

# Lesson 1

## The Linear Regression Model





# Simple linear regression

- **The linear model:**
- Can use a linear model to describe the relationship between  $x$  and  $y$ :

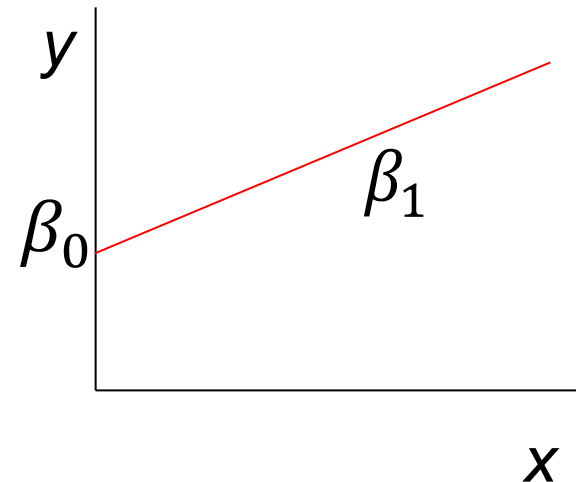
$$y = \beta_0 + \beta_1 x$$

$y_i$  is the response variable

$x_i$  is the predictor variable.

$\beta_0$  is the intercept (the value of  $y$  when  $x = 0$ )

$\beta_1$  is the slope of the regression (the change in  $y$  for every unit of  $x$ )





# Simple linear regression

- The linear **regression** model:
- Can use a linear regression to fit a best linear relationship between  $x$  and  $y$  for some data:

$$y_i = \beta_0 + \beta_1 x_i + \varepsilon_i$$

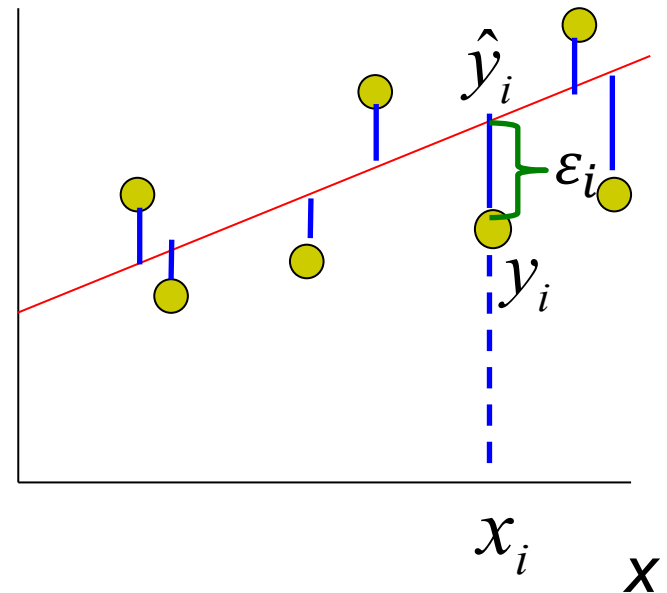
Here:

$i$  are pairs of  $(x, y)$  values measured,

$y_i$  is the response variable

$x_i$  is the predictor variable

$\varepsilon_i$  are the distances between observed  $y_i$  and predicted  $\hat{y}_i$





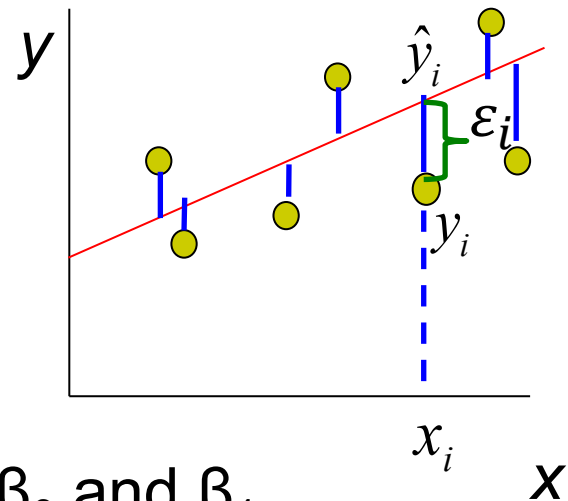
# Regression calculation

- Simple regression uses the method of least squares estimation (OLS):

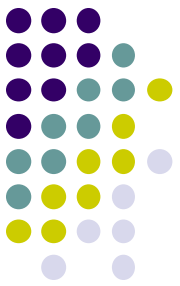
regression fits the best line through the data by minimising the sum of squared errors:

$$\text{Min} \sum_{i=1}^n (y_i - \hat{y}_i)^2 = \text{Min} \sum_{i=1}^n \varepsilon_i^2$$

$$\text{Min} \sum_{i=1}^n (y_i - \hat{\beta}_0 - \hat{\beta}_1 x_i)^2$$



- This requires differentiation w.r.t.  $\beta_0$  and  $\beta_1$ .



# Finding the coefficients, $\beta_i$

- Take partial derivatives wrt to  $\beta_0$  and  $\beta_1$

$$\frac{\partial}{\partial \hat{\beta}_0}: 2 \sum_1^n (y - \hat{\beta}_0 - \hat{\beta}_1 x) = 0 \quad \Leftrightarrow \quad \hat{\beta}_0 n = \sum_1^n y - \hat{\beta}_1 \sum_1^n x$$

$$\frac{\partial}{\partial \hat{\beta}_1}: 2 \sum_1^n x (y - \hat{\beta}_0 - \hat{\beta}_1 x) = 0 \quad \Leftrightarrow \quad \hat{\beta}_0 \sum_1^n x = \sum_1^n xy - \hat{\beta}_1 \sum_1^n x^2$$

- Solve for  $\beta_1$  by equating  $\beta_0$

$$\beta_1 = \frac{\sum_1^n xy - \frac{\sum_1^n y \sum_1^n x}{n}}{\sum_1^n x^2 - \frac{\sum_1^n x \sum_1^n x}{n}}$$

$$n\bar{x} = \sum_1^n x \text{ and } n\bar{y} = \sum_1^n y$$

$$\beta_1 = \frac{\sum_1^n xy - n\bar{x}\bar{y}}{\sum_1^n x^2 - n\bar{x}\bar{x}} = \frac{\sum (x_i - \bar{x})(y_i - \bar{y})}{\sum (x_i - \bar{x})^2}$$



