

Math 5305 Notes

Diagnostics and Remedial Measures

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Model Assumptions

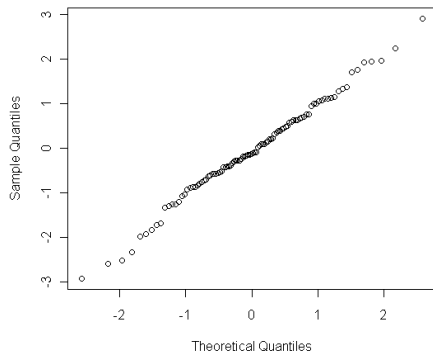
- $Y = X\beta + \epsilon$
- $p < n$ and X has full rank.
- $\epsilon \perp\!\!\!\perp X$
- $\epsilon_1, \dots, \epsilon_n$ are independent
- $E(\epsilon) = 0$
- $\text{Var}(\epsilon_i) = \sigma^2$ for all i
- $\epsilon_1, \dots, \epsilon_n$ are normally distributed

- 1 Error Term Assumptions
- 2 Transformations of Y
- 3 Functional Form
- 4 Model Building

Normality of Errors Diagnostics

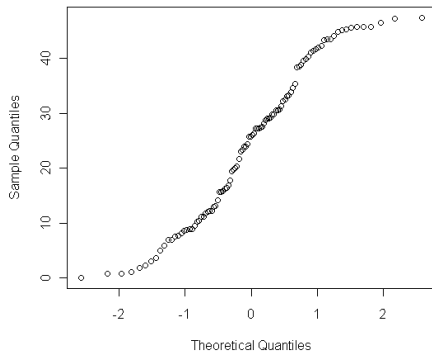
Quantile-Quantile Plot of residuals.

Normal Q-Q Plot



Normally Distributed Residuals

Normal Q-Q Plot



Uniformly Distributed Residuals

R Command: `qqnorm(e)`

Normality of Errors Diagnostics

Shapiro-Wilks Test on Residuals

- Null hypothesis is that the ϵ_j 's are normally distributed.
- Command: `shapiro.test(e)`, where `e` is the vector of residuals.

```
Console ~/ | ↻
> shapiro.test(rnorm(100))

shapiro-wilk normality test

data:  rnorm(100)
W = 0.9844, p-value = 0.2857

> shapiro.test(runif(100))

shapiro-wilk normality test

data:  runif(100)
W = 0.9474, p-value = 0.0005579

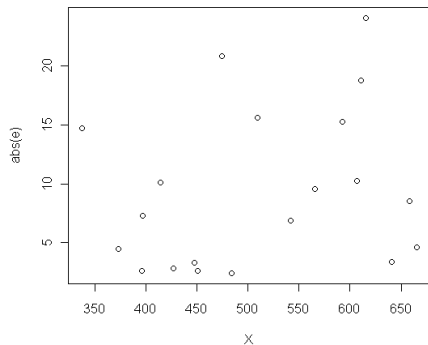
> |
```

- Reject H_0 if p -value is less than α .

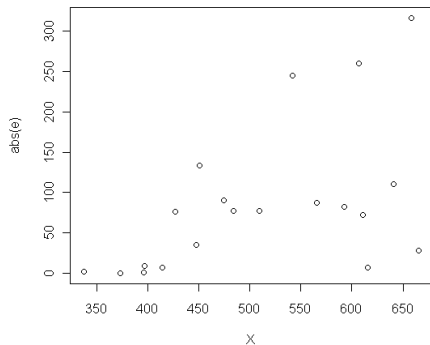
- Transform Y

Constancy of Error Variance Diagnostics

Plot $|e|$ vs. \hat{Y} or X_j .



Constant Error Variance



Nonconstant Error Variance

Brown-Forsythe Test

- The null hypothesis is $\text{Var}(\epsilon_1) = \dots = \text{Var}(\epsilon_n) = \sigma^2$.
- Divide all observations into two groups based on whether \hat{Y} (or X_j) is above or below a certain value.
- Define $e_{i1} = i$ th residual in group 1 and $e_{i2} = i$ th residual in group 2.
- Let n_1 and n_2 be the groups sizes, $n = n_1 + n_2$, and \tilde{e}_1 and \tilde{e}_2 be the medians of the residuals in each group.
- Define $d_{i1} = |e_{i1} - \tilde{e}_1|$ and $d_{i2} = |e_{i2} - \tilde{e}_2|$ for each i .
- Perform a two-sample t -test using the d_{i1} 's and d_{i2} 's.

Brown-Forsythe Test (cont)



$$t = \frac{\bar{d}_1 - \bar{d}_2}{s_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}}$$



$$s_p^2 = \frac{\sum_{i=1}^{n_1} (d_{i1} - \bar{d}_1)^2 + \sum_{i=1}^{n_2} (d_{i2} - \bar{d}_2)^2}{n - 2}$$

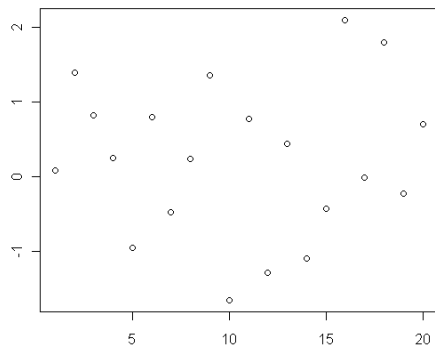
- Reject H_0 if $|t| > t_{\alpha/2}(n - 2)$.

