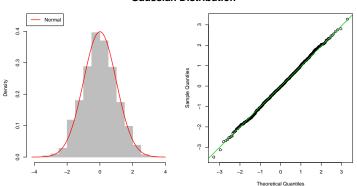
# Chapter 4 Model Adequacy Checking

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# Patterns of Q-Q plot (Normal probability plot)

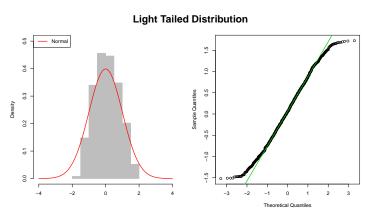
#### 1. Gaussian distribution

#### **Gaussian Distribution**



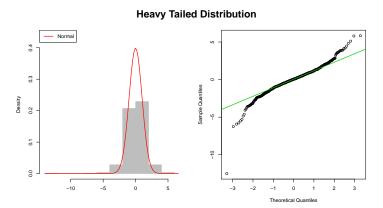
# Patterns of Q-Q plot (Normal probability plot) (cont.)

## 2. Light tailed distribution



## Patterns of Q-Q plot (Normal probability plot) (cont.)

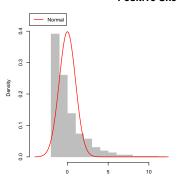
## 3. Heavy tailed distribution

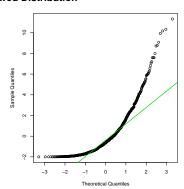


## Patterns of Q-Q plot (Normal probability plot) (cont.)

#### 4. Positive skewed distribution

#### Positive Skewed Distribution

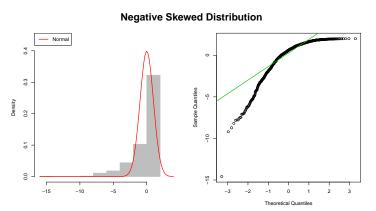




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# Patterns of Q-Q plot (Normal probability plot) (cont.)

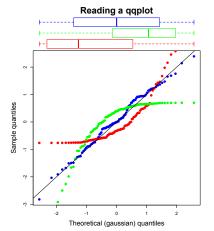
## 5. Negative skewed distribution



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# Patterns of Q-Q plot (Normal probability plot) (cont.)

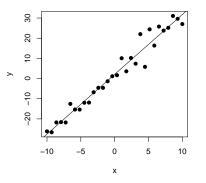
6. Three types of Q-Q plot

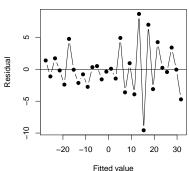


## Patterns of residual plots

1. Non independent errors (negative autocorrelation)

#### Non independent errors(negative autocorrelation)

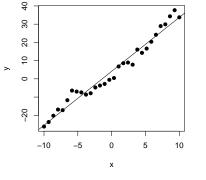


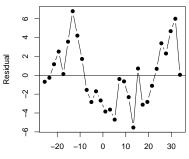


# Patterns of residual plots (cont.)

## 2. Non independent errors (positive autocorrelation)

#### Non independent errors(positive autocorrelation)





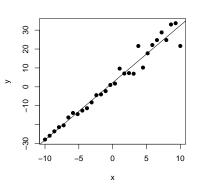
Fitted value

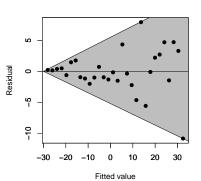
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# Patterns of residual plots (cont.)

## 3. Non constant variance (funnel)

#### Non constant variance(funnel)



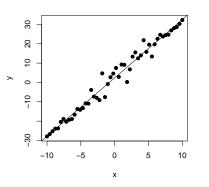


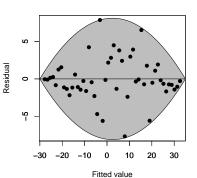
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# Patterns of residual plots (cont.)

4. Non constant variance (double bow)

#### Non constant variance(double bow)

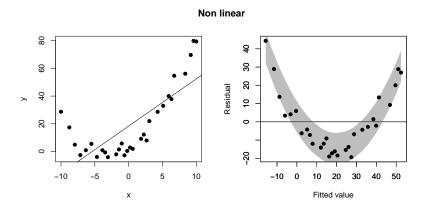




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# Patterns of residual plots (cont.)

