

# The >eR-Biostat initative Making R based education materials in statistics assessable for all

### Linear regression in R

Based on an online course developed by Marc Lavielle

Inria Saclay (Xpop) & Ecole Polytechnique (CMAP) March, 2017



**ER-BioStat** 



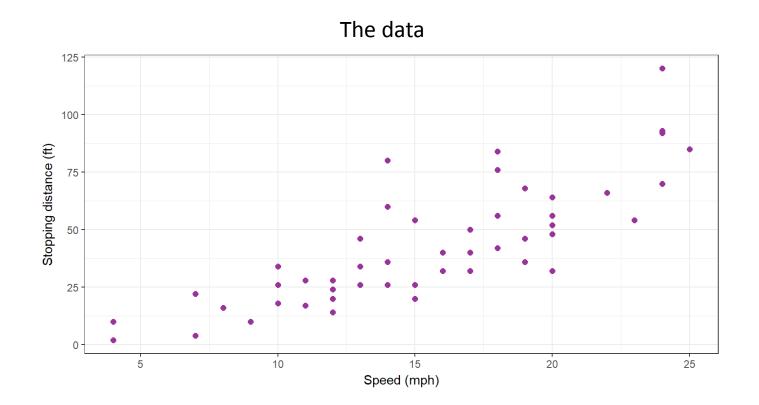


#### Contents

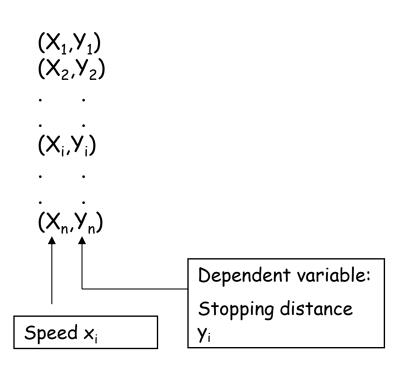
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#### The Car data

- Cars' speed and stopping distance.
- Cars form the 1920s



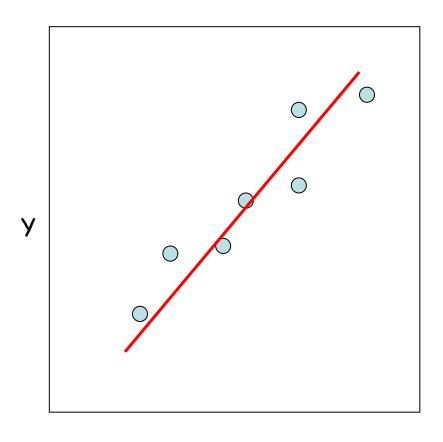
#### Data Structure



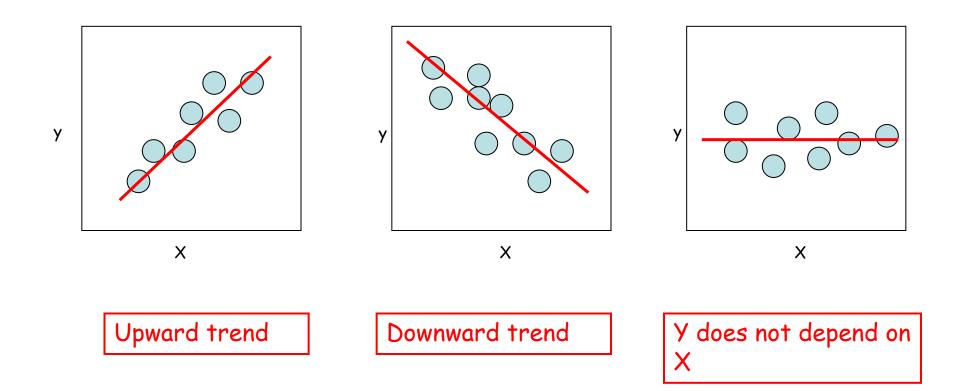
```
The cars data in R
> data(cars)
> head(cars)
  speed dist
2
            10
         22
5
            16
6
            10
Speed x_i
                   Stopping Distance yi
```

## What is a Simple Linear Regression Model?

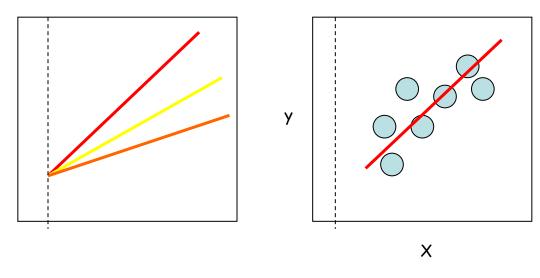
 A regression model is a statistical model which aims to describe the relationship between a predictor (the dose level) and the dependent variable (test score) with a straight line.



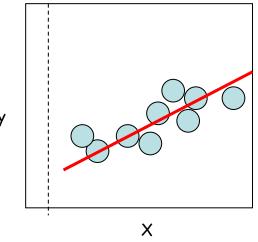
## Properties of Simple Linear Regression Models: Trends



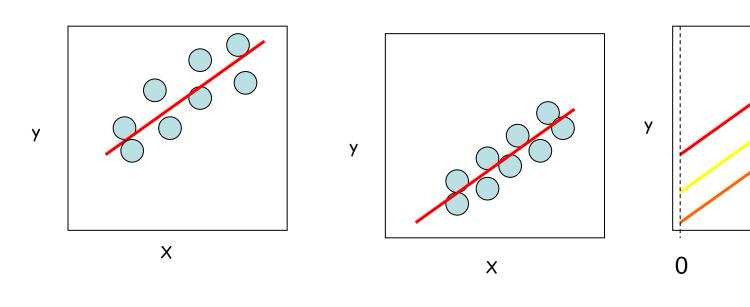
## Properties of Simple Linear Regression Models : Slope



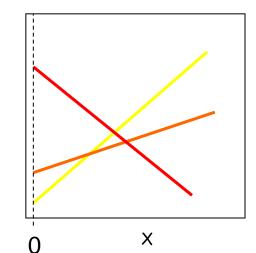
The sploe is the change in the mean of Y for a unit change in X.



## Properties of Simple Linear Regression Models : Intercept



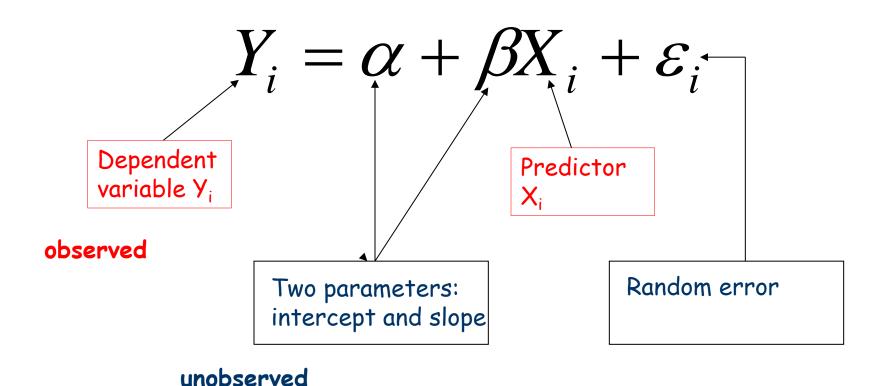
The intercept is the highet of the regression line when x=0.



