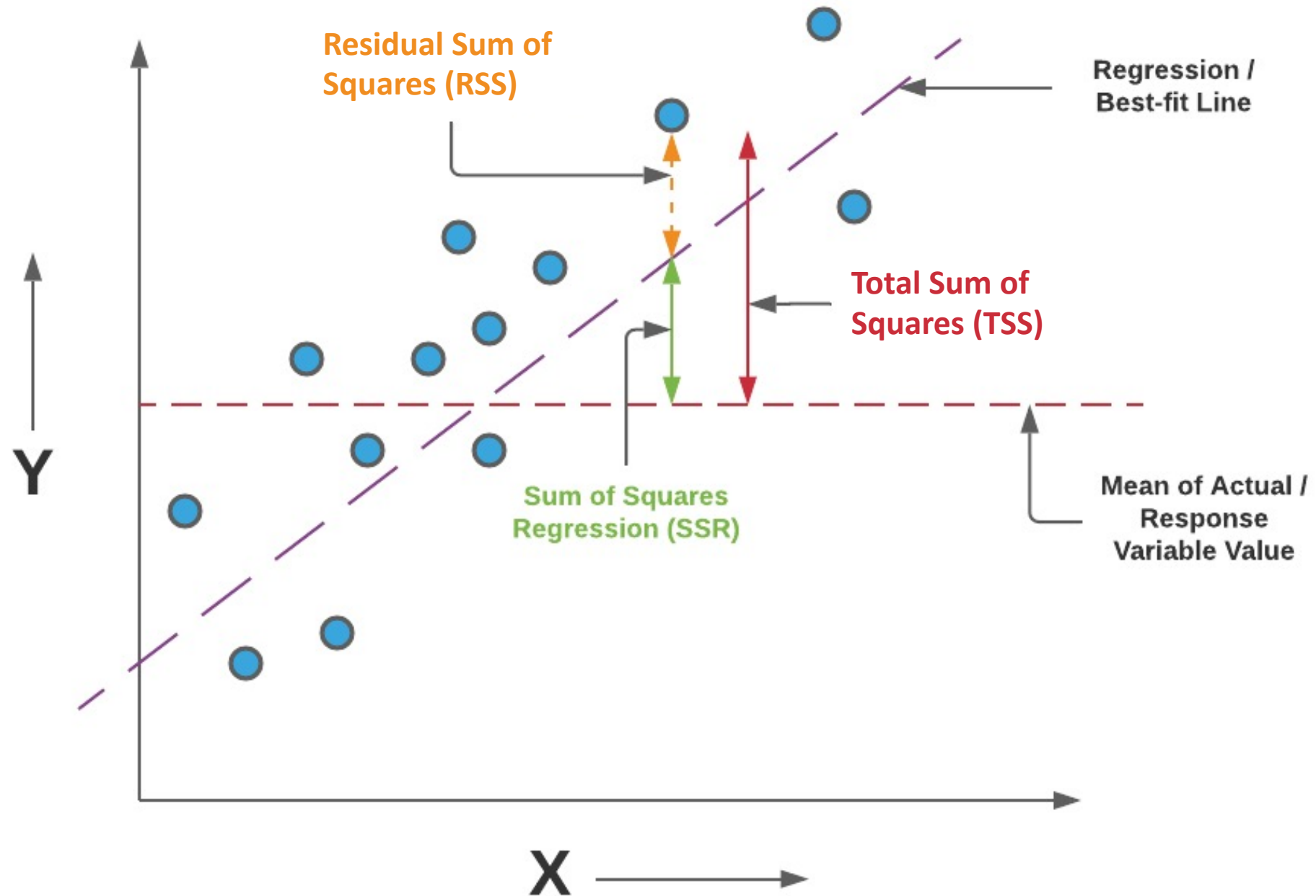
- Using RSE, the standard errors for  $\widehat{\beta}_0$  and  $\widehat{\beta}_1$  are estimated
  - **Significance testing** by t-test  $\left( \frac{\widehat{\beta}_i}{se(\widehat{\beta}_i)} \sim t_{n-1} \right)$

# $R^2$ – Coefficient of Determination

- $R^2$  is measure of **how good is the regression** or best fit line
- It is also termed as **coefficient of determination**

$$R^2 = 1 - \frac{RSS}{TSS}$$
$$TSS = \sum_{i=1}^n (y_i - \bar{y})^2$$

- Measures **the proportion of variation in Y that is explained by the independent variable X** in the regression model
- Greater the value of R-Squared, the better is the regression line as higher is the variance explained by the regression line

