CHAPTER 4

Model Adequacy Checking

Introduction

Assumptions

- 1. Relationship between response and regressors is linear (at least approximately).
- 2. Error term, ε has zero mean
- 3. Error term, ε has constant variance
- Errors are uncorrelated
- 5. Errors are normally distributed (required for tests and intervals)

Residual Analysis

Definition of Residual (= data – fit):

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

Approximate average variance:

$$\frac{\sum (e_i - \overline{e})^2}{n - p} = \frac{\sum e_i^2}{n - p} = \frac{SS_{\text{Res}}}{n - p} = MS_{\text{Res}}$$

• This is important quantity because it is the estimate of variance of residuals, or σ^2 .

- Scaling helps in identifying outliers or extreme values
- Four Methods
 - Standardized Residuals
 - 2. Studentized Residuals
 - 3. PRESS Residuals
 - 4. R-student Residuals

Standardized Residuals

$$d_i = \frac{e_i}{\sqrt{MS_{\text{Res}}}}$$

- d_i's have mean zero and variance approximately equal to 1
- Large values of d_i (d_i > 3) may indicate an outlier

2. Studentized Residuals

- MS_{Res} is only an approximation of the variance of the *i*th residual.
- o Improve scaling by dividing e_i by the exact standard deviation:

$$Var(e_i) = \sigma^2(1 - h_{ii})$$

What is h_{ii} ?

- Elements in the Hat matrix
- In simple linear regression case:

•
$$h_{ii} = \frac{1}{n} + \frac{(x_i - \bar{x})^2}{\sum_{i=1}^n (x_i - \bar{x})^2}$$

- The second part of right hand side of the above equation represents how far the data point is away from the center of the data set.
- If h_{ii} is large, a high leverage point (potential to affect to linear regression).
- If h_{ii} is small, a low leverage point.

Studentized ResidualsThe studentized residuals are then:

$$r_i = \frac{e_i}{\sqrt{\hat{\sigma}^2 (1 - h_{ii})}} = \frac{e_i}{\sqrt{MS_{Res} (1 - h_{ii})}}$$

- \circ r_i 's have mean zero and unit variance.
- Studentized residuals are generally larger than the corresponding standardized residuals.

PRESS Residuals (predicted residual sum of squares)

Examine the differences: $y_i - \hat{y}_{(i)}$

These are the differences between the actual response for the ith data point and the fitted value of the response for the ith data point, using all observations except the ith one.

3. PRESS Residuals

Logic:

- olf the *i*th point is unusual, then it can "overly" influence the regression model.
- o If the ith point is used in fitting the model, then the residual for the ith point will be impacted by the ith point.
- If the ith point is not used in fitting the model, then the residual will better reflect how unusual that point is.

3. PRESS Residuals

Prediction error:
$$e_{(i)} = y_i - \hat{y}_{(i)}$$

Calculate the PRESS residuals using

$$e_{(i)} = \frac{e_i}{1 - h_{ii}}$$

PRESS Residuals

$$Var\left[e_{(i)}\right] = Var\left[\frac{e_i}{1 - h_{ii}}\right]$$

$$= \frac{1}{(1 - h_{ii})^2} \sigma^2 (1 - h_{ii})$$

$$= \frac{\sigma^2}{(1 - h_{ii})}$$

- 3. PRESS Residuals
 - The standardized PRESS residuals are

$$\frac{e_{(i)}}{\sqrt{Var(e_{(i)})}} = \frac{e_i/(1-h_{ii})}{\sqrt{\sigma^2/(1-h_{ii})}} = \frac{e_i}{\sqrt{\sigma^2(1-h_{ii})}}$$

 Note: these are the studentized residuals when MS_{Res} is used as the estimate of the variance.

4. R-Student

- oMS_{Res} is an "internal" estimate of variance
- Ouse a variance estimate that is based on all observations except the ith observation:

$$S_{(i)}^{2} = \frac{(n-p)MS_{\text{Res}} - \frac{e_{i}^{2}}{(1-h_{ii})}}{n-p-1}$$

- 4. R-Student
 - The R-student residual is

$$t_{i} = \frac{e_{i}}{\sqrt{S_{(i)}^{2}(1 - h_{ii})}}$$

This is an externally studentized residual.