## 5. Non linear



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## Example 4.2 The Delivery Time Data

## 1. Various types of residuals

```
> url <- "https://raw.github.com/dongikjang/regression/master/"
> rfun <- getURL(paste(http, "scaled.R",sep=""))</pre>
> eval(parse(text=rfun))
>
> scaled
function(model, type="standardized")
 UseMethod("scaled")
>
> scaled.lm
function(model, type="standardized"){
  switch(type,
         studentized = rstandard(model),
         rstudent = rstudent(model),
         standardized = residuals(model)/summary(model)$sigma
```

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```
> # Data download
> rfun <- getURL(paste(http, "read.xls2.r",sep=""))</pre>
> eval(parse(text=rfun))
> # If OS is Windows then install "xlsReadWrite" package
> # If OS is Mac or Linux then install "gdata" package
>
> library(RCurl)
> tf <- paste(tempfile(), "xls", sep = ".")</pre>
> download.file(paste(url, "Dataset/data-ex-3-1.xls", sep=""), tf, met
 % Total % Received % Xferd Average Speed Time
                                                        Time
                                                                 Time
                                Dload Upload Total Spent Left
                                    0 0 --:--:--:
                  0
                       0
                             0
> data_3.1 <- read.xls2(tf, header=TRUE)</pre>
> View(data_3.1)
> colnames(data_3.1) <- c("obs", "d_time", "n_case", "dista")</pre>
```

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```
> # Linear fit
> lmfit <- lm(d_time~n_case+dista)</pre>
> # standardized residuals
> scaled(lmfit)
          1
-1.54260631 0.35170879 -0.01527661
                                     1.51078203 -0.13634053 -0.0888408
                      10
                                  11
                                              12
                                                           13
2.27635117 0.72907878 0.68645843 -0.18194377
                                                  0.31508443
                                                               0.3275178
         17
                      18
                                  19
                                              20
                                                           21
0.13387449 1.05803019 0.55014821 -1.77573772 -0.80202492 -1.1310194
         25
-0.06522033
```

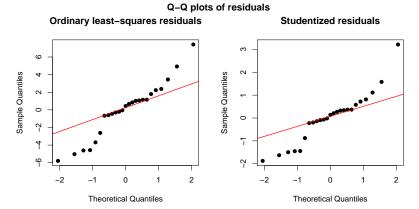
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```
> # standardized. studentized and rstudentized residuals
> residual_mat <- cbind(residuals(lmfit), scaled(lmfit),
                        scaled(lmfit, "studentized"),
                        scaled(lmfit, "rstudent"))
> colnames(residual_mat) <- c("residual", "stadardized",
                              "studentized". "rstudent")
> head(residual mat)
   residual stadardized studentized rstudent
1 -5.0280843 -1.54260631 -1.62767993 -1.69562881
2 1.1463854 0.35170879 0.36484267 0.35753764
3 -0.0497937 -0.01527661 -0.01609165 -0.01572177
4 4.9243539 1.51078203 1.57972040 1.63916491
5 -0.4443983 -0.13634053 -0.14176094 -0.13856493
6 -0.2895743 -0.08884082 -0.09080847 -0.08873728
```

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# Example 4.2 The Delivery Time Data (cont.)

## 2. Q-Q plots of residuals



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```
> par(mfrow=c(1,2), cex.main=1.2, pch=19, cex=1.5)
>
> qqnorm(residuals(lmfit), main='Ordinary least-squares residuals')
> qqline(residuals(lmfit), col=2, lwd=2)
>
> qqnorm(scaled(lmfit, "studentized"), main='Studentized residuals')
> qqline(scaled(lmfit, "studentized"), col=2, lwd=2)
> title(main='Q-Q plots of residuals',line=-1,outer=T)
```

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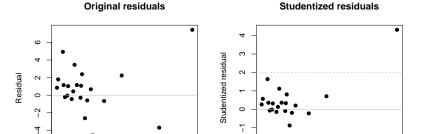
# Example 4.2 The Delivery Time Data (cont.)