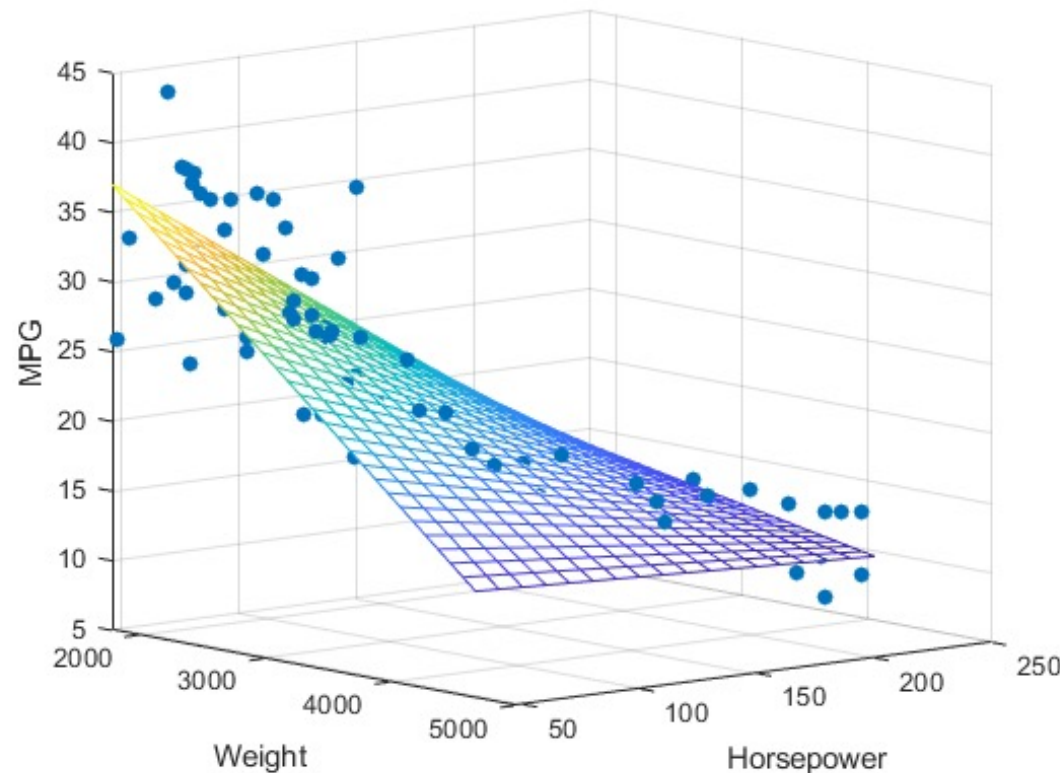


# Multiple Linear Regression

*E.g., quantifying the relative impacts of age, gender, and diet on BMI*

$$Y = \beta_0 + \beta_1 X_1 + \cdots + \beta_p X_p + \varepsilon$$

where  $Y$  is the dependent variable,  $X_1$  to  $X_p$  are  $p$  independent variables,  $\beta_0$  to  $\beta_p$  are the coefficients, and  $\varepsilon$  is the error term





# Multiple Linear Regression (cont.)

- $\widehat{\beta}_0, \widehat{\beta}_1, \dots, \widehat{\beta}_p$  are estimates by **least squares**
- Significance testing by ANOVA
  - $H_0: \widehat{\beta}_0 = \widehat{\beta}_1 = \dots = \widehat{\beta}_p = 0$
- $RSE = \sqrt{\frac{RSS}{n-p-1}}$
- **Adjusted  $R^2$**  is a modified version of  $R^2$  that has been adjusted for the **number of predictors** in the model

$$R_{adj}^2 = \frac{(1 - R^2)(n - 1)}{n - p - 1}$$

*where :*

$R^2$  = *R – squared*

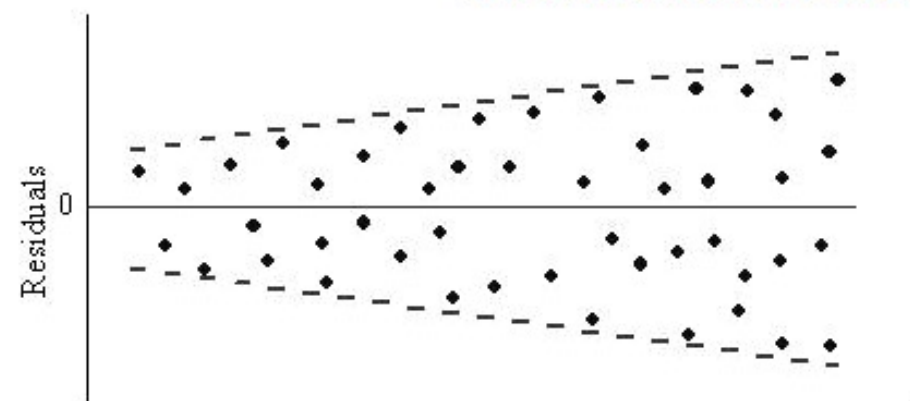
$n$  = *number of samples/rows in the data set*