X

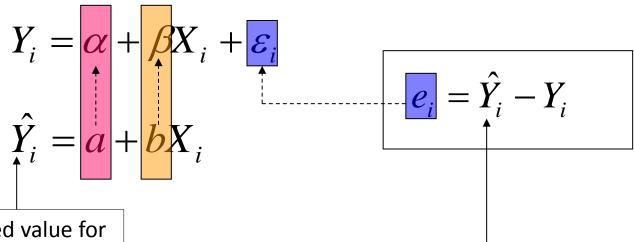
## A Simple Linear Regression Model

 We assume that the relationship between the predictor and the response can be describe with the model:



## Estimation (I)

- We need to estimate the unobserved parameters of the model:
- The estimator for the random error:

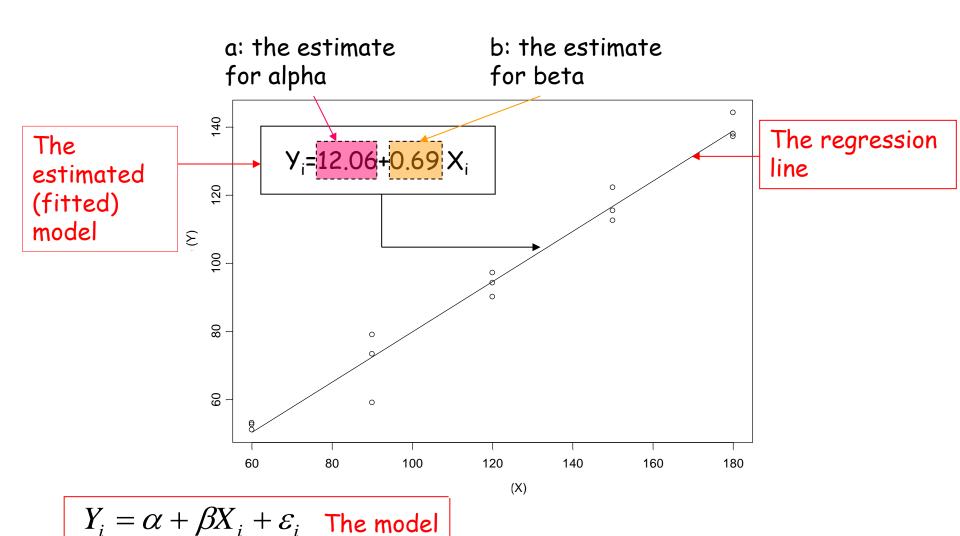


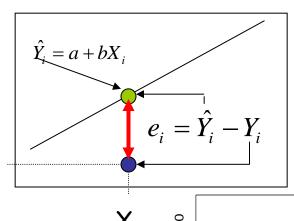
predicted value for the test score

(the estimator for stopping distance)

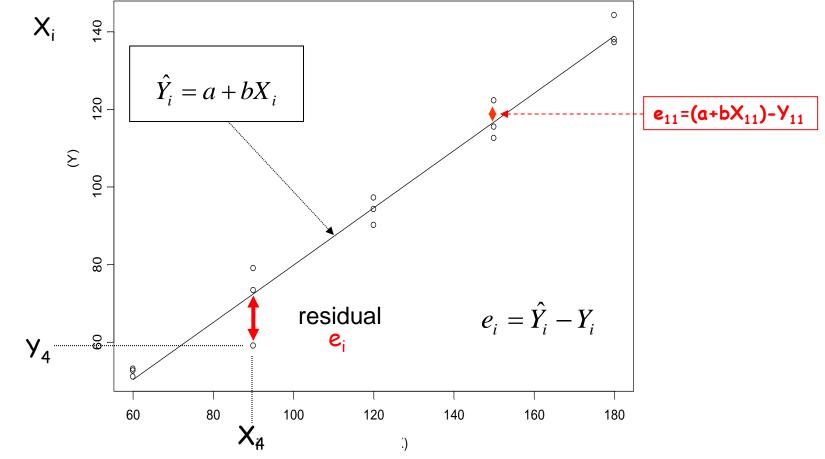
- •a and b are the estimators for alpha and beta
- •e<sub>i</sub> (the residual) is the estimator for the random error

### Regression Model and Data





### The Residuals



## Estimation (II): The Least Squares Criterion

- How to estimate the intercept and slope?
- We want that the fitted model (the line which describes the relationship between Y and X) will be "close" to the data.
- The residual sum of squares = sum (residual) $^2$ .
- The least squares criterion: choose intercept and slope which minimize the residual sum of squares

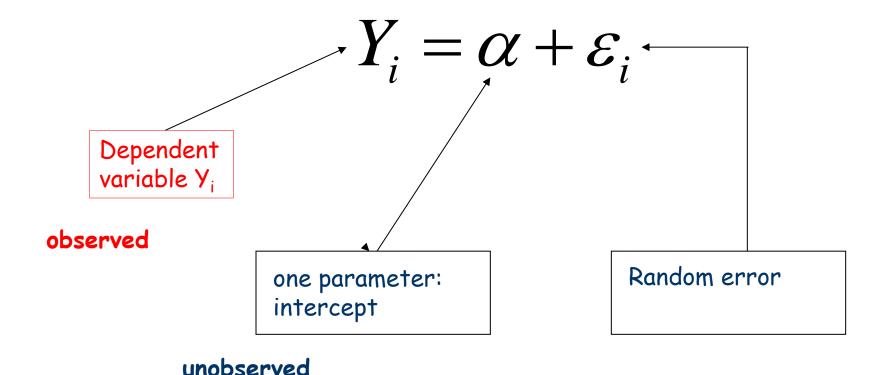
$$RSS = \sum_{i=1}^{n} (\hat{Y}_i - Y_i)^2$$

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## A Simple Linear Regression Model

 We assume that the relationship between the predictor and the response is constant and can be describe with the model:



## Polynomial of degree 0 in R

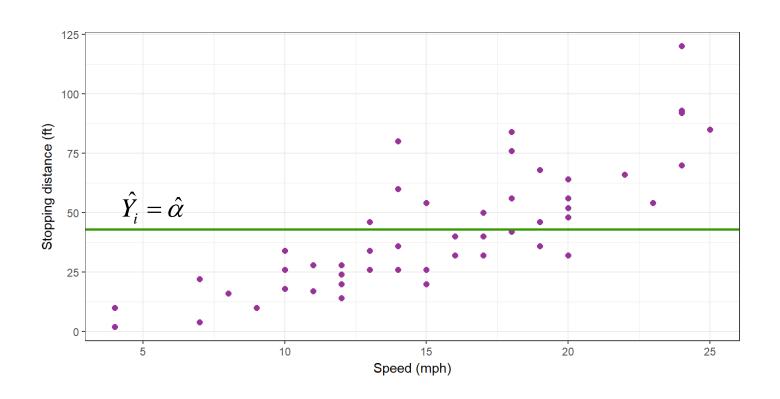
$$\begin{array}{c} \mathbf{attach}(\mathsf{cars}) \\ \mathsf{lm0} <- \ \mathbf{lm}(\mathsf{dist} \ \sim \ \mathbf{1}) \\ Y_i = \alpha + \varepsilon_i \end{array}$$

## R output for the estimated model

```
> summary(lm0)
Call:
lm(formula = dist \sim 1)
Residuals:
  Min 1Q Median 3Q
                             Max
-40.98 -16.98 -6.98 13.02 77.02
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 42.980 < 3.644 11.79 6.38e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '
```

Residual standard error: 25.77 on 49 degrees of freedom

### Data and estimated model

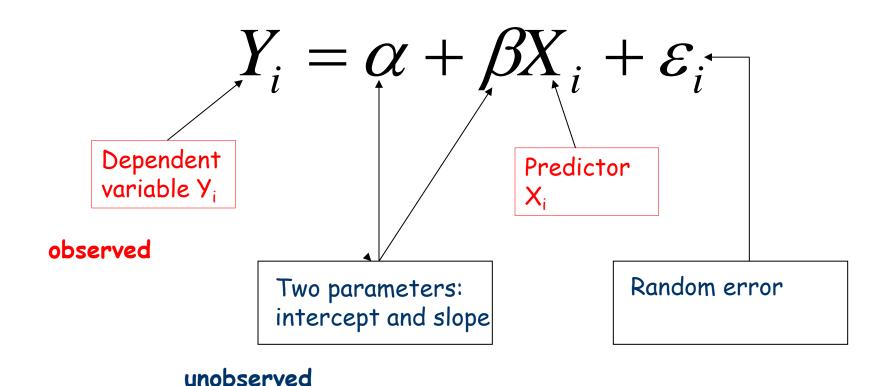


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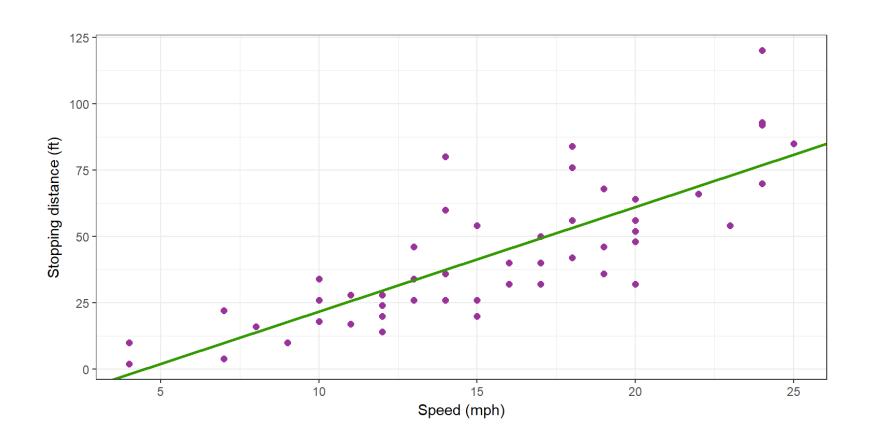
## A Simple Linear Regression Model

 We assume that the relationship between the predictor and the response can be describe with the model:



```
lm1 <- lm(dist ~ speed, data=cars) coef(lm1)</pre>
coef(lm1)
## (Intercept)
                   speed
## -17.579095
                   3.932409
```

## Data and estimated model



#### Estimated model in R

```
> summary(lm1)
Call:
lm(formula = dist ~ speed, data = cars)
Residuals:
   Min 1Q Median 3Q
                                 Max
-29.069 -9.525 -2.272 9.215 43.201
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) -17.5791 6.7584 -2.601 0.0123 *
      3.9324 0.4155 9.464 1.49e-12 ***
speed
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
Residual standard error: 15.38 on 48 degrees of freedom
Multiple R-squared: 0.6511, Adjusted R-squared: 0.6438
F-statistic: 89.57 on 1 and 48 DF, p-value: 1.49e-12
```

## Confidence interval for the parameters

**Summary** 

## **Technical Details (Estimation)**

• A simple linear regression model has the form:

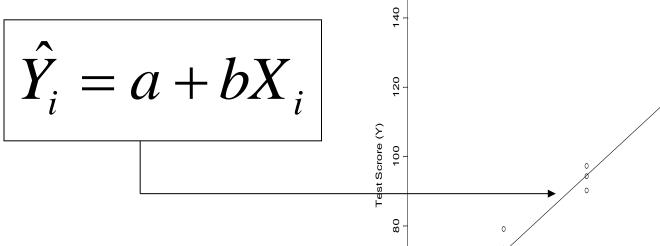
$$Y_i = \alpha + \beta X_i + \varepsilon_i$$

- • $\alpha$  and  $\beta$  are the parameters in the model and  $\mathcal E$  is the random error.
- We can estimate  $\alpha$  and  $\beta$  by minimizing the residual sum of squares

$$RSS = \sum_{i=1}^{n} (\alpha - \beta X_i - Y_i)^2$$

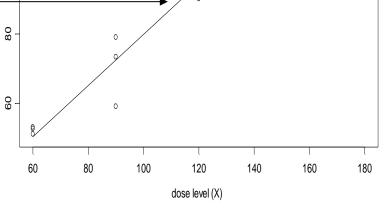
## Technical Details (Estimation)

The estimated model



• The residual

$$e_i = \hat{Y}_i - Y_i$$



## Technical Details (Estimation)

We assume that the relationship between  $Y_i$  and  $X_i$  can be described with a statistical model  $Y_i = \alpha + \beta X_i + \varepsilon_i$ 

We assume that the random error ${\mathcal E}$ is normally distributed.	$\varepsilon \sim N(0, \sigma^2)$
The mean of ${\mathcal E}$ is equal to zero	$E(\varepsilon_i) = 0$
The conditional mean of $Y_i$ (given the value of $X_i$ )	$E(Y_i \mid X_i) = \alpha + \beta X_i$
The estimator for the conditional mean of $Y_i$ (the fitted model=the regression line)	$\hat{E}(Y_i \mid X_i) = a + bX_i = \hat{Y}_i$
The residual: the estimator for ${\mathcal E}$	$e_i = \hat{Y}_i - Y_i$
Least square criterion: choose a and b that minimize the residuals sum of squares	$RSS = \sum_{i=1}^{n} (\alpha - \beta X_i - Y_i)^2$

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## Simple regression model and it's assumtions

We focus on model diagnostic. We consider the following linear regression model

$$Y_i = \alpha + \beta \times X_i + \varepsilon_i$$

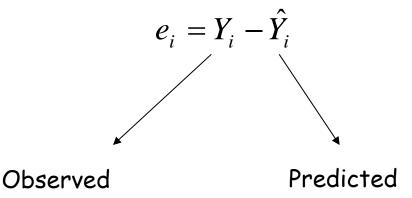
The random error is **assumed** to be normal distributed:

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

We also assume that the variance is constant, i.e., the variance of  $\varepsilon_1, \varepsilon_2, ...., \varepsilon_n$  is equal.

