# 1 Simple Linear Regression

In simple linear regression, we predict values on one variable from the values of a second variable.

- The variable we are predicting is called the *dependent variable* (or response variable) and is referred to as Y.
- The variable we are basing our predictions on is called the **independent** variable (or predictor variable) and is referred to as X.

Remark: When there is only one predictor variable, the prediction method is called simple regression. Linear regression can have more than one predictor variable, i.e. Multiple Linear Regression.

In simple linear regression, the predicted values of Y when plotted as a function of X form a straight line on the scatter plot. This line is known as the **regression line**.

- Suppose we construct our model using n observed values of the response variable  $y_1, y_2, \ldots y_i \ldots y_n$
- For the original data set, there is a predicted value of each case of Y that corresponds to an observed value of Y.
- The difference between an observed value of the dependent variable  $(y_i)$  and the corresponding predicted value  $(\hat{y})$  is called the residual  $(e_i)$ . Each data point from the data set has one residual.
- Simply put, the values of the residuals are derived as follows:

Residual = Observed value - Predicted value

$$e_i = y_i - \hat{y}_i$$

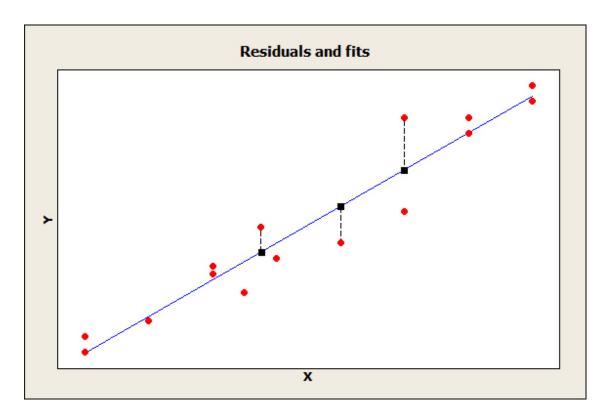


Figure 1:

- For three cases in the graphic above, the observed value (red dot) is linked to its corresponding predicted value (black dot) on the regression line (blue line). The difference (i.e. residual) is depicted using a dashed line. The magnitude of these residuals is of interest.
- The second of the three residuals will have a negative value.
- Ordinary Least Squares is a method of fitting a model, such that the total residual values are minimised.
- Important theoretical assumption underlying the OLS model: the sum of the residuals should equal to zero.

$$\sum e_i = 0$$

- An extension of this is that the expected value of the residuals is 0. E(e) = 0
- Another Important Theoretical Assumption The residuals are normally distributed. (more on that later)

### 1.1 Residual Plots

A residual plot is a graph that shows the residuals on the vertical axis and the independent variable on the horizontal axis. If the points in a residual plot are randomly dispersed around the horizontal axis, a linear regression model is appropriate for the data; otherwise, a non-linear model is more appropriate.

Some important terms in linear regression.

- **Residual:** The difference between the predicted value (based on the regression equation) and the actual, observed value.
- Outlier: In linear regression, an outlier is an observation with large residual. In other words, it is an observation whose dependent-variable value is unusual given its value on the predictor variables. An outlier may indicate a sample peculiarity or may indicate a data entry error or other problem.
- **Leverage:** An observation with an extreme value on a predictor variable is a point with high leverage. Leverage is a measure of how far an independent variable deviates from its mean. High leverage points can have a great amount of effect on the estimate of regression coefficients.
- **Influence:** An observation is said to be influential if removing the observation substantially changes the estimate of the regression coefficients. Influence can be thought of as the product of leverage and outlierness.
- Cook's distance (or Cook's D): A measure that combines the information of leverage and residual of the observation.

# 2 mtcars example

Several data sets, intended as learning tools, are automatically installed when R is installed. Many more are installed within packages to complement learning to use those packages. One of these is the famous **mtcars** data set, which is used in many data mining exercises.

```
> data(mtcars)
> head(mtcars)
                  mpg cyl disp hp drat
                                           wt qsec vs am gear carb
Mazda RX4
                 21.0
                        6 160 110 3.90 2.620 16.46 0
                        6 160 110 3.90 2.875 17.02 0
Mazda RX4 Wag
                 21.0
                                                       1
                                                                 4
Datsun 710
                        4 108 93 3.85 2.320 18.61 1 1
                 22.8
                                                                 1
```

```
Hornet 4 Drive 21.4 6 258 110 3.08 3.215 19.44 1 0 3 1
Hornet Sportabout 18.7 8 360 175 3.15 3.440 17.02 0 0 3 2
Valiant 18.1 6 225 105 2.76 3.460 20.22 1 0 3 1
```

Suppose we fit a model with mpg (miles per gallon) as the response variable and cyl and wt (number of cylinders and weight of the car) as the predictor variables. We will call this fitted model fit.

```
fit <- lm(mpg ~ cyl + wt, data=mtcars)
```