#### 2. Residuals vs Predicted

φ

10

Fitted value



70

7

10 20

Residuals vs predicted for the delivery time data

Fitted value

70

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```
> par(mfrow=c(1,2), cex.main=1.2, pch=19, cex=1.5)
> fit_val <- fitted(lmfit)</pre>
> plot(fit_val, residuals(lmfit), xlab="Fitted value",
        vlab="Residual", main="Original residuals")
> abline(h=0, lty=1, col="grey")
> plot(fit_val, scaled(lmfit, "rstudent"), xlab="Fitted value",
       ylab="Studentized residual", main="Studentized residuals")
> abline(h=c(0,-2,2), lty=c(1,2,2), col="grey")
> title(main='Residuals vs predicted for the delivery time data',
        line=-1,outer=T)
```

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# Example 4.2 The Delivery Time Data (cont.)

## 3. Residuals vs Regressors



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```
> par(mfrow=c(1,2), cex.main=1.2, pch=19, cex=1.5)
> fit_val <- fitted(lmfit)</pre>
> plot(n_case, residuals(lmfit), xlab="Cases",
       vlab="Residual", main="Residuals vs cases")
> abline(h=0, lty=1, col="grey")
> plot(dista, residuals(lmfit), xlab="Distance",
       ylab="Residual", main="Residuals vs distance")
> abline(h=0, lty=1, col="grey")
> title(main='Residuals vs regressors for the delivery time data',
        line=-1,outer=T)
```

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## Example 4.2 The Delivery Time Data (cont.)

## 4. Partial regression plots

- Model:

$$y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \epsilon$$

- Partial residual 1:

$$\hat{y}_i(x_2) = \hat{\theta}_0 + \hat{\theta}_1 x_{i2}$$
  
 $e_i(y|x_2) = y_i - \hat{y}_i(x_2), \quad i = 1, 2, ..., n$ 

- Partial regressor 2:

$$\hat{x}_{i1}(x_2) = \hat{\alpha}_0 + \hat{\alpha}_1 x_{i2} 
e_i(x_1|x_2) = x_{i1} - \hat{x}_{i1}(x_2), \quad i = 1, 2, \dots, n$$

- Partial regression plots: plotting  $e_i(y|x_2)$  against  $e_i(x_1|x_2)$ .

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## Example 4.2 The Delivery Time Data (cont.)

#### 5. Partial residual plots

- The partial residual for regressor  $x_i$ :

$$e_i^*(y|x_i) = e_i + \hat{\beta}_i x_{ii}, \quad i = 1, 2, ..., n$$

where the  $e_i$  are the residuals from the model with all k regressors included.

- Partial residual plots: plotting  $e_i^*(y|x_i)$  against  $x_i$ .

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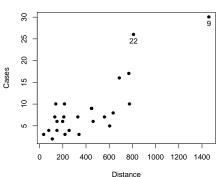
```
> partial <- function(model, part)</pre>
+ UseMethod("partial")
>
> partial <- function(model, part){</pre>
          x <- model$model[, part]
+
          coeff <- model$coefficients[part]</pre>
+
          resi <- c(residuals(model) + x*coeff)</pre>
          return(resi)
+ }
>
> par(mfrow=c(1,2), cex.main=1.2, pch=19, cex=1.5)
> plot(n_case, partial(lmfit, "n_case"), pch=16,cex=1.3,
       xlab='Cases', ylab='Time', main='Time vs Cases')
> plot(dista, partial(lmfit, "dista"), pch=16, cex=1.3,
+ xlab='Distance', ylab='Time', main='Time vs Distance')
> title(main='Partial residual plots for the delivery time data',
        line=-1,outer=T)
```

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## Example 4.2 The Delivery Time Data (cont.)

## 6. Regressor vs Regressor

# Regressor vs regressor for the delivery time data Cases vs Distance



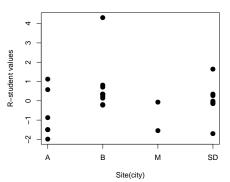
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## Example 4.2 The Delivery Time Data (cont.)

## 7. R-student values by site(city)

#### R-student values by site(city) for the delivery time data



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## Example 4.2 The Delivery Time Data (cont.)