TABLE 4.1 Scaled Residuals for Example 4.1

Observation	$e_i = y_i - \hat{y}_i$	$d_i = e_i / \sqrt{MS_{\rm Res}}$	$r_i = e_i / \sqrt{MS_{\text{Res}}(1 - h_{ii})}$	h_{ii}	$e_{(i)} = e_i/(1 - h_{ii})$	$t_i = e_i / \sqrt{S_{(i)}^2 (1 - h_{ii})}$	$[e_i/(1-h_{ii})^2$
Number, i	(1)	(2)	(3)	(4)	(5)	(6)	(7)
1	-5.0281	-1.5426	-1.6277	0.10180	- 5.5980	-1.6956	31.3373
2	1.1464	0.3517	0.349	0.07070	1.2336	0.3575	1.5218
3	-0.0498	-0.0153	-0.0161	0.09874	-0.0557	-0.0157	0.0031
4	4.9244	1.5108	1.5798	0.05838	5.2297	1.6392	27.3499
5	-0.4444	-0.1363	-0.1418	0.07501	-0.4804	-0.1386	0.2308
6	-0.2896	-0.0888	-0.0908	0.04287	-0.3025	-0.0887	0.0915
7	0.8446	0.2501	0.2704	0.08180	0.9198	0.2646	0.8461
8	1.1566	0.3548	0.3667	0.06373	1.2353	0.3594	1.5260
9	7.4197	2.2763	3.2138	0.49829	14.7888	4.3108	218.7093
10	2.3764	0.7291	0.8133	0.19630	2.9568	0.8068	8.728
11	2.2375	0.6865	0.7181	0.08613	2.4484	0.7099	5.9946
12	-0.5930	-0.1819	-0.1932	0.11366	-0.6690	-0.1890	0.4476
13	1.0270	0.3151	0.3252	0.06113	1.0938	0.3185	1.1965
14	1.0675	0.3275	0.3411	0.07824	1.1581	0.3342	1.3412
15	0.6712	0.2059	0.2103	0.04111	0.7000	0.2057	0.4900
16	-0.6629	-0.2034	-0.2227	0.16594	-0.7948	-0.2178	0.6317
17	0.4364	0.1339	0.1381	0.05943	0.4640	0.1349	0.2153
18	3.4486	1.0580	1.1130	0.09626	3.8159	1.1193	14.5612
19	1.7932	0.5502	0.5787	0.09645	1.9846	0.5698	3.9387
20	-57880	-1.7758	-1.8736	0.10169	-6.4432	-1.9967	41.5150
21	-2.6142	-0.8020	-0.8779	0.16528	- 3.1318	-0.8731	9.8084
22	-3.6865	-1.1310	-1.4500	0.39158	- 6.0591	-1.4896	36.7131
23	-4.6076	-1.4136	-1.4437	0.04126	-4.8059	-1.4825	23.0966
24	-4.5728	-1.4029	-1.4961	0.12061	-5.2000	-1.5422	27.0397
25	-0.2126	-0.0652	-0.0675	0.06664	-0.2278	-0.0660	0.0519
						PRE	SS = 457.4000

R code

- # clean memory
- rm(list=ls())
- # read data
- delivery <- read.csv("eg3.1.delivery.csv",h=T)
- # obtain sample size
- n=dim(delivery)[1]
- names(delivery)
- # visualize data
- pairs (delivery,pch=20)
- # build linear regression
- model1 <- Im(DeliveryTime ~ NumberofCases+Distance, data=delivery)
- # obtain residual
- model1\$residuals
- # obtain SSRes and MSRes, note the MSRes is also our estimate of sigma square.
- SSRes=sum((model1\$residuals-mean(model1\$residuals))^2)
- MSRes=SSRes/(n-3)
- # obtain standardized residuals
- standardized_res=model1\$residuals/sqrt(MSRes)
- # obtain studentized residuals we first obtain leverage hii
- lm.influence(model1)\$hat
- # obtain studentized residuals
- studentized res=model1\$residuals/sqrt(MSRes)/sqrt(1 Im.influence(model1)\$hat)
- # a manual way to obtain studentized_res. We first need to construct X, which is matrix with the first column of 1s.
- X=as.matrix(cbind(1,delivery[,c("NumberofCases","Distance")]))
- # obtain hat matrix H
- H=X%*%solve(t(X)%*%X)%*%t(X)
- # obtain leverage
- diag(H)
- # manually obtain studentized residuals
- studentized_res2=model1\$residuals/sqrt(MSRes)/sqrt(1 diag(H))
- # PRESS residuals
- PRESS_res=model1\$residuals/(1 lm.influence(model1)\$hat)
- # R student
- R_Student=rstudent(model1)
- # a manual way to obtain the R student
- S2_i=((n-3)*MSRes-model1\$residuals^2/(1 diag(H)))/(n-3-1)
- R_Student2=model1\$residuals/sqrt(S2_i)/sqrt(1 diag(H))