# Finding the coefficients, $\beta_i$

- To calculate the slope ( $\beta_1$ ) we need to calculate:

$$\beta_1 = \frac{S_{xy}}{S_{xx}} = \frac{\sum(x_i - \bar{x})(y_i - \bar{y})}{\sum(x_i - \bar{x})^2}$$

- This can also be written as:

$$\beta_1 = (S_{xx})^{-1}S_{xy}$$

- To calculate intercept ( $\beta_0$ ):

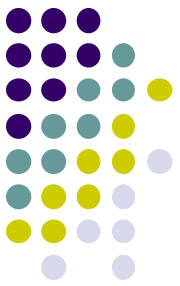
$$\beta_0 = \bar{y} - \beta_1\bar{x}$$

# Estimating the residual variance



- Statistical tests require the residual variance of the model to quantify the uncertainty (s.e.) of  $\beta_i$

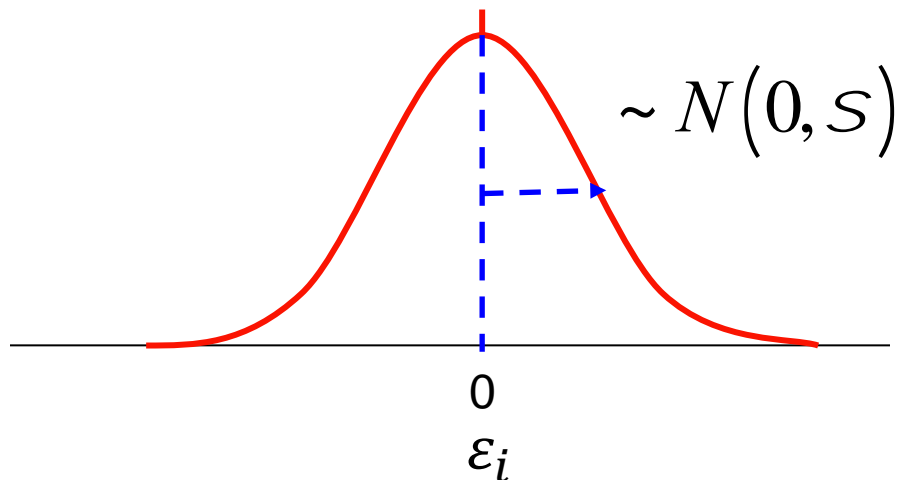
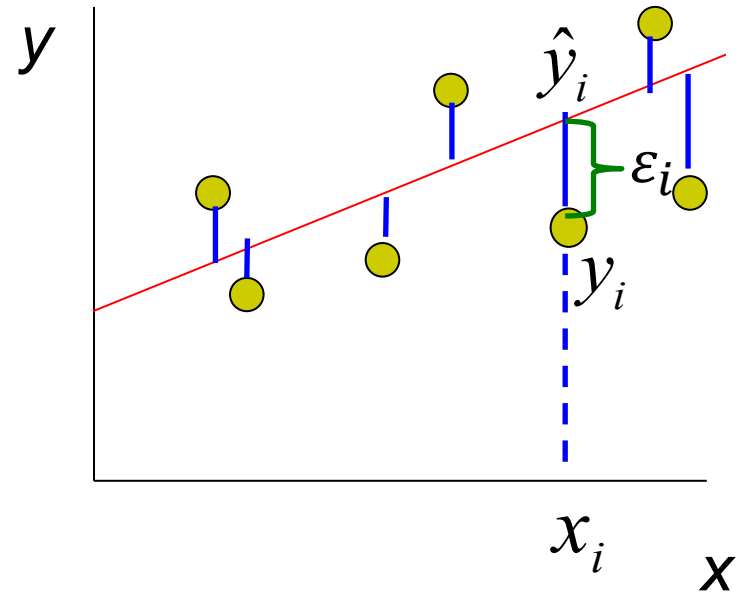
$$\begin{aligned}\hat{\sigma}^2 &= \frac{SS_{res}}{n - 2} = \frac{\sum (y_i - \hat{y}_i)^2}{n - 2} \\ &= \frac{\sum (y_i - (\hat{\beta}_0 + \hat{\beta}_1 x_i))^2}{n - 2}\end{aligned}$$



# The normal distribution

$$y_i = \beta_0 + \beta_1 x_i + \varepsilon_i$$

- The residuals,  $\varepsilon_i$ , of this model are assumed to follow the normal distribution with mean  $\mu$ , and variance  $\sigma^2$ .



$$f(\varepsilon_i) = \frac{1}{\sigma\sqrt{2\pi}} e^{\left(-\frac{1}{2\sigma^2}\varepsilon_i^2\right)}$$



# Hypothesis testing

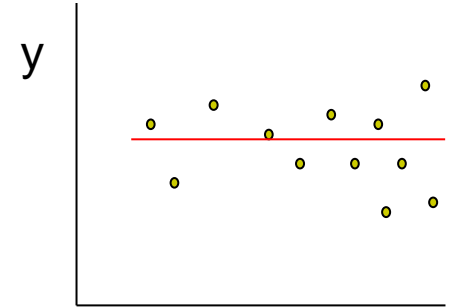


$$y_i = \beta_0 + \beta_1 x_i + \varepsilon_i$$

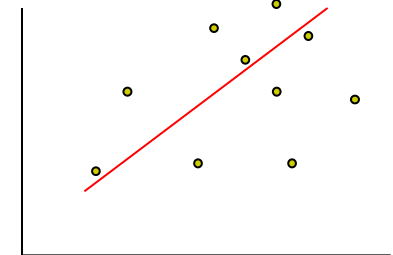
- Main hypothesis: the slope,  $\beta_1 \neq 0$

Coefficients:

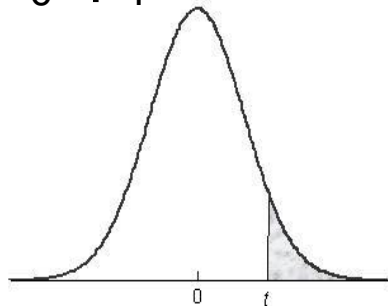
	Estimate	s.e.	t value	Pr(> t )
(Intercept)	22.440	3.4841	6.441	1.55e-05 ***
Temperature	1.015	0.2379	4.268	0.000781 ***



x



- $H_0: \beta_1 = 0$



$$t_{n-2 \text{ df}} = \frac{\widehat{\beta}_1}{se(\widehat{\beta}_1)} = \frac{\widehat{\beta}_1 - 0}{\sqrt{\frac{\sigma^2}{S_{xx}}}}$$