## How to check the model assumptions? (1)

• The random error,  $\varepsilon_i$ , is unknown but we can estimate  $\varepsilon_i$  with the residuals



- The residuals can be used in order to check the model asaumptions.
- We focus on two things:
  - 1) the distribution of e<sub>i</sub>
  - 2) the variability of e<sub>i</sub>

## How to check the model assumptions? (2)

 We assume that the mean of Y<sub>i</sub> is linear with respect to X:

$$E(Y)_i = \alpha + \beta \times X_i$$

 Once again, the residuals can be used in order to check the linearity asaumption.

• This is true only if

$$E(\varepsilon_i) = 0$$

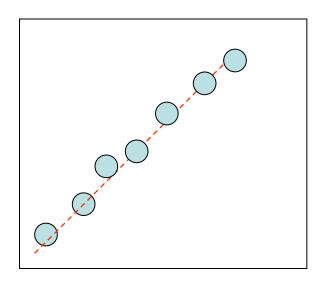
residuals O The residuals in this plot should be around zero and do not show any systematic pattern.

predictor

## Assumption 1: The distribution of e<sub>i</sub>

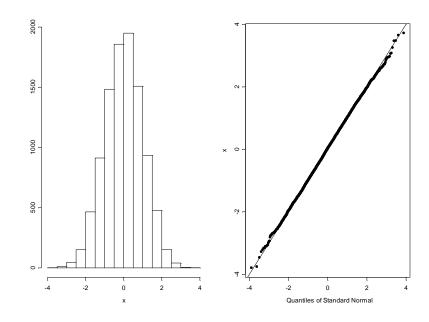
- The distribution of  $e_i$  is expected to be normal with mean zero and variance  $\sigma^2$
- qq-normal plot (or normal probability plot) is a graphical tool that can be used in order to asses the normality assumption.

If the normalty assumption holds we expect qq-normal plot will be a srtight line.



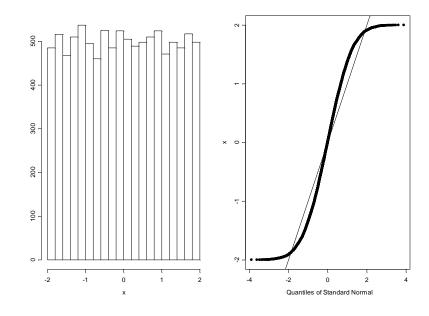
## Example of qqnormal plot form N(0,1)

- Sample of 10000
   observations from N(0,1)
- The qqnormal plot is a stright line.
- If the random error  $\varepsilon_i$  is normal distributed, the qqnormal plot of the residuals should be a stright line.



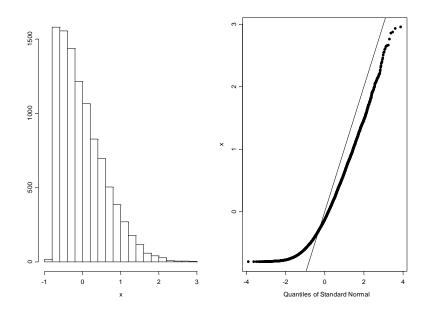
## Example of qqnormal polt from havy tailed distribution

- Sample of 10000
   observations from
   U(-2,2).
- S shape of the qqnormal plot.
- This is an example of a symatric distribution with more observations (relativly to the normal distribution) at the tails.



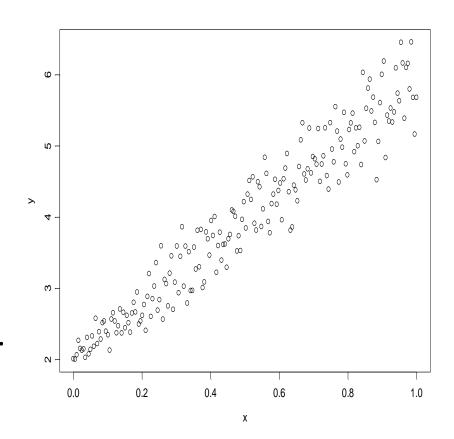
## Example of qqnormal polt from skwed distribution

- Sample of 10000
   observations from a skewed
   distribution.
- The distribution is sewed to the right and the points in the qqplot are not follow the stright line.



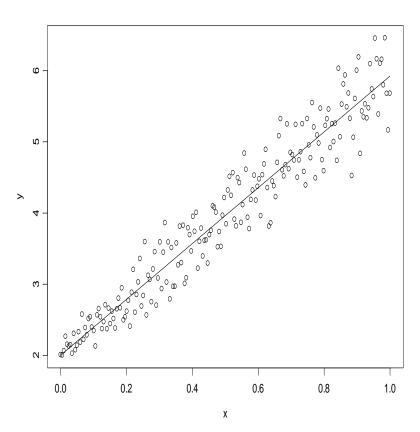
# Assumption 2: Constant variance

- This is an example of a dataset in which the variance is not constant.
- The variance increases when the value of X increases.
- However, there is a linear relatioship between the predictor and the response.



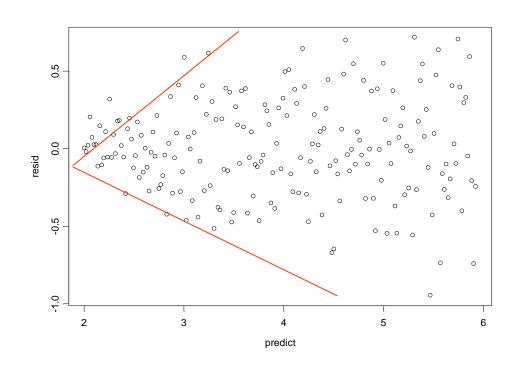
## The data and the resrssion line

 The model seems to fit the data well in the sense that is captures that structure of the mean.