$p$  = *number of predictors/features*

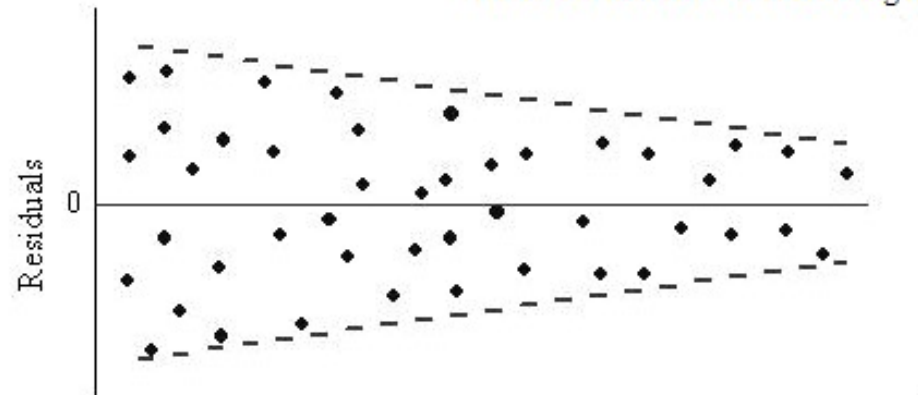
# Linear Regression Assumptions

- There is a **linear relationship** between the independent and dependent variables
- **Normality** – (Q-Q plot / Shapiro-Wilk test)
  - Y values are normally distributed for each X
  - Residuals are normally distributed
- Homoscedasticity (**constant variance**) of the residuals
- **Independence of observations**

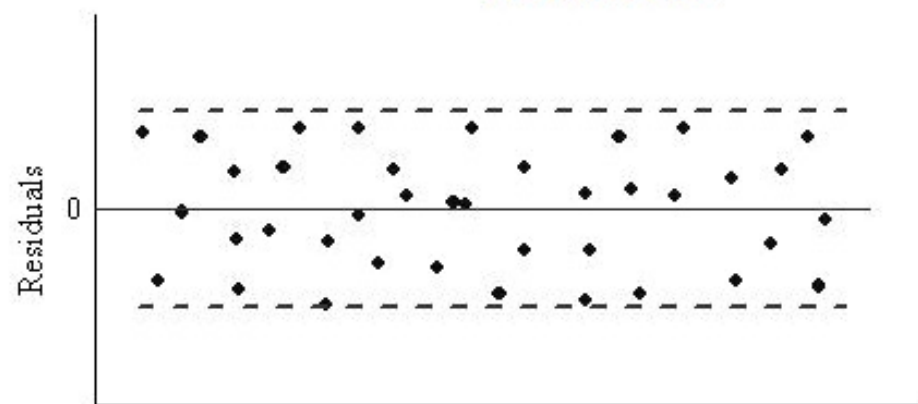
Residuals that show an increasing trend



Residuals that show a decreasing trend

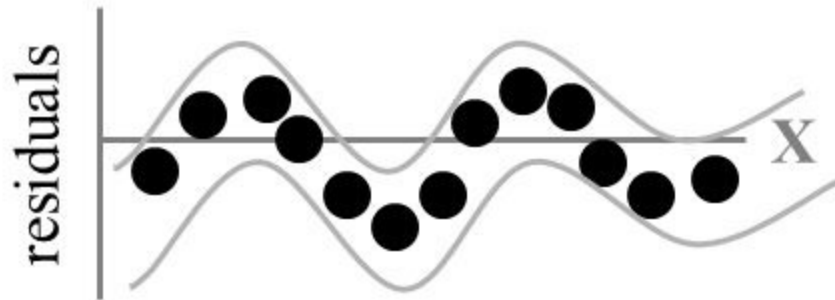
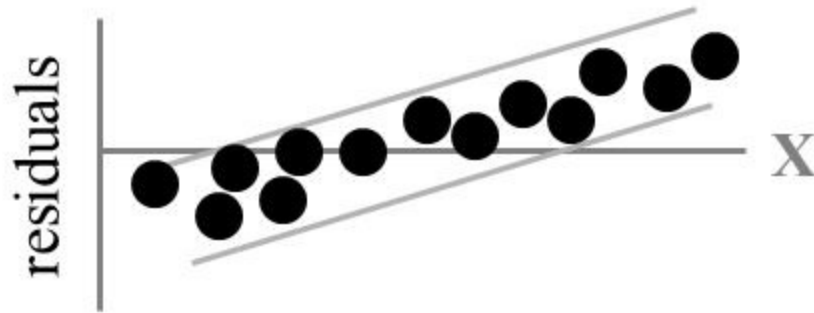