# Interpreting linear models

- Linear models can deal with both continuous and categorical predictor variables simultaneously
- Ordinal predictors = **variates (e.g. age)**
  - (continuous/discrete)
- Categorical predictors = **factors (e.g. sex)**



# Example 1A

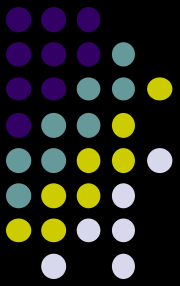
- A researcher measures the maximum swimming speed of 10 brown trout and 10 Canterbury galaxias at a range of temperatures.
- Response ( $Y$ ) = swimming speed
- Predictor Factor = species  $\rightarrow$  dummy variable ( $D$ )
- Predictor Variate ( $X_1$ ) = temperature



# Example 1A



		$X_1$	Y	D
Obs. No.	Species	Temperature (°C)	Speed (cm/s)	Dummy Variable
1	Trout	3	48.1	0
2	Trout	6	51.2	0
3	Trout	11	73.1	0
4	Trout	12	78.1	0
5	Trout	15	81.1	0
6	Trout	21	94.5	0
7	Trout	24	99.0	0
8	Trout	26	115.3	0
9	Trout	28	113.7	0
10	Trout	30	118.7	0
11	Galaxias	4	26.1	1
12	Galaxias	9	33.7	1
13	Galaxias	12	31.4	1
14	Galaxias	14	38.5	1
15	Galaxias	15	36.9	1
16	Galaxias	17	39.1	1
17	Galaxias	18	42.2	1
18	Galaxias	21	43.3	1
19	Galaxias	25	58.9	1
20	Galaxias	28	54.3	1



## Example 1.1

Open the data “fishspeed.csv” in R

(1) Run speed as a function of temperature using `lm()`.  
Write out the model.

(2) Run speed as a function of species using `lm()`.  
Write out the model.

(3) Compare fishspeed of species using `t.test()`  
(please specify: `var.equal=TRUE`)

(4) Compare (2) and (3)



# Example 1A: some output

- Note that the `lm()` output gives you the  $\beta$  estimates with standard errors plus t-tests evaluating significance

Coefficients:

	Estimate	s.e.	t value	Pr(> t )
(Intercept)	17.340	11.967	1.449	0.1667
Temperature	2.979	0.677	4.401	0.0004 ***

- The output also gives you an  $R^2$  value, which is a goodness-of-fit measure (how well model explains data)
- It also gives you the residual standard error =  $\sqrt{\sigma^2}$ , from which the coefficient s.e.'s are derived.



# Matrix calculations

- How regression models actually calculate the parameters for linear models



# Linear models in Matrix notation

- Linear models written in expanded form

$$y_i = \beta_0 + \beta_1 x_i + \varepsilon_i$$

- Or in compound matrix notation

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$$

- Where  $\mathbf{y}$  is the vector of response values, capital  $\mathbf{X}$  is a matrix of predictor values,  $\boldsymbol{\beta}$  is a vector of regression coefficients ( $\beta_0, \beta_1$ ), and  $\boldsymbol{\varepsilon}$  is a vector of residuals.
- Matrix algebra allows us to solve linear regressions simply and powerfully





# Finding the coefficients, $\beta_i$

- Recall for simple linear regression (one predictor):

$$\beta_1 = (S_{xx})^{-1}S_{xy}$$

- This can be solved for p predictors using matrix algebra:

$$\hat{\beta} = (X'X)^{-1}X'y$$



$$\hat{\beta} = (X'X)^{-1}X'y$$



Int.	X1	X2	Y
1	7	560	16.68
1	3	220	11.50
1	3	340	12.03
1	4	80	14.88
1	6	150	13.75
1	7	330	18.11
1	2	110	8.00
1	7	210	17.83
1	30	1460	79.24
1	5	605	21.50
1	16	688	40.33
1	10	215	21.00
1	4	255	13.50
1	6	462	19.75
1	9	448	24.00
1	10	776	29.00
1	6	200	15.35
1	7	132	19.00
1	3	36	9.50
1	17	770	35.10
1	10	140	17.90
1	26	810	52.32
1	9	450	18.75
1	8	635	19.83
1	4	150	10.75

The  $X'X$  matrix is

$$X'X = \begin{bmatrix} 1 & 1 & \dots & 1 \\ 7 & 3 & \dots & 4 \\ 560 & 220 & \dots & 150 \end{bmatrix} \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix} \begin{bmatrix} 7 \\ 3 \\ \vdots \\ 4 \end{bmatrix} \begin{bmatrix} 560 \\ 220 \\ \vdots \\ 150 \end{bmatrix} = \begin{bmatrix} 25 & 219 & 10,232 \\ 219 & 3,055 & 133,899 \\ 10,232 & 133,899 & 6,725,688 \end{bmatrix}$$

and the  $X'y$  vector is

$$X'y = \begin{bmatrix} 1 & 1 & \dots & 1 \\ 7 & 3 & \dots & 4 \\ 560 & 220 & \dots & 150 \end{bmatrix} \begin{bmatrix} 16.68 \\ 11.50 \\ \vdots \\ 10.75 \end{bmatrix} = \begin{bmatrix} 559.60 \\ 7,375.44 \\ 337,072.00 \end{bmatrix}$$

The least-squares estimator of  $\beta$  is

$$\hat{\beta} = (X'X)^{-1}X'y$$

or

$$\begin{bmatrix} \hat{\beta}_0 \\ \hat{\beta}_1 \\ \hat{\beta}_2 \end{bmatrix} = \begin{bmatrix} 25 & 219 & 10,232 \\ 219 & 3,055 & 133,899 \\ 10,232 & 133,899 & 6,725,688 \end{bmatrix}^{-1} \begin{bmatrix} 559.60 \\ 7,375.44 \\ 337,072.00 \end{bmatrix}$$

$$= \begin{bmatrix} 0.11321518 & -0.00444859 & -0.00008367 \\ -0.00444859 & 0.00274378 & -0.00004786 \\ -0.00008367 & -0.00004786 & 0.00000123 \end{bmatrix} \begin{bmatrix} 559.60 \\ 7,375.44 \\ 337,072.00 \end{bmatrix}$$

$$= \begin{bmatrix} 2.34123115 \\ 1.61590712 \\ 0.01438483 \end{bmatrix}$$



# Finding the variance, $\hat{\sigma}^2$

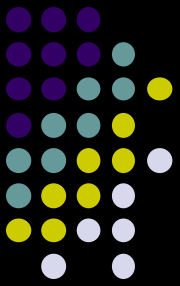
- We can calculate the residual variance for n sampling units and p coefficients  $\hat{\beta}$ :

$$\hat{\sigma}^2 = \frac{SS_{res}}{n - p} = \frac{y'y - \hat{\beta}'X'y}{n - p}$$