# > summary(fit)

### Call:

lm(formula = mpg ~ cyl + wt, data = mtcars)

### Residuals:

Min 1Q Median 3Q Max -4.2893 -1.5512 -0.4684 1.5743 6.1004

### Coefficients:

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

(Intercept) 39.6863 1.7150 23.141 < 2e-16 \*\*\*

cyl -1.5078 0.4147 -3.636 0.001064 \*\*

wt -3.1910 0.7569 -4.216 0.000222 \*\*\*

\_\_\_

Signif. codes: 0 \*\*\* 0.001 \*\* 0.01 \* 0.05 . 0.1 1

Residual standard error: 2.568 on 29 degrees of freedom Multiple R-squared: 0.8302, Adjusted R-squared: 0.8185

F-statistic: 70.91 on 2 and 29 DF, p-value: 6.809e-12

## residuals(fit1)

```
R File Edit View Misc Packages Windows Help
> residuals(fit)
        Mazda RX4
                      Mazda RX4 Wag
                                              Datsun 710
                                                            Hornet 4 Drive
                                                                               Hornet Sportabout
                        -0.46544677
        -1.27914467
                                              -3.45202624
                                                                   1.01948376
                                                                                       2.05304242
       Merc 230
-0.80351937
                          Merc 280
-0.46254751
                                              Merc 280C -1.86254751
                                                                   Merc 450SE
                                                                                      Merc 450SL
1.57842434
                                                                   1.76335487
  Chrysler Imperial
                                              Honda Civic
                                                               Toyota Corolla
                                                                                 Toyota Corona
                         5.76505710
Fiat X1-9
        4.13184435
                                               1.89833840
                                                                   6.10035227
                                                                                     -4.28933528
                                          Porsche 914-2
   Pontiac Firebird
                                                                 Lotus Europa
                                                                                Ford Pantera L
        3.84538614
                         -0.18055052
                                              -0.82640123
                                                                   1.57285924
                                                                                     -1.70852005
>
             McAfee LiveSafe – Internet Security
```

Figure 2:

```
> sum(residuals(fit))
[1] 1.096345e-15

> #Shapiro-Wilk Test for Normality
> shapiro.test(resid(fit))

Shapiro-Wilk normality test

data: resid(fit)
W = 0.9375, p-value = 0.06341
```

## 2.1 resid - Extracting Model Residuals

- residuals is a generic function which extracts model residuals from objects returned by modeling functions.
- The abbreviated form resid is an alias for residuals. It is intended to encourage users to access object components through an accessor function rather than by directly referencing an object slot.
- All object classes which are returned by model fitting functions should provide a residuals method. (Note that the method is for residuals and not resid.)
- Methods can make use of naresid methods to compensate for the omission of missing values. The default, nls and smooth.spline methods do.

residuals(fit)		
resid(fit)		

# 3 Some Important Definitions

## 3.1 Homoscedasticity

- *Homoscedascity* is the technical term to describe the variance of the residuals being constant across the range of predicted values.
- *Heteroscedascity* is the converse scenario: the variance differs along the range of values.

Suppose you plot the individual residuals against the predicted value, the variance of the residuals predicted value should be constant.

Consider the red arrows in the picture below, intended to indicate the variance of the residuals at that part of the number line. For the OLS summption to be valid, the length of the red lines should be more or less the same.

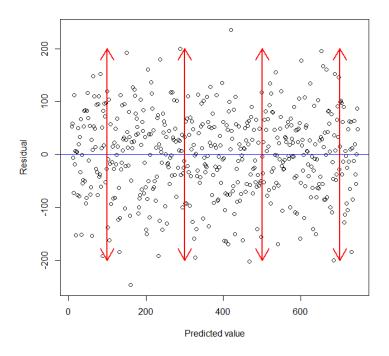


Figure 3:

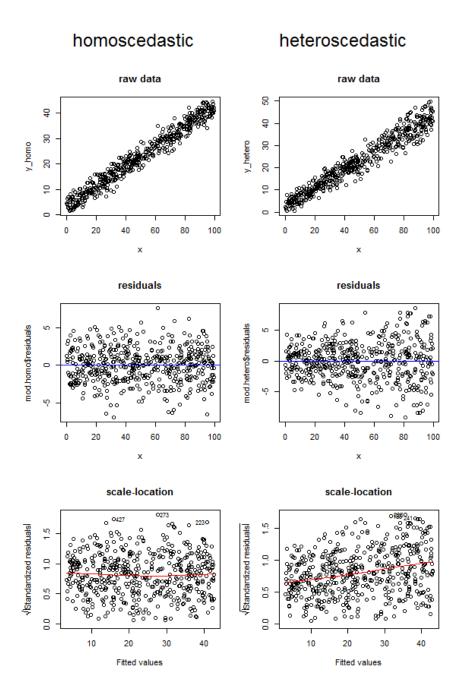


Figure 4:

#### 3.2 More Definitions

To understand a diagnostic plot called the residual-leverage plot, we need to understand three things:

- Leverage,
- Sstandardized residuals, and
- Cook's distance.