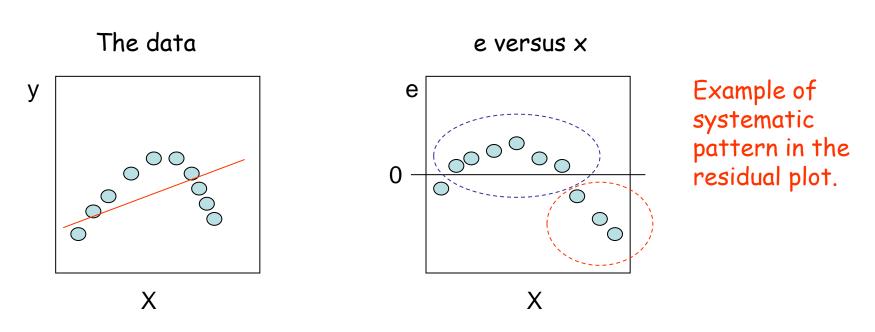


# Residuals plot: Residuals versus the predicted values

 In this plot we can see clearly a pattern. As the predicted values increase the variability among the residuals increase (a "megaphone" shape).

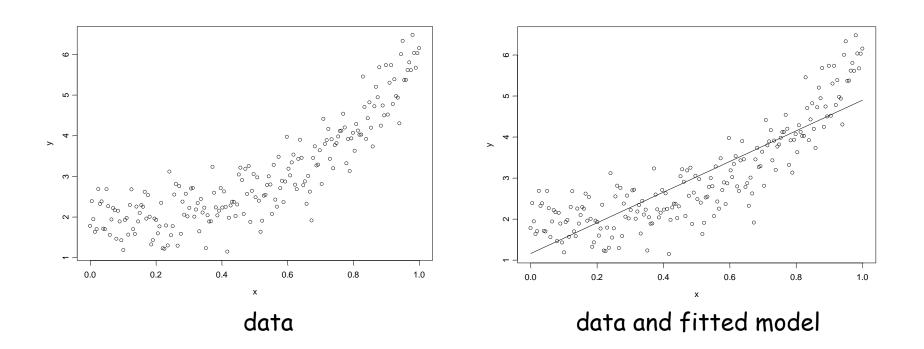


# **Assumptio 3: Linearity**



The scatterplot of the data reveals that the association between the response and the predictor is not linear. The residuals plot (in the right) reveals a clear pattern among the residuals which depends on the value of X.

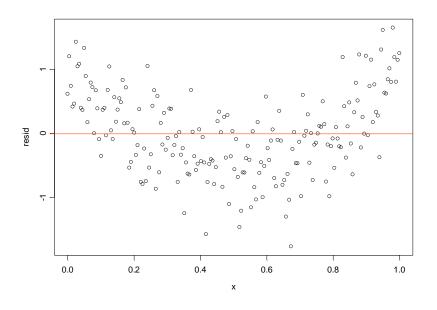
# Systematic patterns



The model underestimates the value of Y when the value of X is relatively small or large.

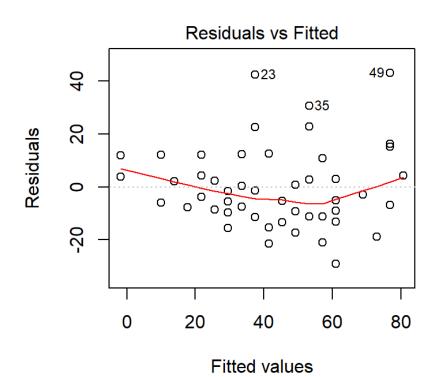
# Linearity: Residuals plot

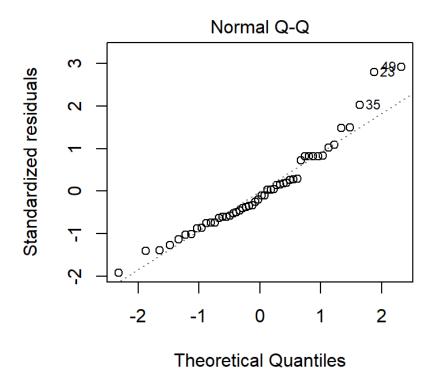
This plot reveals clear systematic pattern among the residuals: the residuals are positive for small and large value of X and negative in the middle. This means that there is structure in the data that the linear regression model did not capture.



# Diagnostic plots in R

```
par(mfrow = c(1, 2))
plot(lm1, which=c(1,2))
```





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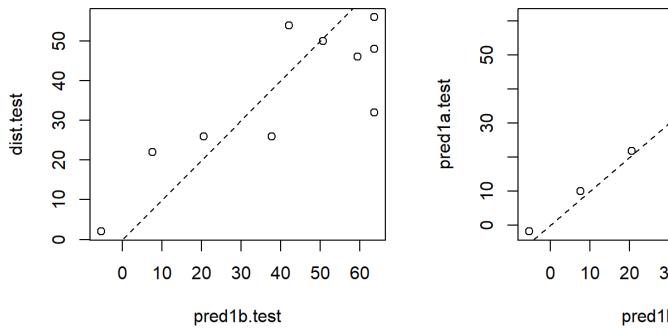
# Cross-validation for prediction

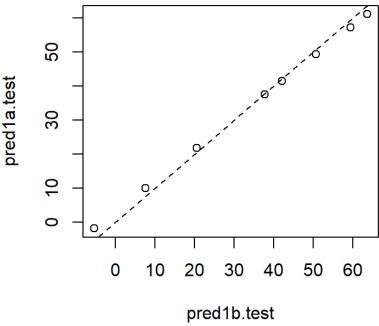
 A common practice is to split the dataset into a 80:20 sample (training:test), then, build the model on the 80% sample and then use the model thus built to predict the reponse variable on test data.

#### Predicted values for the test data

```
> set.seed(100)
> n <- nrow(cars)</pre>
> i.training <- sort(sample(n,round(n*0.8)))</pre>
> cars.training <- cars[i.training,]</pre>
> cars.test <- cars[-i.training,]</pre>
> pred1a.test <- predict((1m1, )newdata=cars.test)</pre>
>
> lm1.training <- lm(dist ~ speed, data=cars.training)</pre>
> pred1b.test <- predict(Im1.training, newdata=cars.test)</pre>
> data.frame(cars.test, pred1a.test, pred1b.test)
   speed dist pred1a.test pred1b.test
                           -5.392776
       4
                -1.849460
           22 9.947766 7.555787
                21.744993 20.504349
      10
           26
           26
                37.474628 37.769100
20
      14
26
      15
           54 41.407036
                           42.085287
31
      17
           50
                49.271854
                           50.717663
37
                57.136672
                           59.350038
      19
           46
39
      20
           32
                61.069080
                           63.666225
      20
                61.069080
                           63.666225
40
           48
42
           56
                61.069080
                            63.666225
      20
```