- Why?

$$\begin{aligned} SS_{res} &= (y - \hat{\beta}X)'(y - \hat{\beta}X) \\ &= y'y - \hat{\beta}'X'y - y'\hat{\beta}X + \hat{\beta}'X'X\hat{\beta} \\ &= y'y - 2\hat{\beta}'X'y + \hat{\beta}'X'X\hat{\beta} \quad \text{where } X'X\hat{\beta} = X'y \end{aligned}$$

$$\text{Thus: } \hat{\sigma}^2 = \frac{SS_{res}}{n-p} = \frac{y'y - \hat{\beta}'X'y}{n-p}$$



# Example1.2

## Matrix OLS solution of LM

Use the fishspeed data to solve the beta coefficients for:

- (1) Speed ~ Temperature
- (2) Speed ~ Species



# Multi-linear models

- The underlying linear models can be extended from simple linear models

$$y_i = \beta_0 + \beta_1 x_i + \varepsilon_i$$

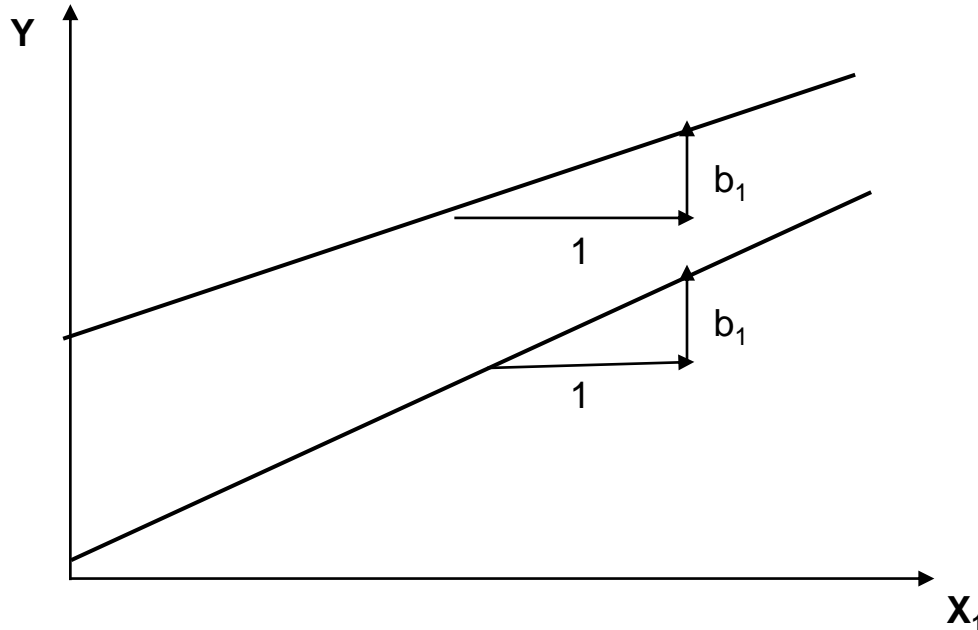
- to models with multiple variables and interactions.

$$y_{ijk} = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_1 x_2 + \varepsilon_{ijk}$$

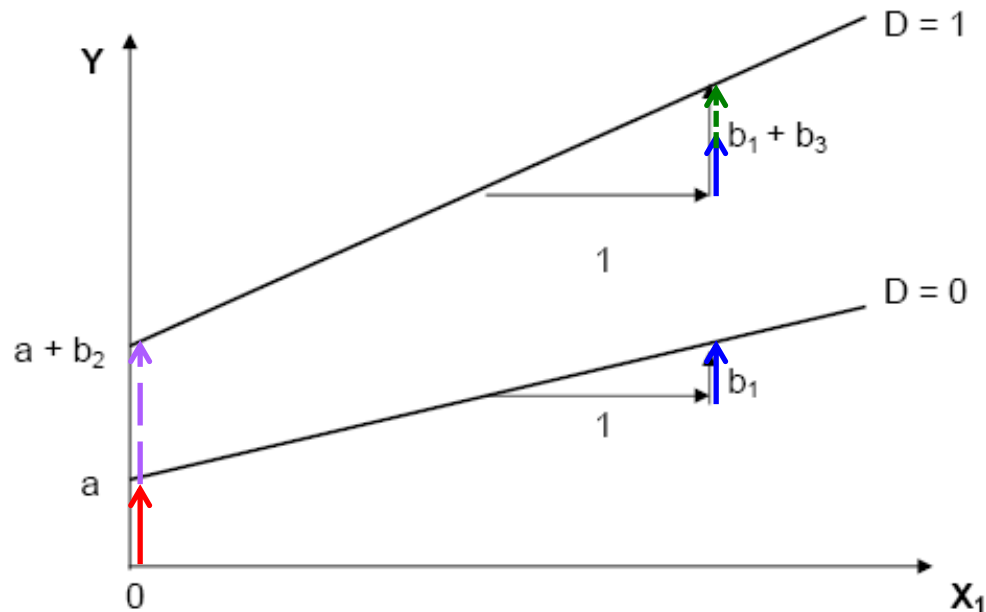
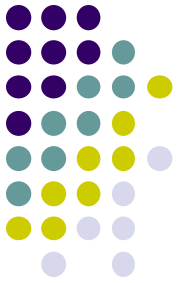
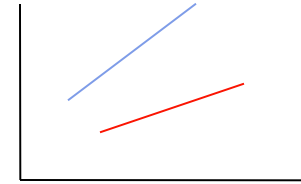
# Interpreting multilinear models



- Regression models with factors and variables have separate regression lines for each factor level



# Modelling interactions



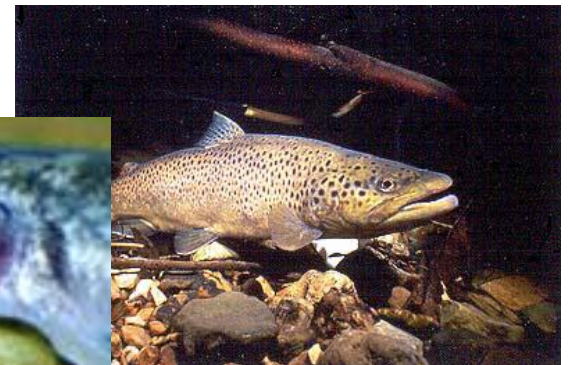
- $a$  = intercept for level 1 ( $D=0$ )
- $b_1$  = regression slope for level 1
- $b_2$  = difference in intercepts of the two lines =  $Y_1 - Y_2$  when  $X = 0$
- $b_3$  = difference in slopes of the two lines



# Fishspeed Example

- A researcher measures the maximum swimming speed of 10 brown trout and 10 Canterbury galaxias at a range of temperatures.
- NOW he wants to know whether the two fish species respond differently to temperature.