#### 3.2.1 Leverage

To understand leverage, recognize that *Ordinary Least Squares* regression fits a line that will pass through the centre of your data,  $(\bar{x}, \bar{y})$ . The line can be shallowly or steeply sloped, but it will pivot around that point like a lever on a fulcrum.

This analogy can be taken fairly literally: because OLS seeks to minimize the vertical distances between the data and the line, the data points that are further out towards the extremes of X will push / pull harder on the lever (i.e., the regression line); they have more leverage. One result of this could be that the results you get are driven by a few data points; that these diagnostic plots are intended to identify.

### 3.2.2 Standardization

Another result of the fact that points further out on X have more leverage is that they tend to be closer to the regression line (or more accurately: the regression line is fit so as to be closer to them) than points that are near  $\bar{x}$ . In other words, the residual standard deviation can differ at different points on X (even if the error standard deviation is constant). To correct for this, residuals are often standardized so that they have constant variance (assuming the underlying data generating process is homoscedastic, of course).

#### 3.2.3 Cook's Distance

One way to think about whether or not the results you have were driven by a given data point is to calculate how far the predicted values for your data would move if your model were fit without the data point in question.

This calculated total distance is called **Cook's distance**. Fortunately, you don't have to rerun your regression model N times to find out how far the predicted values will move, Cook's D is a function of the leverage and standardized residual associated with each data point.

With these facts in mind, consider the plots associated with four different situations:

- 1. a dataset where everything is fine
- 2. a dataset with a high-leverage, but low-standardized residual point
- 3. a dataset with a low-leverage, but high-standardized residual point
- 4. a dataset with a high-leverage, high-standardized residual point

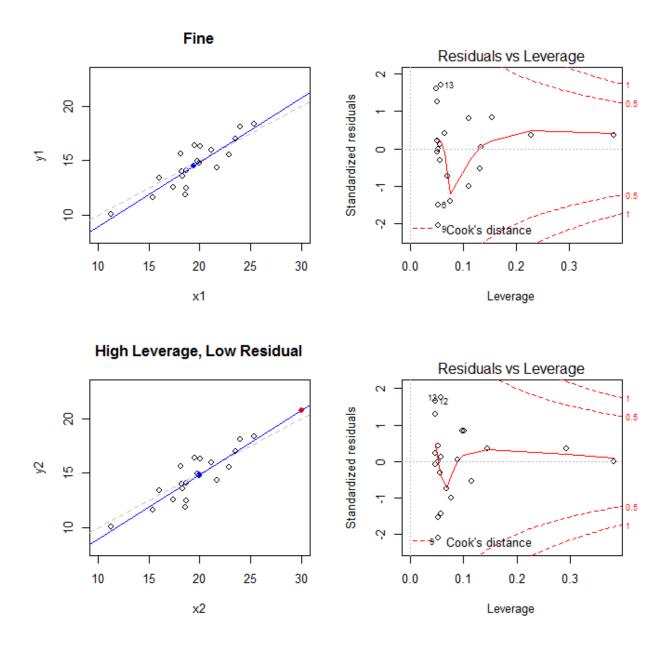


Figure 5:

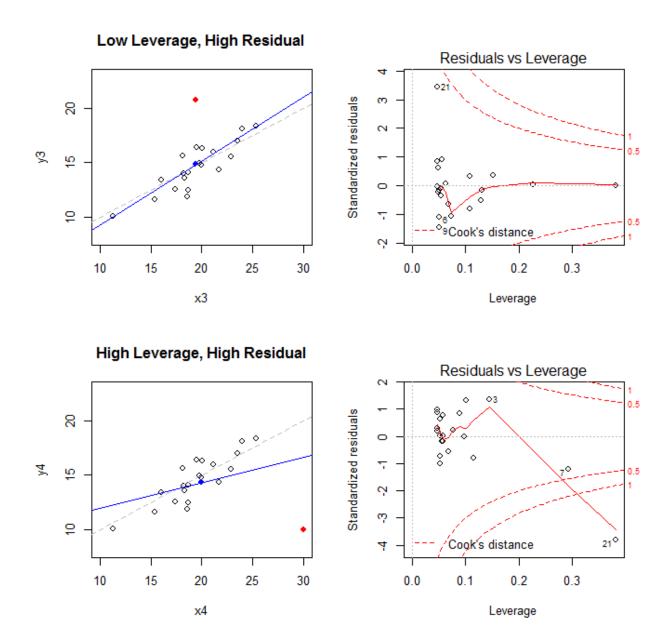


Figure 6:

- The plots on the left show the data, the center of the data with a blue dot, the underlying data generating process with a dashed gray line, the model fit with a blue line, and the special point with a red dot.
- On the right are the corresponding residual-leverage plots; the special point is 21.
- The model is badly distorted primarily in the fourth case where there is a point with high leverage and a large (negative) standardized residual.