# Predicted values for two test dataset and the observed data



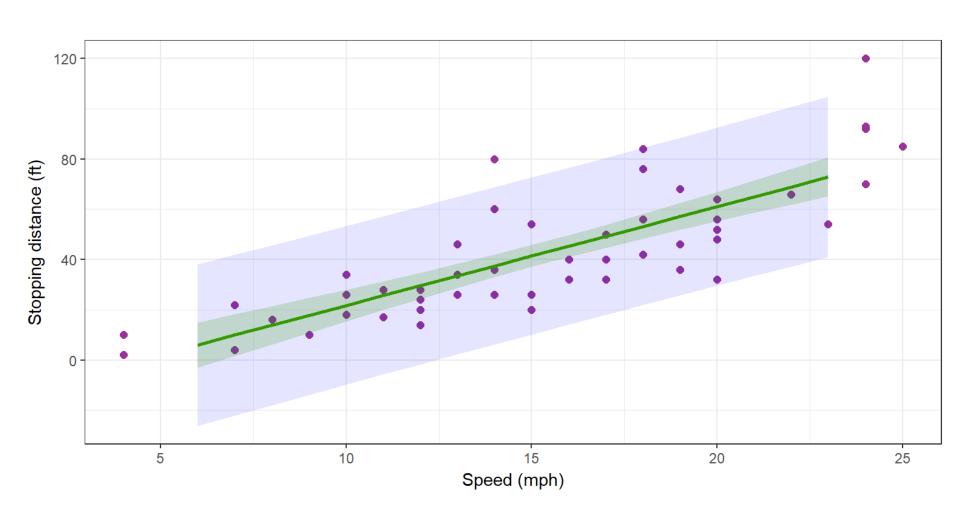


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# Confidence intervals for regression line

# A prediction interval for a new measured distance

# Graphical display for the two confidence intervals



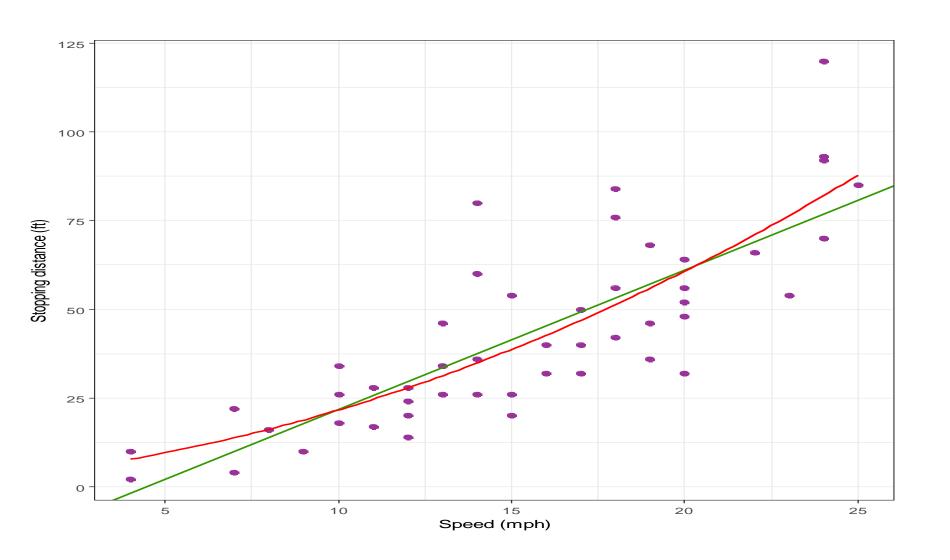
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# Polynomial of degree 2

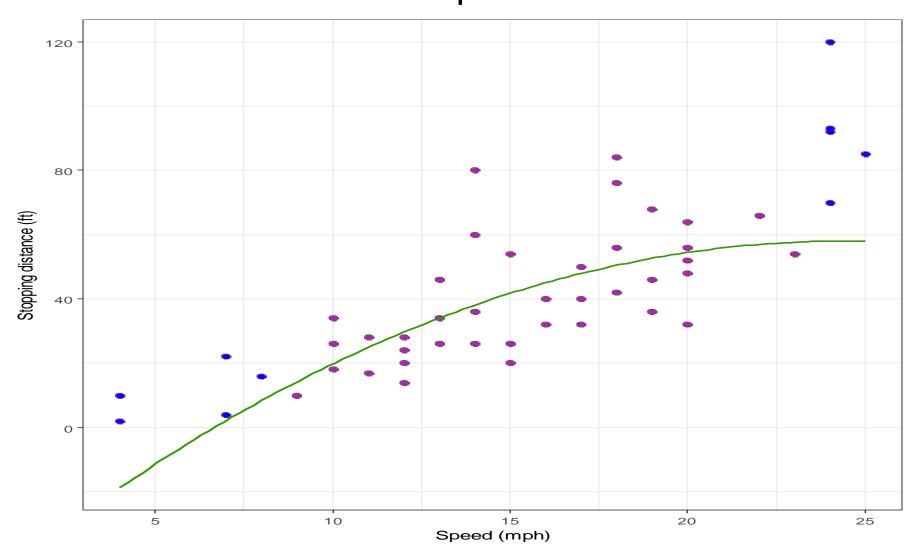
```
> lm2 <- lm(dist ~ speed + I(speed^2))</pre>
> summary(lm2)
Call:
lm(formula = dist ~ speed + I(speed^2))
Residuals:
   Min 1Q Median 3Q
                                 Max
-28.720 -9.184 -3.188 4.628 45.152
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 2.47014 14.81716 0.167 0.868
speed
     0.91329 2.03422 0.449 0.656
I(speed^2) 0.09996 0.06597 1.515 0.136
Residual standard error: 15.18 on 47 degrees of freedom
Multiple R-squared: 0.6673, Adjusted R-squared: 0.6532
F-statistic: 47.14 on 2 and 47 DF, p-value: 5.852e-12
```

## Covariance matrix of the vector of estimates

## Data and estimated model



# Predicted model with the 80% most central data points

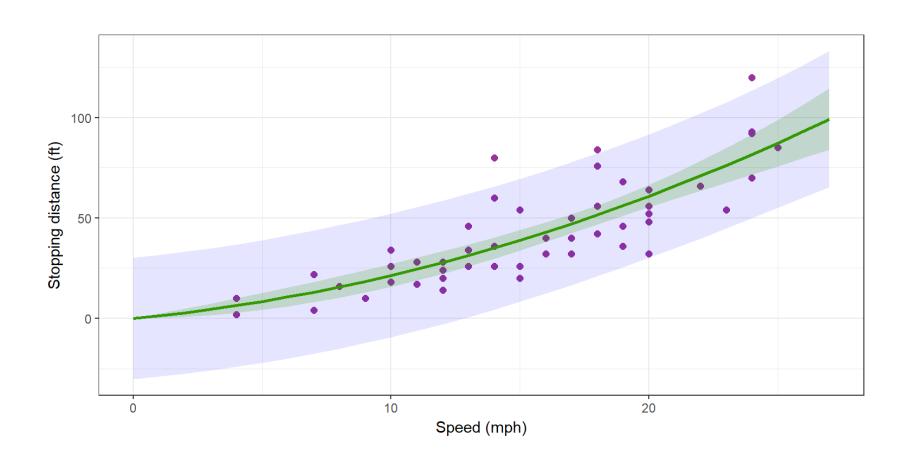


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# Fitting a polynomial without intercept

```
> lm2.noint <- lm(dist \sim -1 + speed + I(speed^2))
> coef(lm2.noint)
     speed I(speed^2)
1.23902996 0.09013877
> X <- model.matrix(lm2.noint)</pre>
> head(X)
  speed I(speed^2)
                  16
                  16
                                            Design matrix
3
                  49
                 49
5
                 64
                 81
```