R code

- # plot all residual and leverage
- # partition the canvas into 6 columns.
- par(mfrow=c(1,6))
- plot(model1\$fitted.values,model1\$residuals,pch=20,ylab="residual",xlab="fitted value")
- abline(h=0,col="grey")
- plot(model1\$fitted.values,standardized_res,pch=20,ylab="standardized residual",xlab="fitted value")
- abline(h=0,col="grey")
- plot(model1\$fitted.values,studentized_res,pch=20,ylab="studentized residual",xlab="fitted value")
- abline(h=0,col="grey")
- plot(model1\$fitted.values,PRESS_res,pch=20,ylab="PRESS residual",xlab="fitted value")
- abline(h=0,col="grey")
- plot(model1\$fitted.values,R_Student,pch=20,ylab="R student",xlab="fitted value")
- abline(h=0,col="grey")
- plot(model1\$fitted.values,lm.influence(model1)\$hat,pch=20,ylab="leverage",xlab="fitted value")

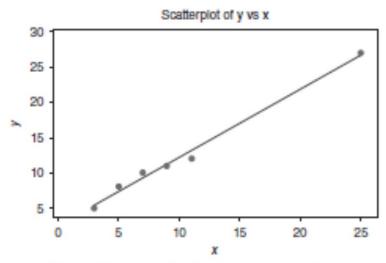


Figure 4.1 Example of a pure leverage point.

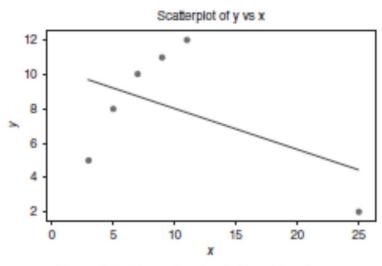


Figure 4.2 Example of an influential point.

- Normal Probability Plot of Residuals
 - Checks the normality assumption
- Residuals against Fitted values, \hat{y}_i
 - Checks for nonconstant variance
 - Checks for nonlinearity
 - Look for potential outliers

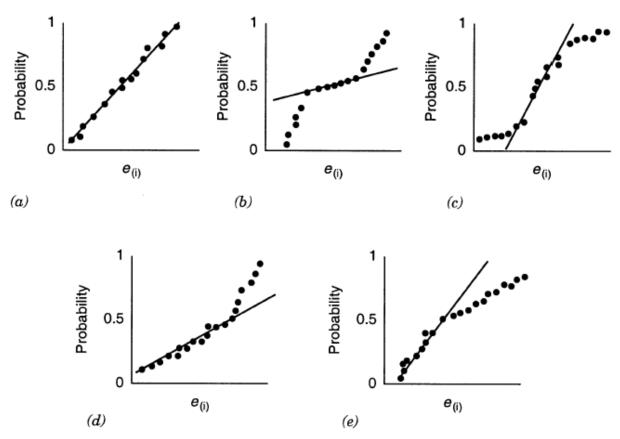


Figure 4.1 Normal probability plots: (a) ideal; (b) heavy-tailed distribution; (c) light-tailed distribution; (d) positive skew; (e) negative skew.

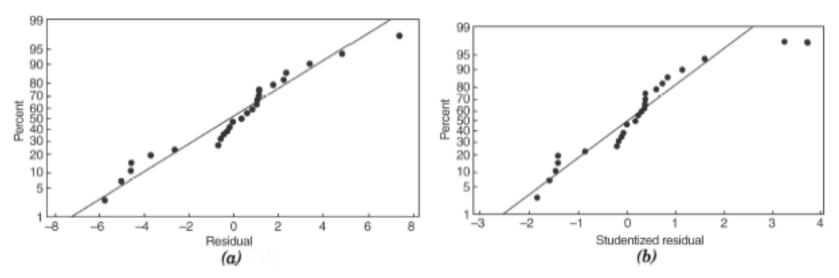


Figure 4.2 Normal probability plots of the residuals for the delivery time data: (a) ordinary least-squares residuals; (b) studentized residuals.