## 4 Diagnostic Plots for Linear Models with R

There are six plots (selectable by which) are currently available:

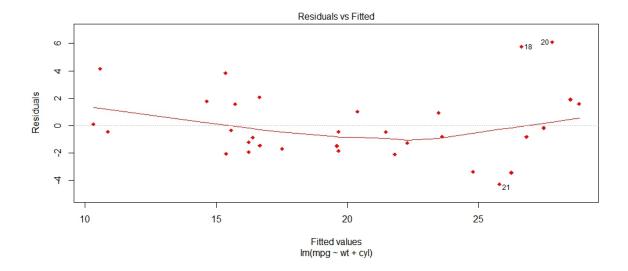
- 1. a plot of residuals against fitted values,
- 2. a Scale-Location plot of  $sqrt( \mid residuals \mid )$  against fitted values,
- 3. a Normal Q-Q plot,
- 4. a plot of Cook's distances versus row labels,
- 5. a plot of residuals against leverages,
- 6. a plot of Cook's distances against leverage/(1-leverage).

By default, the first three and 5 are provided, if you just type something like plot(fit).

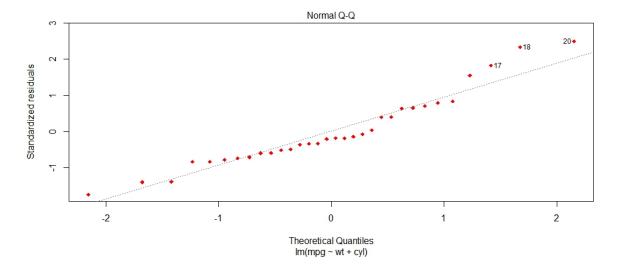
```
plot(lm(mpg~wt+cyl), which=c(1), pch=18, col="red")
plot(lm(mpg~wt+cyl), which=c(2), pch=18, col="red")
plot(lm(mpg~wt+cyl), which=c(3), pch=18, col="red")
plot(lm(mpg~wt+cyl), which=c(4), pch=18, col="red")
plot(lm(mpg~wt+cyl), which=c(5), pch=18, col="red")
plot(lm(mpg~wt+cyl), which=c(6), pch=18, col="red")
```

- The **Scale-Location** plot, also called Spread-Location (or S-L plot), takes the square root of the absolute residuals in order to diminish skewness  $(\operatorname{sqrt}(|E|))$  is much less skewed than |E| for Gaussian zero-mean E).
- Plot 5 The Residual-Leverage plot shows contours of equal Cook's distance, for values of cook.levels (by default 0.5 and 1) and omits cases with leverage one with a warning. If the leverages are constant (as is typically the case in a balanced aov situation) the plot uses factor level combinations instead of the leverages for the x-axis.

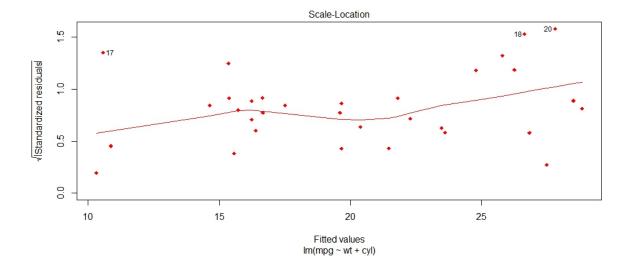
(The factor levels are ordered by mean fitted value.)



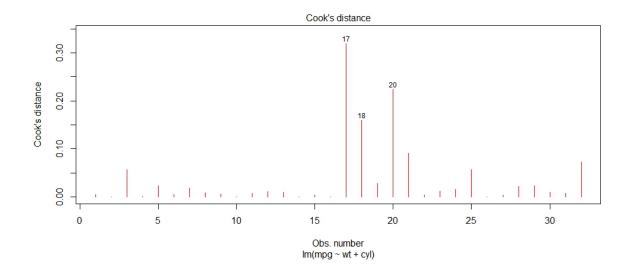
Plot 2: Normal Probability Plot

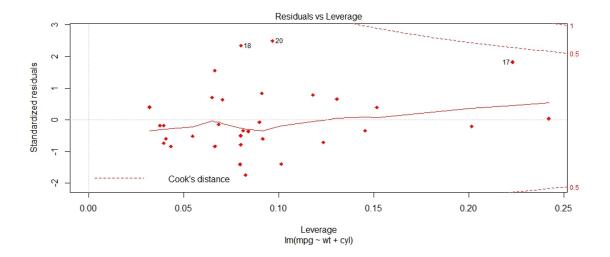


This plot is used to assess the validity of the normality of the residuals.