Constant variance

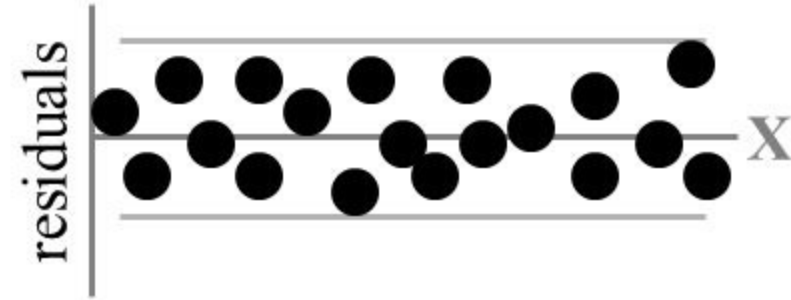


# Residual Analysis for Independence

**Not Independent**



**Independent**



# Linear Regression - Example

## **Prognostic factors for body fat**

- Number of observed individuals:  $n = 241$
- Dependent variable: body fat = percental body fat
- We are interested in the influence of three independent variables:
  - BMI in  $\text{kg/m}^2$
  - Waist circumference (abdomen) in cm.
  - Waist/hip-ratio

# Example - Prognostic factors for body fat - Multiple Linear Regression

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	-60.045	5.365	-11.192	0.000
bmi	0.123	0.236	0.519	0.605
abdomen	0.438	0.105	4.183	0.000
waist_hip_ratio	38.468	10.262	3.749	0.000

$$R^2 = 0.681, R^2_{\text{adj}} = 0.677$$

the proportion of the variation in the dependent variable that is predictable from the independent variable

$$\text{Estimated Body Fat} = -60.045 + 0.123 * \text{bmi} + 0.438 * \text{abdomen} + 38.468 * \text{waist\_hip\_ratio}$$

# Example - Prognostic factors for body fat - Multiple Linear Regression

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	-60.045	5.365	-11.192	0.000
bmi	0.123	0.236	0.519	0.605
abdomen	0.438	0.105	4.183	0.000
waist_hip_ratio	38.468	10.262	3.749	0.000

- For a person with bmi = 0, abdomen = 0, waist\_hip\_ratio = 0, the body fat is estimated to be -60.045 ( $p < 0.001$ )
- (Keeping all other variables the same) with one unit increase in bmi, body fat increases by 0.123 (not significant since  $p > 0.05$ )
- With 95% confidence, it can be stated that with one unit increase in abdomen, body fat increases by 0.438 ( $p < 0.001$ )
- With one unit increase in waist\_hip\_ratio, body fat increases by 38.468 ( $p < 0.001$ )



# Example II

- We'll analyze the prostate cancer dataset
- The main aim of collecting this data set was to inspect the associations between **prostate-specific antigen (PSA)** and **prognostic clinical measurements** in men advanced prostate cancer
- Data were collected on 97 men who were about to undergo radical prostatectomies

*\*PSA was transformed to logPSA for “normalization”*

# Example II – Model 1

$$\log PSA = 1.8 + 0.07 * \textit{vol} + 0.77 * I(\textit{invasion} = 1)$$

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	1.8035	0.1141	15.81	<0.001
vol	0.0725	0.0133	5.43	<0.001
invasion1	0.7755	0.2541	3.05	0.003

Adjusted R-squared: 0.472

## Example II – Model 2

$$\log PSA = 1.67 + 0.1021 * vol + 1.326 * I(invasion = 1) - 0.056 * I(invasion = 1) * vol$$

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	1.6673	0.1289	12.94	<0.001
vol	0.1021	0.0191	5.35	<0.001
invasion1	1.326	0.3588	3.7	<0.001
vol:invasion1	-0.056	0.0262	-2.13	0.0354

Adjusted R-squared: 0.491

For a patient with invasion, there is an additional -0.056 change in PSA when vol changes one unit  
= For a patient with invasion, one unit change in volume results in (0.1021 – 0.056) change in PSA

## Example II – Model 3

$$\log PSA = 1.55 + 0.076 * \mathbf{vol} + 0.45 * I(\mathbf{Gleason} = 7) + 0.9 * I(\mathbf{Gleason} = 8)$$

(compared to **Gleason = 6**)

	Estimate	Std. Error	t value	Pr(> t )
<b>(Intercept)</b>	1.5523	0.1548	10.02	< 2e-16
<b>vol</b>	0.0758	0.0131	5.79	9.30E-08
<b>Gleason7</b>	0.4521	0.1928	2.34	0.0212
<b>Gleason8</b>	0.9043	0.2747	3.29	0.0014

Adjusted R-squared: 0.48

# Brief Summary

- Regression
  - Understand the relationship between variables
  - Predict the value of one variable based on other variables
- Linear regression is a method for estimating the linear relationship between the dependent and independent variables