=> INTERACTION

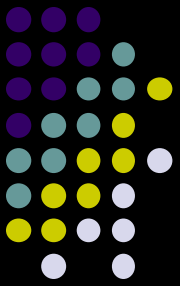




# Recall the data



Obs. No.	Species	$X_1$	Y	D
		Temperature (°C)	Speed (cm/s)	Dummy Variable
1	Trout	3	48.1	0
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# Example: 1.3

## The Fishspeed data

- Formulate the linear model
- Run the regression model in R
- Write out the solution model



# Results: coefficients

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	22.4405	3.4841	6.441	1.55e-05	***
Temperature	1.0152	0.2379	4.268	0.000781	***
SpeciesTrout	18.3559	4.2037	4.367	0.000645	***
SpeciesTrout:Temperature	1.6259	0.2660	6.113	2.68e-05	***

- Model for galaxias:

$$\text{Speed} = 22.440 + (1.015 * \text{Temp})$$

- Model for trout:

$$\text{Speed} = 22.440 + (1.015 * \text{Temp}) + (18.356) + (1.626 * \text{Temp})$$

$$\text{Speed} = 40.796 + (2.641 * \text{Temp})$$

# Results: coefficients

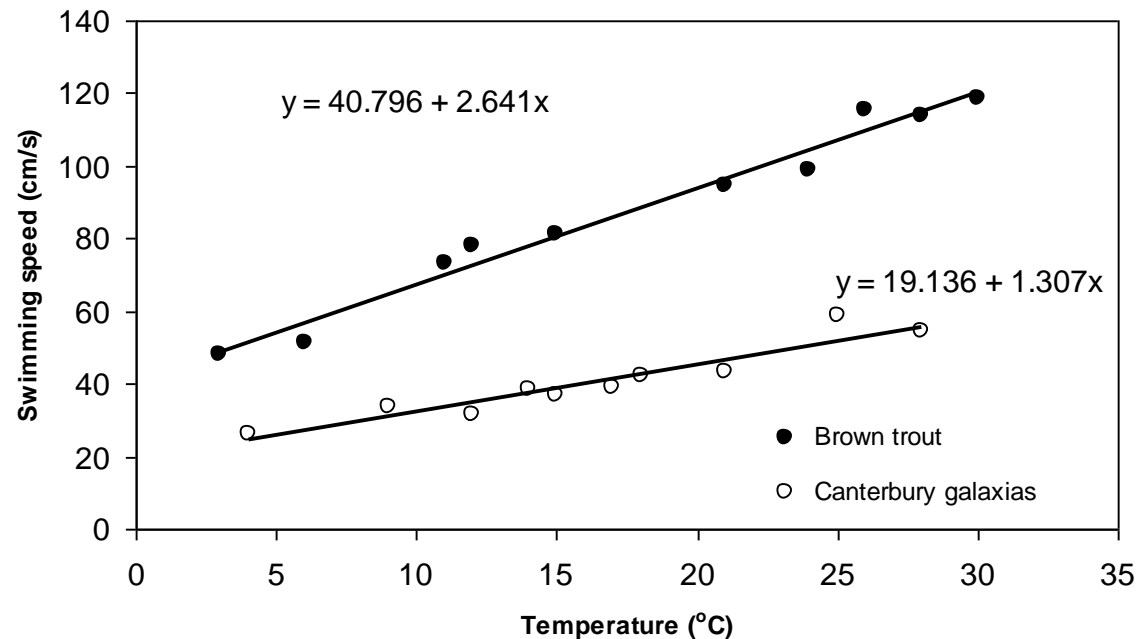


Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
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SpeciesTrout:Temperature	1.6259	0.2660	6.113	2.68e-05	***

- $b_1 > 0$ : swimming speed of galaxias increases with temp.
- $b_2 > 0$ : swimming speed of trout is greater than that of galaxias at 0°C (intercept)
- $b_3 > 0$ : temperature has higher effect on swimming speed of trout than on galaxias.  
i.e. Temperature effect depends on species

# Results



- $b_1 > 0$ : swimming speed of galaxias increases with temp.
- $b_2 > 0$ : swimming speed of trout is greater than that of galaxias at 0°C (intercept)
- $b_3 > 0$ : temperature has higher effect on swimming speed of trout than on galaxias.



# Multilevel Linear Models

- Models with factors are evaluated by considering differences between particular levels and a chosen reference level (the default case)
- When a factor has more than 2 levels, we need to change the default case multiple times to ensure that we cover all pairwise comparisons



## Exercise: 1.4



## Exercise: 1.5





# Principle of marginality

The principle of marginality states that the main effects of a model are marginal to high order terms (such as an interaction). Therefore models should include all lower-order relatives of that higher order term (e.g. the main effects that comprise the interaction).

In other words, the *main effects*, of species and temperature are *marginal* to the species\*temperature interaction.

# Assumptions of linear models



- IMPORTANT!
- Things we need to check (consider) when using linear regression models



# Assumptions of linear models

$$y_{ijk} = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_1 x_2 + \varepsilon_{ijk}$$