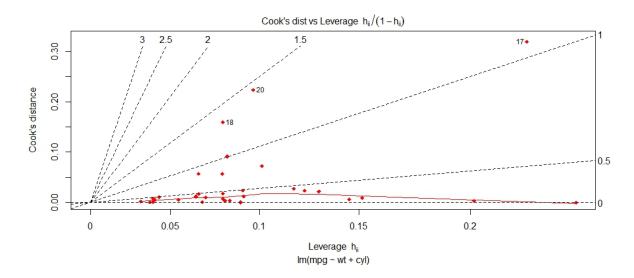


Plot 5: Cook's Distance





Plot 6 : Cook's Distance vs Leverage



Plot the four default plots together:

```
par(mfrow=c(4,1))
plot(fittedmodel)
par(opar)
```

### 5 Outliers and Influential Observations

"Outliers are sample values that cause surprise in relation to the majority of the sample" (W.N. Venables and B.D. Ripley. 2002. Modern applied statistics with S. New York: Springer, p.119).

Crucially, surprise is in the mind of the beholder and is dependent on some tacit or explicit model of the data.

There may be another model under which the outlier is not surprising at all, say if the data really are lognormal or gamma rather than normal.

#### 5.1 Outliers

Data points that diverge in a big way from the overall pattern are called outliers. There are four ways that a data point might be considered an outlier.

- It could have an extreme X value compared to other data points.
- It could have an extreme Y value compared to other data points.
- It could have extreme X and Y values.
- It might be distant from the rest of the data, even without extreme X or Y values.

After a regression line has been computed for a group of data, a point which lies far from the line (and thus has a large residual value) is known as an outlier. Such points may represent erroneous data, or may indicate a poorly fitting regression line. If a point lies far from the other data in the horizontal direction, it is known as an influential observation. The reason for this distinction is that these points have may have a significant impact on the slope of the regression line.

### 6 Cook's Distance

- In statistics, Cook's distance or Cook's D is a commonly used estimate of the influence of a data point when performing least squares regression analysis.
- Cook's distance is useful for identifying outliers in the X values (observations for predictor variables). It also shows the influence of each observation on the fitted response values.

- Cook's distance is the scaled change in fitted values. Each resulting element in a diagnostic calculation is the normalized change in the vector of coefficients due to the deletion of an observation.
- In a practical ordinary least squares analysis, Cook's distance can be used in several ways: to indicate data points that are particularly worth checking for validity; to indicate regions of the design space where it would be good to be able to obtain more data points.
- It is named after the American statistician R. Dennis Cook, who introduced the concept in 1977.
- Cook's distance measures the effect of deleting a given observation. Data points with large residuals (outliers) and/or high leverage may distort the outcome and accuracy of a regression.
- Points with a large Cook's distance are considered to merit closer examination in the analysis.
- Influential cases are not usually a problem when their removal from the dataset would leave the parameter estimates essentially unchanged: the ones we worry about are those whose presence really does change the results.
- Cook's D is a good measure of the influence of an observation and is proportional to the sum of the squared differences between predictions made with all observations in the analysis and predictions made leaving out the observation in question. If the predictions are the same with or without the observation in question, then the observation has no influence on the regression model. If the predictions differ greatly when the observation is not included in the analysis, then the observation is influential.

### Cook's Distance Formula

It is calculated as:

$$D_i = \frac{\sum_{j=1}^{n} (\hat{Y}_j - \hat{Y}_{j(i)})^2}{p \text{ MSE}},$$

where:

- $\hat{Y}_j$  is the prediction from the full regression model for observation j;
- $\hat{Y}_{j(i)}$  is the prediction for observation j from a refitted regression model in which observation i has been omitted;
- p is the number of fitted parameters in the model;
- MSE is the mean square error of the regression model.

For the case of simple linear regression, the following are the algebraically equivalent expressions

$$D_{i} = \frac{e_{i}^{2}}{p \text{ MSE}} \left[ \frac{h_{ii}}{(1 - h_{ii})^{2}} \right],$$
$$D_{i} = \frac{(\hat{\beta} - \hat{\beta}^{(-i)})^{T} (X^{T} X)(\hat{\beta} - \hat{\beta}^{(-i)})}{(1 + p)s^{2}},$$

where:

 $\bullet$   $h_{ii}$  is the leverage, i.e., the i-th diagonal element of the hat matrix

$$\mathbf{X} \left( \mathbf{X}^T \mathbf{X} \right)^{-1} \mathbf{X}^T$$

•  $e_i$  is the residual (i.e., the difference between the observed value and the value fitted by the proposed model).