#### 8. PRESS statistics

$$PRESS = \sum_{i=1}^{n} [y_i - \hat{y}_{(i)}]^2 = \sum_{i=1}^{n} \left(\frac{e_i}{1 - h_{ii}}\right)^2$$

```
> press <- function(obj){
+ sum((resid(obj)/(1-hatvalues(obj)))^2)
</pre>
```

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## Example 4.2 The Delivery Time Data (cont.)

-  $R^2$  for prediction based on PRESS

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

```
> 1-press(lmfit)/sum((d_time-mean(d_time))^2)
[1] 0.9206438
```

- Using PRESS to compare Models

```
> press(lm(d_time ~ n_case))
[1] 733.55
> press(lm(d_time ~ n_case + dista))
[1] 459.0393
```

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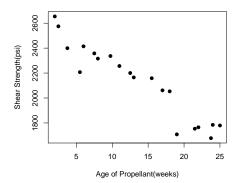
## Example 4.7 The Rocket Propellant Data

#### 1. Data and Plots

obs	yi	xi
1	2158.70	15.50
2	1678.15	23.75
3	2316.00	8.00
4	2061.30	17.00
5	2207.50	5.50
:	:	:
19	2654.20	2.00
20	1753.70	21.50

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## Example 4.7 The Rocket Propellant Data (cont.)



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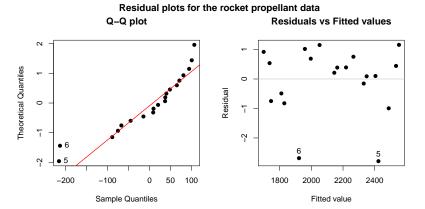
## Example 4.7 The Rocket Propellant Data (cont.)

```
> tf <- paste(tempfile(), "xls", sep = ".")</pre>
> download.file(paste(url, "Dataset/data-ex-2-1.xls", sep=""), tf, met
 % Total % Received % Xferd Average Speed Time
                                                       Time
                                                                Time
                                Dload Upload Total
                                                       Spent Left
                       0
                             0
                                    0 0 --:--:
> data_2.1 <- read.xls2(tf, header=TRUE)</pre>
> colnames(data_2.1) <- c("obs", "yi", "xi")</pre>
> attach(data 2.1)
>
> par(mfrow=c(1,1), pch=16, cex=1.4)
> plot(xi, yi, pch=19, xlab="Age of Propellant(weeks)",
      vlab="Shear Strength(psi)")
```

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## Example 4.7 The Rocket Propellant Data (cont.)

#### 2. Detection and treatment of outliers



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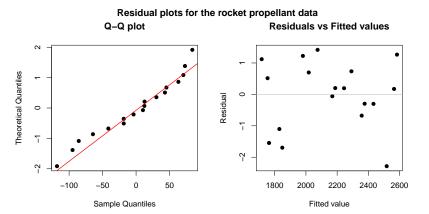
# Example 4.7 The Rocket Propellant Data (cont.)

```
> lmfit <- lm(yi~xi)</pre>
> par(mfrow=c(1,2), cex.main=1.2, pch=19, cex=1.5)
> qqnorm(residuals(lmfit), datax=TRUE, main="Q-Q plot")
> qqline(residuals(lmfit), datax=TRUE, col=2, lwd=2)
> identify(sort(residuals(lmfit)), qnorm(1:length(xi)/length(xi)),
           (1:length(xi))[order(residuals(lmfit))])
> fit_val <- fitted(lmfit)</pre>
> plot(fit_val, scaled(lmfit, "rstudent"), xlab="Fitted value",
       vlab="Residual", main="Residuals vs Fitted values")
> identify(fit_val, scaled(lmfit, "rstudent"), 1:length(xi))
> abline(h=0, lty=1, col="grey")
> title(main="Residual plots for the rocket propellant data",
        line=-1,outer=T)
```

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## Example 4.7 The Rocket Propellant Data (cont.)