R code

- # generate QQ plot
- qqnorm(model1\$residuals)
- qqline(model1\$residuals)

- Residuals against Regressors in the model
 - Checks for nonconstant variance
 - Look for nonlinearity
- Residuals against Regressors not in the model
 - olf a pattern appears, could indicate that adding that regressor might improve the model fit.
- Residuals against time order
 - Check for Correlated errors

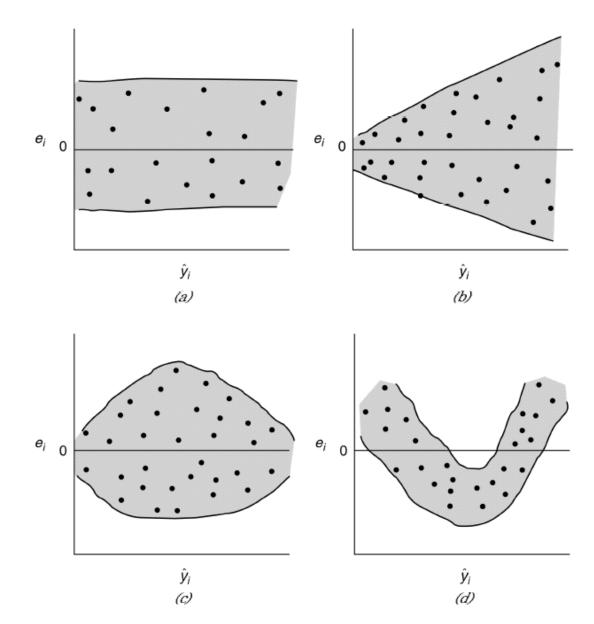


Figure 4.3 Patterns for residual plots: a) satisfactory; b) funnel; c) double bow; d) nonlinear.

R code

- # generate residual plot, NumberofCases vs residuals
- plot(delivery\$NumberofCases,model1\$residuals)
- # generate residual plot, Distance vs residuals
- plot(delivery\$Distance,model1\$residuals)
- # generate residual plot, fitted values vs residual
- plot(model1\$fitted.values,model1\$residuals)

Example 4.4 The Delivery Time Data

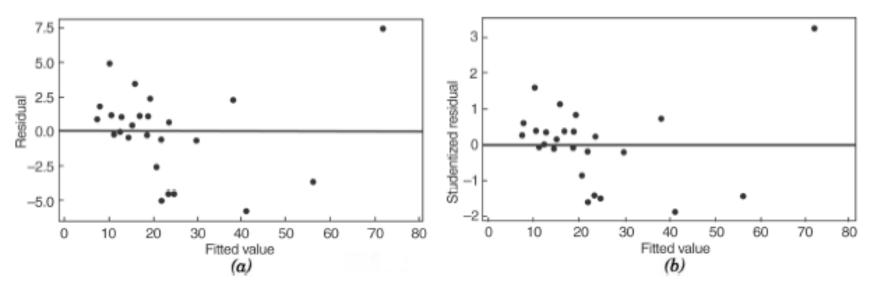


Figure 4.4 Plot of residuals versus predicted for the delivery time data: (a) original residuals; (b) studentized residuals.

Example 4.4 The Delivery Time Data

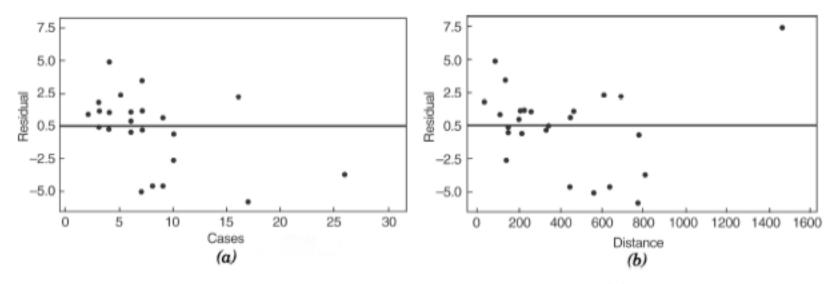


Figure 4.5 Plot of residuals versus the regressors for the delivery time data: (a) residuals versus cases; (b) residuals versus distance.

Plot of Residuals in Time Sequence

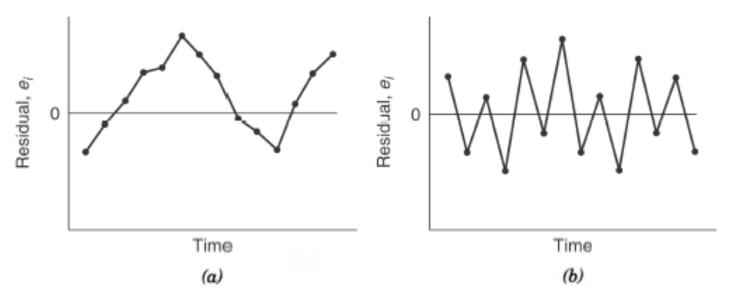


Figure 4.6 Prototype residual plots against time displaying autocorrelation in the errors: (a) positive autocorrelation; (b) negative autocorrelation.