# R code for computing Cook's Distance

```
attach(mtcars)
fit = lm(mpg ~ cyl + wt )
cooks.distance(fit)
plot(cooks.distance(fit),type="b",pch=18,col="red")

N = 32
k = 2
cutoff = 4/ (N-k-1)
abline(h=cutoff,lty=2)
```

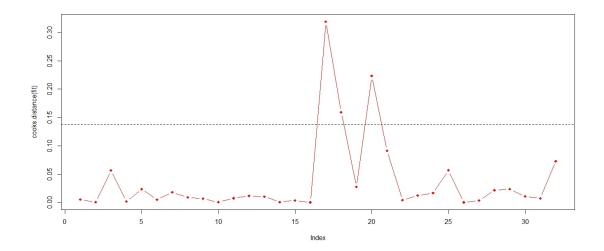


Figure 7:

### 6.1 Interpreting Cook's Distance

A common rule of thumb is that an observation with a value of Cook's D over 1.0 has too much influence. As with all rules of thumb, this rule should be applied judiciously and not thoughtlessly. (John Fox) (1991). Regression Diagnostics: An Introduction. Sage Publications.

- Some texts tell you that points for which Cook's distance is higher than 1 are to be considered as influential.
- Other texts give you a threshold of 4/N or

$$\frac{4}{(N-k-1)},$$

where N is the number of observations and k the number of explanatory variables.

- The R help file advises that an observation with Cook's distance larger than three times the mean Cook's distance might be an outlier. .
- John Fox (mentioned above), in his booklet on regression diagnostics is rather cautious when it comes to giving numerical thresholds. He advises the use of graphics and to examine in closer details the points with "values of D that are substantially larger than the rest". According to Fox, thresholds should just be used to enhance graphical displays.

#### Remark

- Cook's distance refers to how far, on average, predicted y-values will move if the observation in question is dropped from the data set.
- **DFBETA**: *DFBETA* refers to how much a parameter estimate changes if the observation in question is dropped from the data set.
- Note that with k covariates, there will be k+1 DFBETAs (the intercept,  $\beta_0$ , and  $\beta_1$  for each covariate).
- Cook's distance is arguably more important if you are doing predictive modeling, whereas *DFBETA* is more important in explanatory modeling.
- **DFFITS**: Although the raw values resulting from the equations are different, Cook's distance and *DFFITS* are conceptually identical and there is a closed-form formula to convert one value to the other.

# 7 Regression Deletion Diagnostics

This suite of functions can be used to compute some of the regression (leave-one-out deletion) diagnostics for linear and generalized linear models discussed in Belsley, Kuh and Welsch (1980), Cook and Weisberg (1982), etc.

### **Details**

- The primary high-level function is influence.measures which produces a class "infl" object tabular display showing the DFBETAS for each model variable, DFFITS, covariance ratios, Cook's distances and the diagonal elements of the hat matrix. Cases which are influential with respect to any of these measures are marked with an asterisk.
- The functions dfbetas, dffits, covratio and cooks.distance provide direct access to the corresponding diagnostic quantities.

- Functions rstandard and rstudent give the standardized and Studentized residuals respectively.
- (These functions re-normalize the residuals to have unit variance, using an overall and leave-one-out measure of the error variance respectively.)
- Values for generalized linear models are approximations, as described in Williams (1987) (except that Cook's distances are scaled as F rather than as chi-square values). The approximations can be poor when some cases have large influence.
- The optional infl, res and sd arguments are there to encourage the use of these direct access functions, in situations where, e.g., the underlying basic influence measures (from lm.influence or the generic influence) are already available.
- Note that cases with weights == 0 are dropped from all these functions, but that if a linear model has been fitted with na.action = na.exclude, suitable values are filled in for the cases excluded during fitting.
- The function hat() exists mainly for S (version 2) compatibility; we recommend using hatvalues() instead.

```
Usage
influence.measures(model)

rstandard(model, ...)

## S3 method for class 'lm'
rstandard(model, infl = lm.influence(model, do.coef = FALSE),
sd = sqrt(deviance(model)/df.residual(model)), ...)
```

```
## S3 method for class 'glm'
rstandard(model, infl = influence(model, do.coef = FALSE),
type = c("deviance", "pearson"), ...)
```

```
rstudent(model, ...)

## S3 method for class 'lm'
rstudent(model, infl = lm.influence(model, do.coef = FALSE),
res = infl$wt.res, ...)

## S3 method for class 'glm'
rstudent(model, infl = influence(model, do.coef = FALSE), ...)

dffits(model, infl = , res = )
```

## 8 Standardized and Studentized Residuals

The standardized residual is the residual divided by its standard deviation.

Plot the standardized residual of the simple linear regression model of the data set faithful against the independent variable waiting.