# Data, predicted model, prediction intervals and confidence intervals



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# Orthogonal polynomials

```
> lm2.poly <- lm(dist ~ poly(speed, degree=2, raw=T))</pre>
> M <- model.matrix(lm2.poly)</pre>
> head(M)
  (Intercept) poly(speed, degree = 2, raw = T)1
2
3
4
5
                                                     8
                                                            Design
                                                     9
6
                                                            matrix
  poly(speed, degree = 2, raw = T)2
                                      16
2
                                      16
3
                                      49
                                      49
5
                                      64
6
                                      81
```

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  - Fitting a polynomial without intercept
  - Using orthogonal polynomials
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  - t-test
  - Analysis-of-variance (anova)
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#### T-test

- Fitting polynomial models
  - Fitting a polynomial of degree 0
  - Fitting a polynomial of degree 1
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  - Some diagnostic plots
  - The predictive performance of the model
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# anova(model1, model2)

```
> anova(lm0, lm1)  
Analysis of Variance Table  
Model 1: dist ~ 1   Y_i = \alpha + \varepsilon_i  
Model 2: dist ~ speed Y_i = \alpha + \beta \times X_i + \varepsilon_i  
Res.Df RSS Df Sum of Sq F Pr(>F)  
1   49 32539  
2   48 11354 1   21186 89.567 1.49e-12 ***  
---  
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

# anova(model1, model2)

```
> anova(lm1, lm2)
Analysis of Variance Table

Model 1: dist ~ speed
Model 2: dist ~ speed + I(speed^2)
   Res.Df   RSS Df Sum of Sq     F Pr(>F)
1     48 11354
2     47 10825     1     528.81 2.296 0.1364
```

- Fitting polynomial models
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# Likelihood ratio test (LRT)

```
> logLik(lm0)
'log Lik.' -232.9012 (df=2)
> logLik(lm1)
'log Lik.' -206.5784 (df=3)
> logLik(lm2)
'log Lik.' -205.386 (df=4)
>
> dl <- 2*as.numeric(logLik(lm1) - logLik(lm0))
> 1 pobisa(dl 1)
> 1-pchisq(dl,1)
[1] 3.995693e-13
>
> dl <- 2*as.numeric(logLik(lm2) - logLik(lm1))
> 1-pchisa(dl.1)
[1] 0.122521
```

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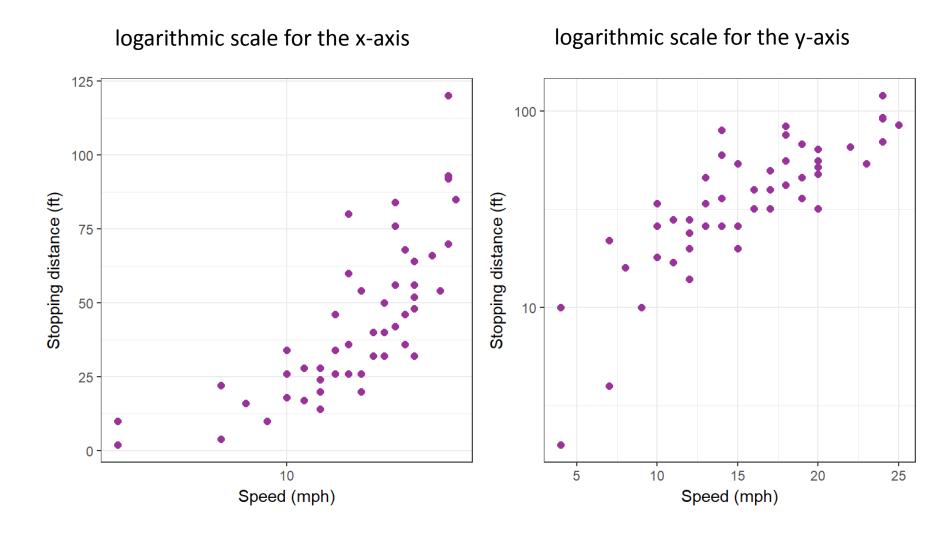
#### Information criteria

```
> AIC(lm0,lm1,lm2)
    df     AIC
lm0    2 469.8024
lm1    3 419.1569
lm2    4 418.7721

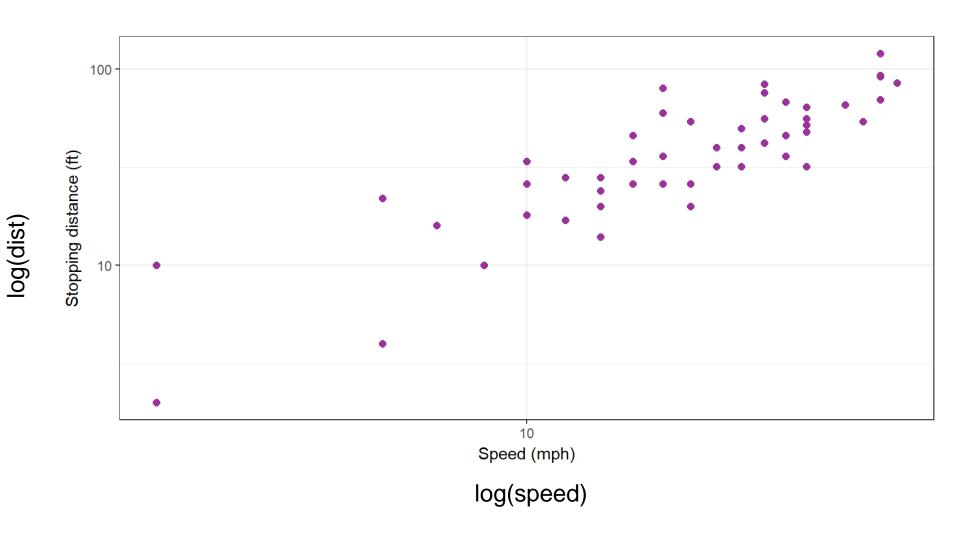
> BIC(lm0,lm1,lm2)
    df     BIC
lm0    2 473.6265
lm1    3 424.8929
lm2    4 426.4202
```

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# Logarithmic scale



# The data using log-log transformation.



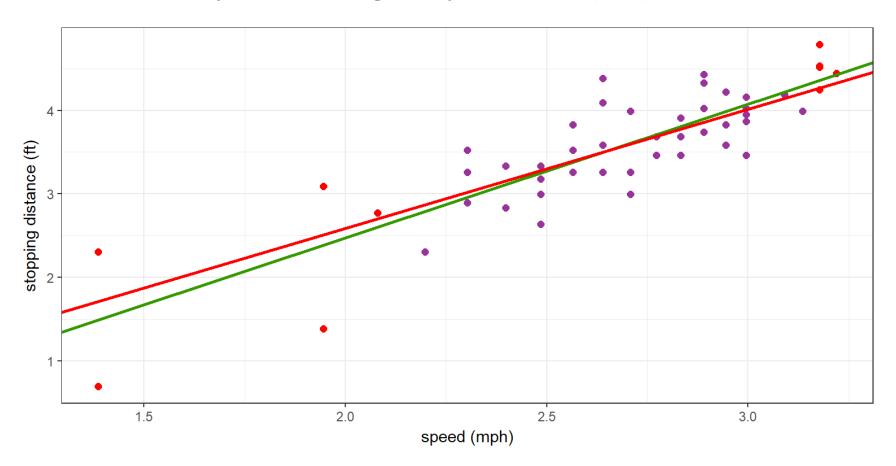
# Regression model for log(dist) and log(speed)