R code (rocket propellant example)

- # example
- rm(list=ls())
- rocket <- read.delim("Data-ex-2-1 (Rocket Prop).txt",h=T)
- n=dim(rocket)[1]
- plot(rocket\$x,rocket\$y,pch=20)
- model1 <- Im(y ~ x, data=rocket)
- abline(model1,col="blue")
- # residual
- model1\$residuals
- #standardized residuals
- SSRes=sum((model1\$residuals-mean(model1\$residuals))^2)
- MSRes=SSRes/(n-3)
- standardized res=model1\$residuals/sqrt(MSRes)
- # studentized residuals
- studentized res=model1\$residuals/sqrt(MSRes)/sqrt(1 Im.influence(model1)\$hat)
- # PRESS residuals
- PRESS_res=model1\$residuals/(1 Im.influence(model1)\$hat)
- # R student
- R Student=rstudent(model1)
- # plot all residual and leverage
- # partition the canvas into 6 columns.
- par(mfrow=c(1,6))
- plot(model1\$fitted.values,model1\$residuals,pch=20,ylab="residual",xlab="fitted value")
- abline(h=0,col="grey")
- plot(model1\$fitted.values,standardized_res,pch=20,ylab="standardized residual",xlab="fitted value")
- abline(h=0,col="grey")
- plot(model1\$fitted.values,studentized res,pch=20,ylab="studentized residual",xlab="fitted value")
- abline(h=0,col="grey")
- plot(model1\$fitted.values,PRESS res,pch=20,ylab="PRESS residual",xlab="fitted value")
- abline(h=0,col="grey")
- plot(model1\$fitted.values,R Student,pch=20,ylab="R student",xlab="fitted value")
- abline(h=0,col="grey")
- plot(model1\$fitted.values,lm.influence(model1)\$hat,pch=20,ylab="leverage",xlab="fitted value")

- Partial Regression Plots
 - Purpose:
 - $_{\circ}$ To determine if the correct relationship between y and x_i has been identified
 - To determine the marginal contribution of a variable, given all other variables are in the model.

- Partial Regression Plots Method:
 - Regress y against all variables except x_i and calculate residuals
 - Regress x_i against all other regressor variables and calculate residuals
 - Plot these two sets of residuals against each other.

- Partial Regression Plots Interpretation:
 - If the plot appears to be linear, then a linear relationship between y and x_i seems reasonable
 - If plot is curvilinear, may need x_i² or 1/x_i instead
 - If x_i is a candidate variable, and a horizontal "band" appears, then that variable adds no new information.

Example 4.5

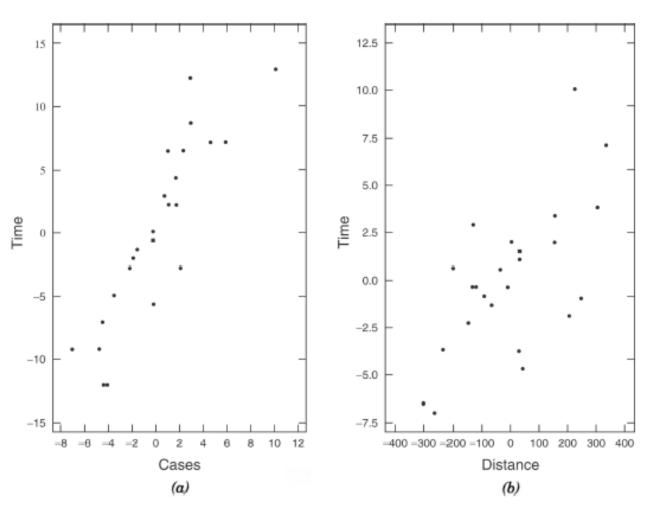
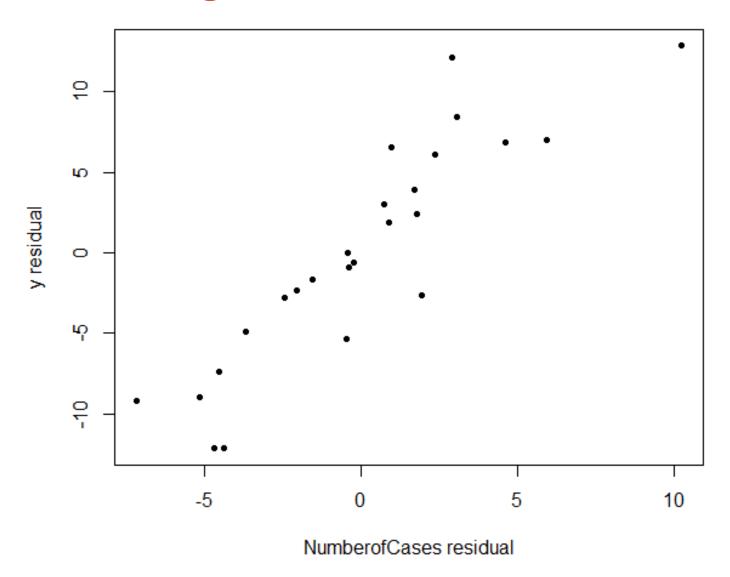