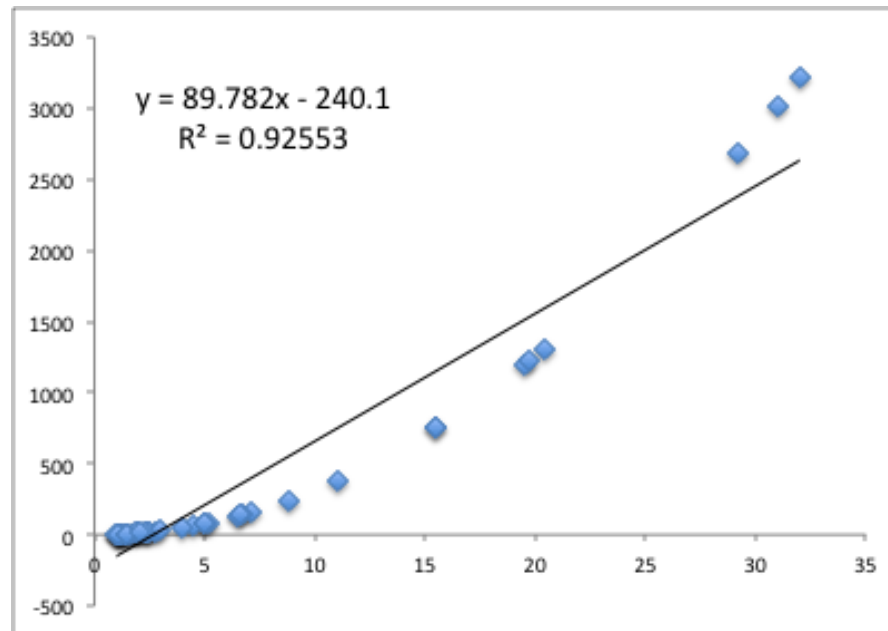
Linear models make many assumptions, including:

1. The model makes biological sense/ physical sense
2. Additivity (terms are added together)
3. Linearity
4. Independence of errors (LATER)
5. Homoscedasticity – equal variance of errors
6. Normality of errors.

# Linearity



- If the relationship is not linear, then the fitted model will not fit the data properly across the domain of values



# Linearity



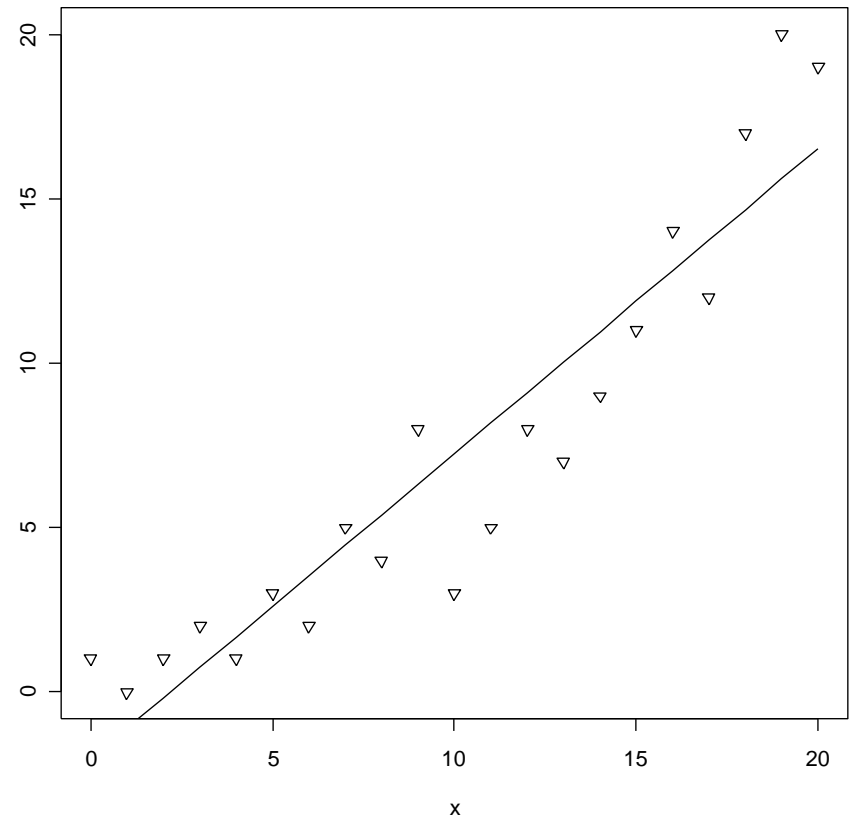
- Linearity:
- How do we check it?

- (1) Plot the raw data:

```
> plot(x, y)
```

```
> model <- lm(y ~ x)
```

```
> lines(x, model$fittedy)
```





# Linearity

- What's the solution?
- Transform the response

$$\ln y = a + b_1 x$$

- Transform the predictors  
e.g. Polynomial regression

$$y = a + b_1 x + b_2 x^2$$

- THINK ABOUT WHAT RELATIONSHIP IS NATURALLY MEANINGFUL (**ASSUMPTION 1**)



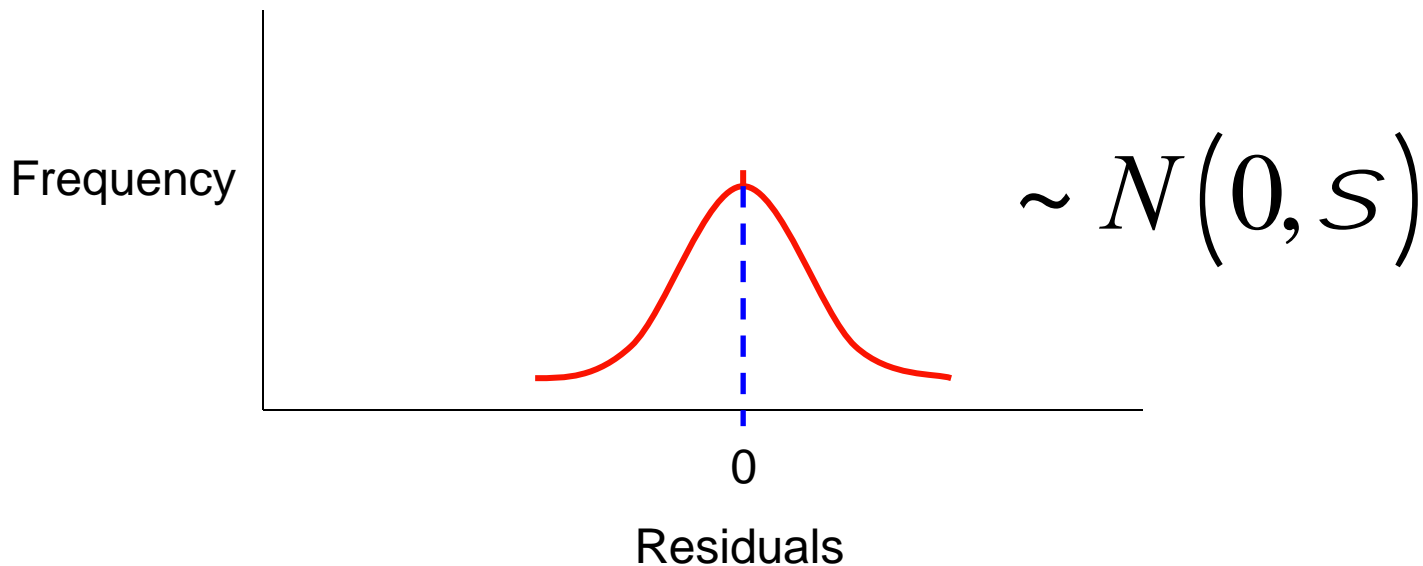
# Distribution of residuals

- The parametric tests are mathematically derived, based on assumed distributions (e.g. normal, F)
- Which means the tests can only be trusted when the data being tested follow the assumed distributions
- e.g. t-tests and F-tests assume residuals are approximately normally distributed

# Normality



.. the residuals of the model should be normally distributed  
Why? Because t-tests and ANOVA assume normal residuals!