We apply the lm function to a formula that describes the variable eruptions by the variable waiting, and save the linear regression model in a new variable eruption.lm. Then we compute the standardized residual with the rstandard function.

```
> eruption.lm = lm(eruptions ~ waiting, data=faithful)
> eruption.stdres = rstandard(eruption.lm)
```

We now plot the standardized residual against the observed values of the variable waiting.

```
> plot(faithful$waiting, eruption.stdres,
+    ylab="Standardized Residuals",
+    xlab="Waiting Time",
+    main="Old Faithful Eruptions")
> abline(0, 0)  # the horizon
```

# 9 Leverage and Influence

#### 9.1 Influence

The influence of an observation can be thought of in terms of how much the predicted scores for other observations would differ if the observation in question were not included.

### 9.2 Influential Points

An influential point is an outlier that greatly affects the slope of the regression line. One way to test the influence of an outlier is to compute the regression equation with and without the outlier.

This type of analysis is illustrated below. The scatter plots are identical, except that the plot on the right includes an outlier. The slope is flatter when the outlier is present (-3.32 vs. -4.10), so this outlier would be considered an influential point.

### 9.3 Without Outlier

• Regression equation:  $\hat{y} = 104.78 - 4.10x$ 

• Coefficient of determination:  $R^2 = 0.94$ 

• Regression equation:  $\hat{y} = 97.51 - 3.32x$ 

• Coefficient of determination:  $R^2 = 0.55$ 

The charts below compare regression statistics for another data set with and without an outlier. Here, the chart on the right has a single outlier, located at the high end of the X axis (where x = 24). As a result of that single outlier, the slope of the regression line changes greatly, from -2.5 to -1.6; so the outlier would be considered an influential point.

Sometimes, an influential point will cause the coefficient of determination to be bigger; sometimes, smaller. In the first example above, the coefficient of determination is smaller when the influential point is present (0.94 vs. 0.55). In the second example, it is bigger (0.46 vs. 0.52).

If your data set includes an influential point, here are some things to consider.

• An influential point may represent bad data, possibly the result of measurement error. If possible, check the validity of the data point.

• Compare the decisions that would be made based on regression equations defined with and without the influential point. If the equations lead to contrary decisions, use caution.

### 9.4 Summary of Influence Statistics

### 9.5 rstudent

The studentized residual RSTUDENT is estimated by  $s(i)^2$  without the ith observation, not by  $s^2$ . For example,

RSTUDENT = 
$$\frac{r_i}{s_{(i)}\sqrt{(1-h_i)}}$$

Observations with RSTUDENT larger than 2 in absolute value may need some attention.

- Studentized Residuals Residuals divided by their estimated standard errors (like t-statistics). Observations with values larger than 3 in absolute value are considered outliers.
- Leverage Values (Hat Diag) Measure of how far an observation is from the others in terms of the levels of the independent variables (not the dependent variable). Observations with values larger than 2(k+1)/n are considered to be potentially highly influential, where k is the number of predictors and n is the sample size.
- **DFFITS** Measure of how much an observation has effected its fitted value from the regression model. Values larger than  $2\sqrt{(k+1)/n}$  in absolute value are considered highly influential.
- **DFBETAS** Measure of how much an observation has effected the estimate of a regression coefficient (there is one DFBETA for each regression coefficient, including the intercept). Values larger than 2/sqrt(n) in absolute value are considered highly influential.

The measure that measures how much impact each observation has on a particular predictor is DFBETAs The DFBETA for a predictor and for a particular observation is the difference between the regression coefficient calculated for all of the data and the regression coefficient calculated with the observation deleted, scaled by the standard error calculated with the observation deleted.

• Cooks D Measure of aggregate impact of each observation on the group of regression coefficients, as well as the group of fitted values. Values larger than 4/n are considered highly influential.

# 10 Other Measures of Influence

The impact of an observation on a regression fitting can be determined by the difference between the estimated regression coefficient of a model with all observations and the estimated coefficient when the particular observation is deleted. The measure DFBETA is the studentized value of this difference.

Influence arises at two stages of the LME model. Firstly when V is estimated by  $\hat{V}$ , and subsequent estimations of the fixed and random regression coefficients  $\beta$  and u, given  $\hat{V}$ .

#### 10.0.1 **DFBETA**

A group of measures that measures how much impact each observation has on a particular predictor are the DFBETAs. The DFBETA for a predictor and for a particular observation is the difference between the regression coefficient calculated for all of the data and the regression coefficient calculated with the observation deleted, scaled by the standard error calculated with the observation deleted.

$$DFBETA_a = \hat{\beta} - \hat{\beta}_{(a)} \tag{1}$$

$$= B(Y - Y_{\bar{a}} \tag{2})$$

For k predictors in the mode, there ar k+1 dfbetas.