Figure 4.7 Partial regression plots for the delivery time data.

R code

- # generate partial regression plot
- # suppose we first have one regression with only Distance as covariate.
- model2 <- Im(DeliveryTime ~ Distance, data=delivery)
- # now we are considering if we want to add a new covariate NumberofCases. So we first regress NumberofCases on Distance and obtain residuals.
- model3 <- Im(NumberofCases ~ Distance, data=delivery)
- # we plot the residual from model3 against the residuals from model2 and see if there is a linear relationship. If yes, then we should include this new variable Number of Cases.
- plot(model3\$residuals,model2\$residuals,pch=20,ylab="y residual", xlab="NumberofCases residual")

Partial Regression Plot



- Partial Regression Plots
 Comments:
 - Use with caution, they only suggest possible relationships
 - Do not generally detect interaction effects
 - If multicollinearity is present, regression plots could give incorrect information
 - The slope of the partial regression plot is the regression coefficient for the variable of interest

Other Residual Plotting and Analysis Methods

- Plotting regressors against each other can give information about the relationship between the two:
 - o may indicate correlation between the regressors.
 - o may uncover remote points

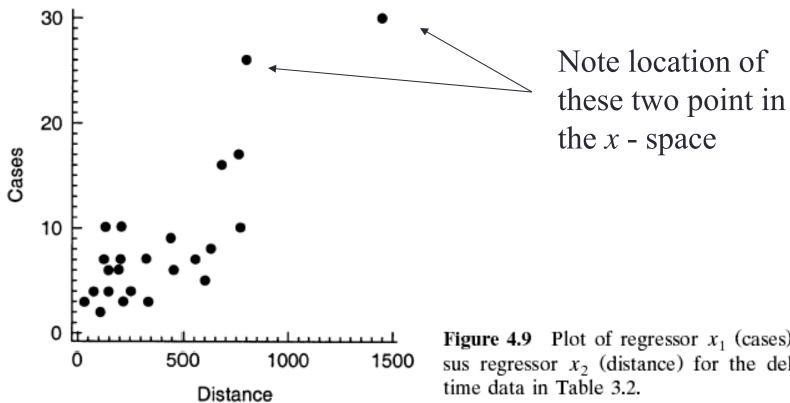


Figure 4.9 Plot of regressor x_1 (cases) versus regressor x_2 (distance) for the delivery

The PRESS Statistic

- PRESS Residual:
 - Prediction Error Sum of Squares (PRESS) Statistic:

$$e_{(i)} = y_i - \hat{y}_{(i)}$$

$$PRESS = \sum (y_i - \hat{y}_{(i)})^2$$

$$= \sum \left(\frac{e_i}{1 - h_{ii}}\right)^2$$

A small value of the PRESS Statistic is desired.

The PRESS Statistic

R² for Prediction Based on PRESS

$$R_{prediction}^2 = 1 - \frac{PRESS}{SS_T}$$

- Interpretation:
 - We expect the model to explain about R²% of the variability in prediction of a new observation.
- PRESS is a valuable statistic for comparison of models.

R code

- # To obtain R_square_prediction, we first calculate PRESS (prediction error sum of square)
- PRESS=sum(PRESS_res^2)
- # then obtain SST
- SST=sum((delivery\$DeliveryTimemean(delivery\$DeliveryTime))^2)
- # finally, we obtain R square prediction
- R_square_pred=1-PRESS/SST

Outliers

- An outlier is an observation that is considerably different from the others
- Formal tests for outliers
- Points with large residuals may be outliers
- Impact can be assessed by removing the points and refitting