#### Observations 5 and 6 are removed



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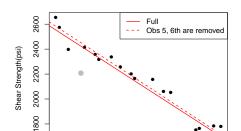
## Example 4.7 The Rocket Propellant Data (cont.)

```
> lmfit <- lm(vi[-c(5,6)]~xi[-c(5,6)])
> par(mfrow=c(1,2), cex.main=1.2, pch=19, cex=1.5)
> qqnorm(residuals(lmfit), datax=TRUE, main="Q-Q plot")
> gqline(residuals(lmfit), datax=TRUE, col=2, lwd=2)
> identify(sort(residuals(lmfit)), qnorm(1:length(xi)/length(xi)),
           (1:length(xi))[order(residuals(lmfit))])
> fit_val <- fitted(lmfit)</pre>
> plot(fit_val, scaled(lmfit, "rstudent"), xlab="Fitted value",
       vlab="Residual", main="Residuals vs Fitted values")
> abline(h=0, lty=1, col="grey")
> title(main="Residual plots for the rocket propellant data",
        line=-1.outer=T)
```

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## Example 4.7 The Rocket Propellant Data (cont.)

5



10

15

Age of Propellant(weeks)

20

Treatment of outliers

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# Example 4.7 The Rocket Propellant Data (cont.)

```
> par(mfrow=c(1,1), pch=16, cex=1.4)
> lmfit <- lm(vi~xi)</pre>
> plot(xi, yi, xlab="Age of Propellant(weeks)",
       vlab="Shear Strength(psi)")
> abline(lmfit, col=2, lwd=2)
> points(xi[5:6], yi[5:6], col="grey", cex=1.5, pch=19)
> lmfit <- lm(yi[-c(5,6)]~xi[-c(5,6)])
> abline(lmfit, col=2, lwd=2, lty=2)
> legend("topright", legend=c("Full", "Obs 5, 6th are removed"),
        col=2, lty=1:2, lwd=2)
```

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## Lack of Fit of the Regression Model

$$\sum_{i=1}^{m} \sum_{j=1}^{n_i} (y_{ij} - \hat{y}_i)^2 = \sum_{i=1}^{m} \sum_{j=1}^{n_i} (y_{ij} - \bar{y}_i)^2 + \sum_{i=1}^{m} \sum_{j=1}^{n_i} (\bar{y}_i - \hat{y}_i)^2$$

$$SS_{RES} = SS_{PE} + SS_{LOF}$$

$$F_0 = \frac{SS_{LOF}/(m-2)}{SS_{PE}/(n-m)} = \frac{MS_{LOF}}{MS_{PE}} \sim F_{(m-2,n-m)}$$

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## Lack of Fit of the Regression Model (cont.)

```
SSpe <- function(model, lof){ #SSpe function
        lmfit <- lm(model)</pre>
        y <- model.response(lmfit$model)</pre>
        x <- factor(lof)
        SSpe <- sum(xtabs(y^2x)-xtabs(y^x)^2/table(x))
        SSres <- sum(residuals(lmfit)^2)
        SSlof <- SSres- SSpe
        out <- matrix(NA, 3, 5)
        colnames(out) <- c("Sum Sq", "Df", "Mean Sq", "F value", "Pr(>F)")
        rownames(out) <- c("SSlof", "SSpe", "SSres")</pre>
        out[,1] <- c(SSlof, SSpe, SSres)</pre>
        out[,2] <- c(length(levels(x))-2, length(x)-length(levels(x)),
                      length(x)-2)
        out[1:2,3] <- out[1:2,1]/out[1:2,2]
        out[1,4] <- out[1,3]/out[2,3]
        out[1,5] <- pf(out[1,4], out[1,2], out[2,2], lower.tail=F)
        printCoefmat(out, digits=4, na.print="")
```

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# Lack of Fit of the Regression Model (cont.)

## 1. Using SSpe function

```
> x < c(1,1,2,3.3,3.3,4,4,4,4.7,5,5.6,5.6,5.6,6,6,6,6.5,6.9)
> y < c(10.84, 9.30, 16.35, 22.88, 24.35, 24.56, 25.86, 29.16, 24.59,
         22.25,25.90,27.2,25.61,25.45,26.56,21.03,21.46)
>
> SSpe(y~x, x)
                   Df Mean Sq F value Pr(>F)
       Sum Sq
SSlof 234.571 8.000
                      29.321
                               13.19 0.00139 **
SSpe
    15.563 7.000
                       2.223
SSres 250.134
               15.000
Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1 1
>
```

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## Lack of Fit of the Regression Model (cont.)

2. Using anova function (restricted method)

> f1 <- lm(y ~x)