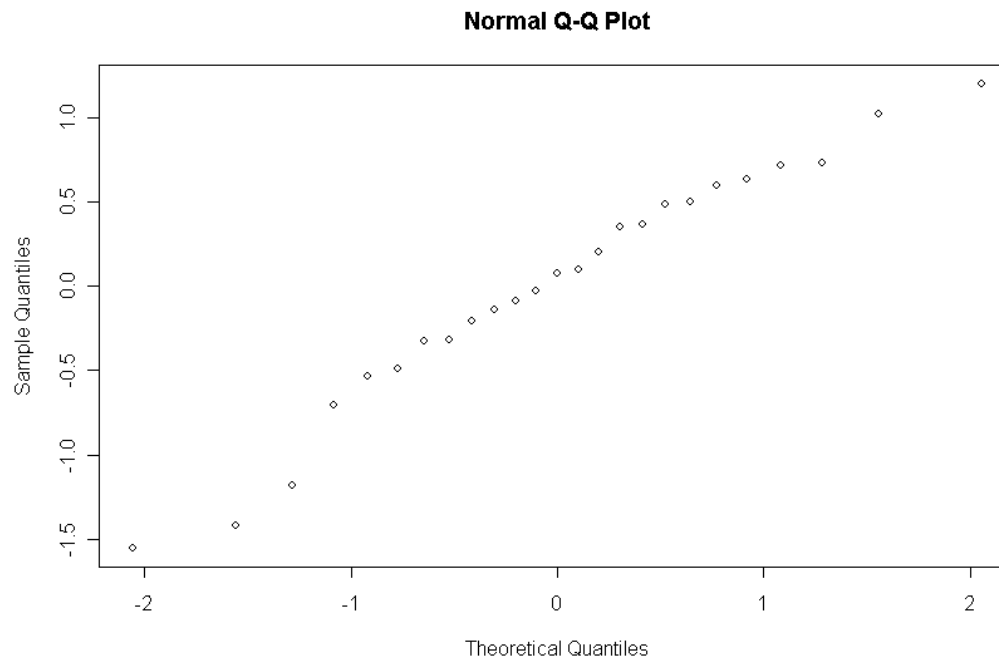




# Normality



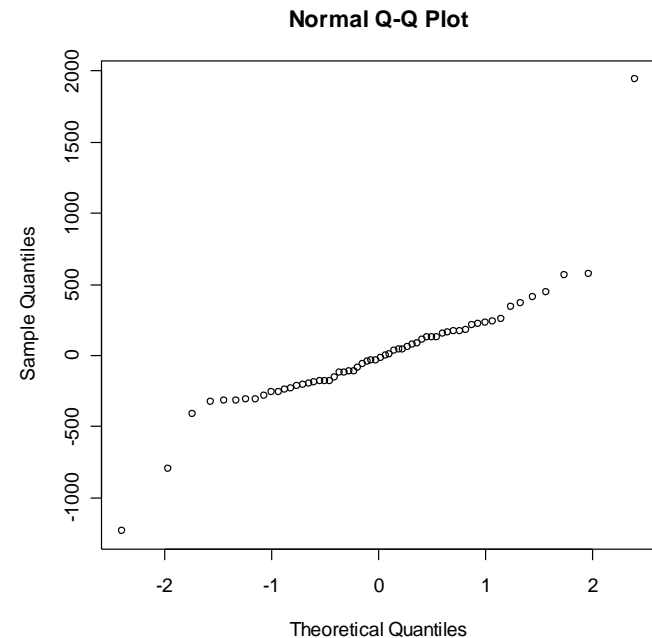
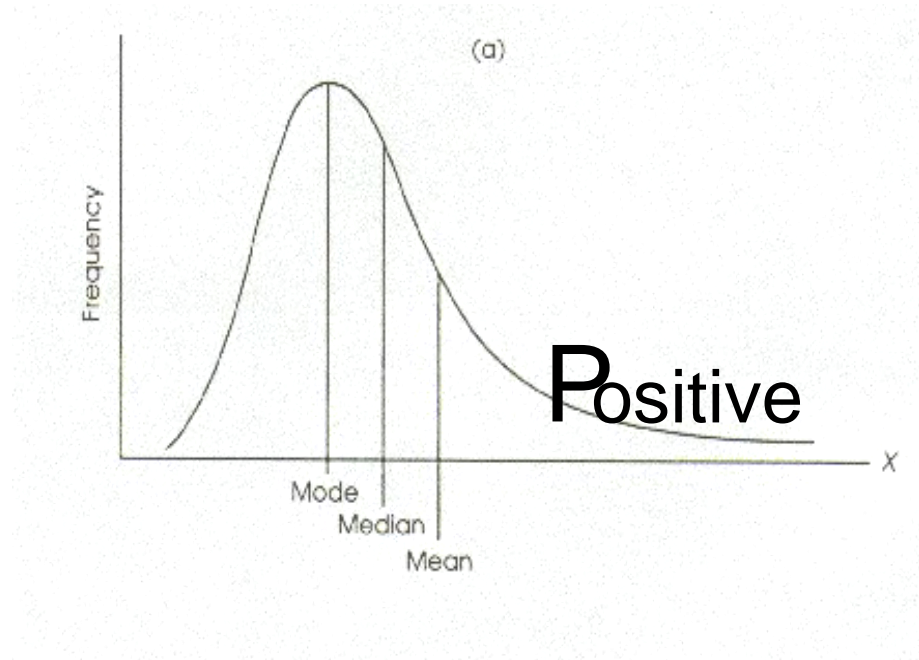
- **How do we check it?**
- Normality of residuals can be checked using a q-q plot





# Distribution of residuals

- Most common problem is positive skew:



- **Solution?** Transform response variable

# Homoscedasticity

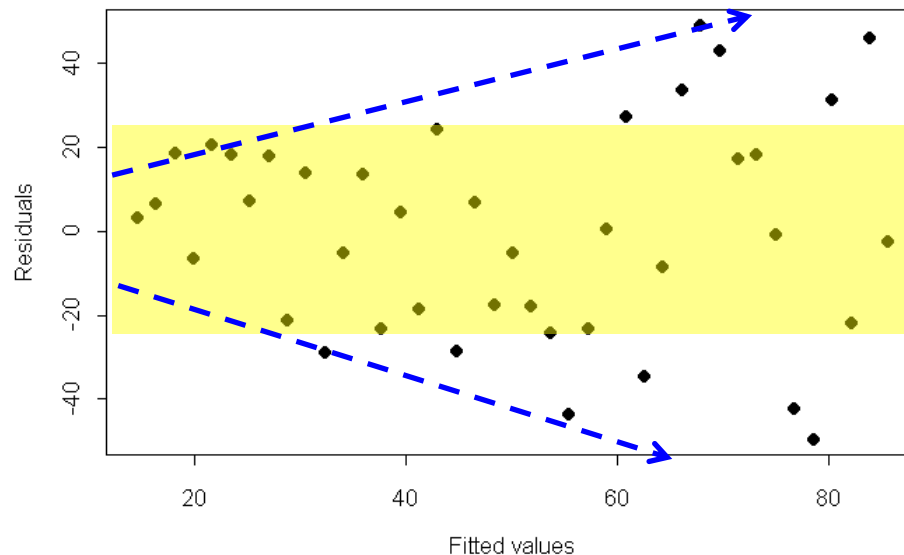


- Homogeneity of variances (Homoscedasticity):
- i.e. the residuals ( $e_i$ 's) should have the same variance across values of the response variable
- If not, then parameter estimates are likely to be unreliable



# Homoscedasticity

- How do we check it?
- 1) Plot residuals ( $y_i - \hat{y}$ ) against fitted values ( $\hat{y}$ )
- Residuals should be evenly spread across the range of fitted values





# Homoscedasticity

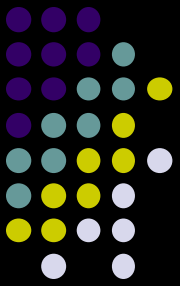
- **What can cause the assumption to be broken?**
- Skew (response or predictors require transformation)
- Outliers
  
- **What's the solution?**
- Transform the response variable or predictor variables



# Checking diagnostics

- This is easily done in R for `lm()` models
- Use the `plot()` function on the derived `lm()` object and it will plot the errors for you.

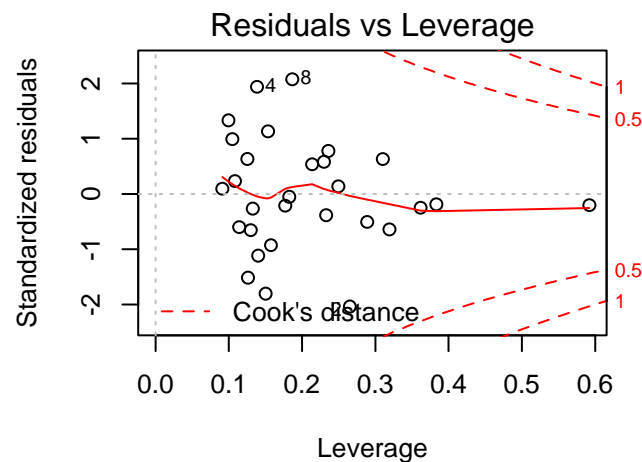
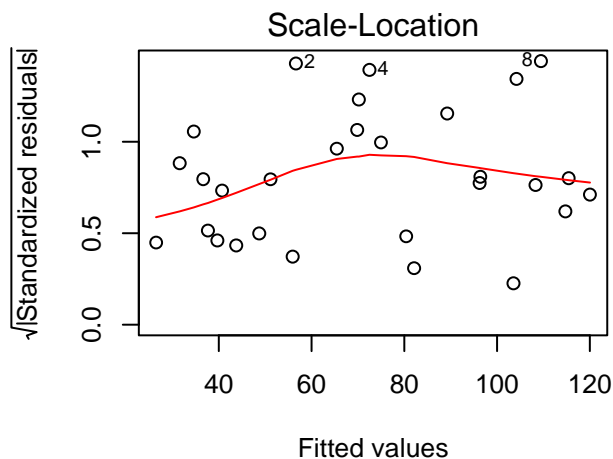
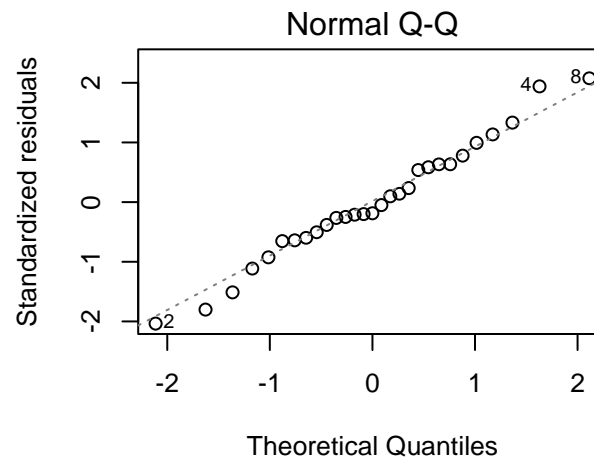
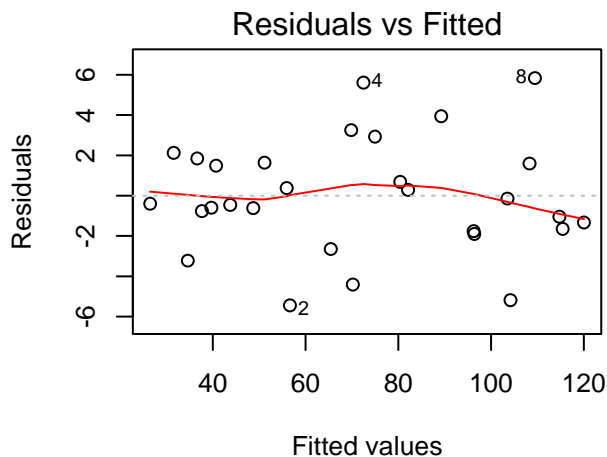
## Example: 1.6



Return to the fishspeed2 problem.

Run diagnostic plots for the model  
you constructed there

# Diagnostics plot for `lm()` in R







**End of Lecture 1**