### 10.0.2 **DFFITS**

DFFITS is a statistical measured designed to a show how influential an observation is in a statistical model. It is closely related to the studentized residual.

$$DFFITS = \frac{\widehat{y_i} - \widehat{y_{i(k)}}}{s_{(k)}\sqrt{h_{ii}}}$$

#### 10.0.3 COVRATIO

The COVRATIO statistic measures the change in the determinant of the covariance matrix of the estimates by deleting the ith observation:

$$COVRATIO = \frac{det(s_{(i)}^{2}(X'_{(i)}X_{(i)})^{-1}))}{det(s^{2}(X'X)^{-1})}$$

Observations with

$$|COVRATIO - 1| \ge \frac{3k}{n}$$

where k is the number of parameters in the model and n is the number of observations used to fit the model, are worth further investigation.

10.1 Influential Observations : DFBeta and DFBetas

```
R File Edit View Misc Packages Windows Help
> influence.measures(fit)
Influence measures of
        lm(formula = mpg ~ cyl + wt, data = mtcars) :
                   dfb.1_ dfb.cyl
-0.05324 -0.05593
                                     dfb.wt
                                              dffit cov.r
Mazda RX4
                                    0.07928 -0.1218 1.144 0.005077 0.0548
Mazda RX4 Waq
                   -0.01548 -0.00901
                                    0.01430 -0.0359 1.150 0.000444 0.0376
Datsun 710
                   0.02369 -0.01203 0.00931 0.0725 1.129 0.001803 0.0321
Hornet 4 Drive
                   -0.07203 0.21124 -0.14301 0.2643 1.136 0.023527 0.0912
Hornet Sportabout
Valiant
                   -0.02443 0.05191 -0.05731 -0.1213 1.116 0.005021 0.0407
Duster 360
                   0.07463 -0.17174  0.10163 -0.2300 1.132 0.017873 0.0801
Merc 240D
                   0.07726 -0.14506  0.11221  0.1628  1.289  0.009103  0.1517
                   -0.06771 0.12181 -0.09254 -0.1376 1.285 0.006506 0.1455
Merc 230
Merc 280
                   -0.00776  0.01506  -0.01648  -0.0367  1.153  0.000464  0.0396
Merc 280C
                   Merc 450SE
                   -0.09362
                            0.06992 0.01625
                                             0.1856 1.127 0.011685 0.0650
Merc 450SL
                   -0.06782
                            0.11419 -0.05099
                                             0.1738 1.146 0.010288 0.0705
Merc 450SLC
                   0.01596 -0.02424
                                    0.00940 -0.0389 1.190 0.000523 0.0685
Cadillac Fleetwood
                    0.04866
                           0.03982 -0.08368 -0.1015 1.385 0.003550 0.2015
Lincoln Continental -0.00967 -0.00917
                                     0.01786
                                             0.0209 1.466 0.000150 0.2421
                   -0.48080 -0.42853
                                     0.86142
                                             1.0217 0.991 0.318936 0.2230
Chrysler Imperial
Fiat 128
                    0.68604 -0.31604 -0.06239
                                             0.7543 0.644 0.159299 0.0802
                                                                            - 🛡 😽 📶 🐠
```

Figure 8:

# 11 Influential Points in Regression

```
inflm.fit <- influence.measures(fit)
which(apply(inflm.fit$is.inf, 1, any))</pre>
```

```
dfbeta(model, ...)
## S3 method for class 'lm'
dfbeta(model, infl = lm.influence(model, do.coef = TRUE), ...)

dfbetas(model, ...)
## S3 method for class 'lm'
dfbetas(model, infl = lm.influence(model, do.coef = TRUE), ...)

covratio(model, infl = lm.influence(model, do.coef = FALSE),
    res = weighted.residuals(model))
```

```
cooks.distance(model, ...)
## S3 method for class 'lm'
cooks.distance(model, infl = lm.influence(model, do.coef = FALSE),
res = weighted.residuals(model),
sd = sqrt(deviance(model)/df.residual(model)),
hat = infl$hat, ...)
## S3 method for class 'glm'
cooks.distance(model, infl = influence(model, do.coef = FALSE),
res = infl$pear.res,
dispersion = summary(model)$dispersion,
hat = infl$hat, ...)
```

```
hatvalues(model, ...)
## S3 method for class 'lm'
```

```
hatvalues(model, infl = lm.influence(model, do.coef = FALSE), ...)
hat(x, intercept = TRUE)
```

## Arguments

model an R object, typically returned by lm or glm.

infl influence structure as returned by lm.influence or influence (the larres (possibly weighted) residuals, with proper default.

sd standard deviation to use, see default.

dispersion dispersion (for glm objects) to use, see default.

hat hat values H[i,i], see default.

type type of residuals for glm method for rstandard.

x the X or design matrix.

intercept should an intercept column be prepended to x?

... further arguments passed to or from other methods.