```
> lm1.log <- lm(log(dist) ~ log(speed)) log(Y_i) = \alpha + \beta \times log(X_i) + \varepsilon_i
> summary(lm1.log)
Call:
lm(formula = log(dist) \sim log(speed))
Residuals:
     Min 10 Median
                                 30
                                         Max
-1.00215 -0.24578 -0.02898 0.20717 0.88289
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) -0.7297 0.3758 -1.941 0.0581.
log(speed) 1.6024 0.1395 11.484 2.26e-15 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
Residual standard error: 0.4053 on 48 degrees of freedom
Multiple R-squared: 0.7331, Adjusted R-squared: 0.7276
F-statistic: 131.9 on 1 and 48 DF, p-value: 2.259e-15
```

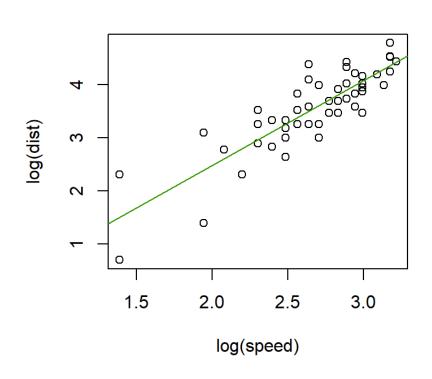
- Fitting polynomial models
  - Fitting a polynomial of degree 0
  - Fitting a polynomial of degree 1
  - Numerical results
  - Some diagnostic plots
  - The predictive performance of the model
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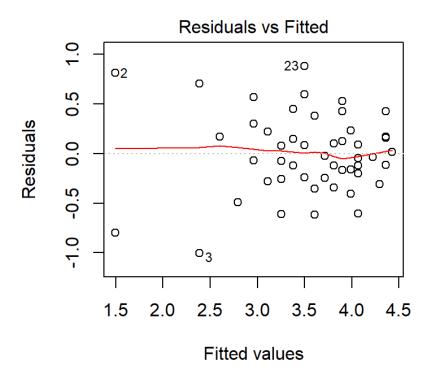
## Data and estimated models

A model fitted using the complete data (green) and a model fitted when only the training sample is used (red).



# Diagnostic plots

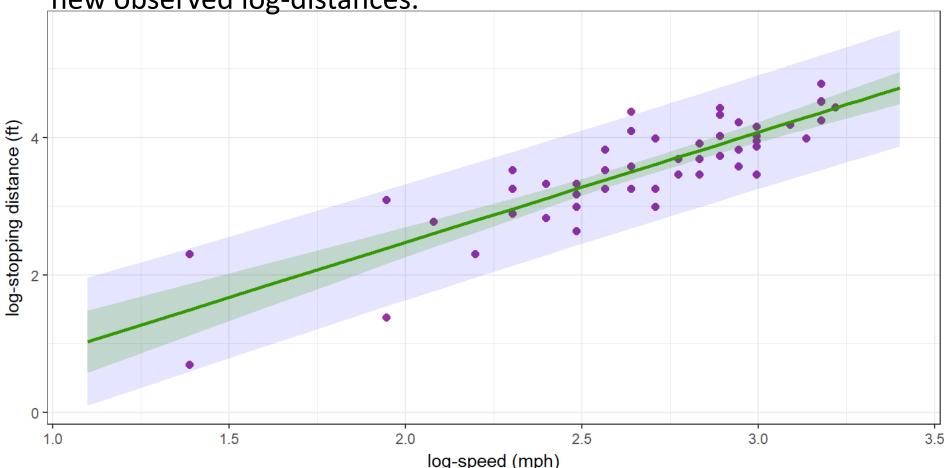




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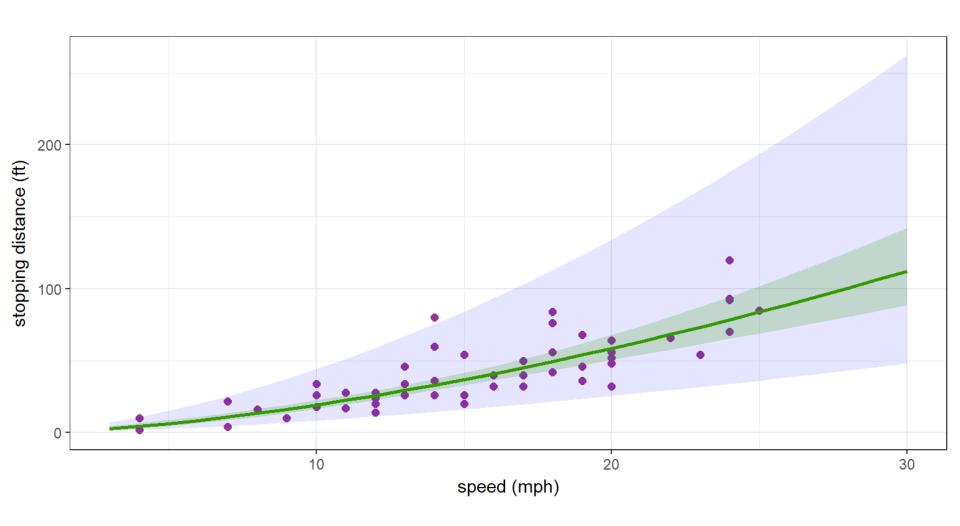
# Confidence and prediction intervals

Confidence interval for the regression line (of the model lm(log(dist) ~ log(speed))) and the prediction interval for new observed log-distances.



# Confidence and prediction intervals

Original scale.



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# Model comparison

```
> logLik(lm1.log)
'log Lik.' -24.76592 (df=3)
> logLik(lm1.log) - sum(log(dist))
'log Lik.' -201.5613 (df=3)
> AIC(lm1.log) - 2*sum(log(1/dist))
[1] 409.1226
> BIC(lm1.log) - 2*sum(log(1/dist))
[1] 414.8587
```