Constancy of Error Variance Remedial Measures

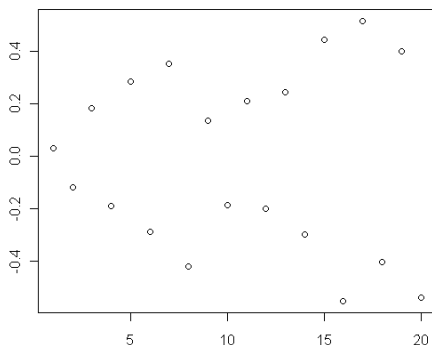
- Transform Y
- Use GLS

Independence of Errors Diagnostics

- Were the data collected in time order?
- Durbin-Watson Test
- Sequence plot: plot $\epsilon_1, \dots, \epsilon_n$ vs. $1, \dots, n$.



No Auto Correlation



Autocorrelated Residuals

Independence of Errors Remedial Measures

- If data were collected in time order, and the Durbin-Watson test/sequence plot show evidence of autocorrelation, use time series analysis.
- If there is a structural reason to believe the ϵ_i 's are dependent, use GLS.

Outline

- 1 Error Term Assumptions
- 2 Transformations of Y**
- 3 Functional Form
- 4 Model Building

Theorem

- Consider an OLS model $Y = X\beta + \epsilon$.
- Ch4 assumptions hold and ϵ is normally distributed.
- Then the maximum likelihood estimators for β and σ^2 are

$$\hat{\beta} = (X'X)^{-1}X'Y \text{ and}$$

$$\hat{\sigma}^2 = \frac{1}{n}\|e\|^2.$$

- If L is the likelihood function, then

$$-2\ln(L(\hat{\beta}, \hat{\sigma}^2)) = n\ln(2\pi) + n\ln(\|e\|^2) - n\ln(n) + n$$

- For linear models with normal disturbance terms, maximizing likelihood is equivalent to minimizing residual sum of squares, $\|e\|^2$.

Transformations of Y

- Problem: error terms not normal or have nonconstant variance.
- Possible solution: transform Y
- Assuming values of Y are **nonnegative**, possible transformations include

$$\tilde{Y}_i = \sqrt{Y_i}$$

$$\tilde{Y}_i = \ln Y_i$$

$$\tilde{Y}_i = \frac{1}{Y_i}$$

- We would then fit the model

$$\tilde{Y} = X\beta + \epsilon$$

Box-Cox Transformations

- Assume Y values are nonnegative. If not, add a constant to all Y values.
- Given a power parameter $\lambda \in \mathbb{R}$, the Box-Cox transformation is

$$\tilde{Y} = \begin{cases} Y^\lambda, & \text{if } \lambda \neq 0 \\ \ln(Y), & \text{if } \lambda = 0 \end{cases}$$

- The model becomes

$$\tilde{Y} = X\beta + \epsilon$$

- λ is estimated with maximum likelihood (least squares).

- Consider a range of values for λ , such as $-2, -1.9, -1.8, \dots, 1.8, 1.9, 2.0$.
- For each value of λ in this range, perform the following steps.
 - ▶ Standardize Y as follows:

$$W_i = \begin{cases} K_1(Y_i^\lambda - 1), & \text{if } \lambda \neq 0 \\ K_2(\ln(Y_i)), & \text{if } \lambda = 0, \end{cases}$$

where

$$K_2 = \left(\prod_{i=1}^n Y_i \right)^{\frac{1}{n}}$$

$$K_1 = \frac{1}{\lambda K_2^{\lambda-1}}.$$

- ▶ Fit the model $W = X\beta + \epsilon$ and compute $\|e\|^2$.
- The value of λ leading to the smallest value of $\|e\|^2$ is the MLE.

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Overall Measures of Fit



$$R^2 = 1 - \frac{\|e\|^2}{\|Y - \bar{Y}\|^2}$$

- Adjusted R^2

$$R_a^2 = 1 - \frac{n-1}{n-p} \frac{\|e\|^2}{\|Y - \bar{Y}\|^2} = 1 - \frac{\hat{\sigma}^2}{\text{Var}(Y)}$$

- Aikake Information Criterion

$$\text{AIC} = 2p - 2\ln(L) = 2p + n\ln(2\pi) + n\ln(\|e\|^2) - n\ln(n) + n$$

- How R calculates AIC for linear models

$$2p + n\ln(\|e\|^2) - n\ln(n)$$

Leave One Out Cross-validation (LOOCV)

- For each $i = 1, \dots, n$, fit a model based on the other observations $1, \dots, i - 1, i + 1, \dots, n$.
- Use this model to predict Y_i , and call this prediction \hat{Y}_i .
- Find the prediction sum of square errors (PRESS)

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

Measures of Fit Based on Cross-validation

Delete- d Cross-validation

- Choose an integer d between 1 and n . A value that has been suggested by Shao (1997) is

$$d = n(1 - (\ln(n) - 1)^{-1}).$$

- Repeat the following process a large number (say 1000) times:
 - Randomly select d rows of the data and remove them.
 - Fit a model to the remaining $n - d$ rows.
 - Use this model to predict the values of Y_i for the removed rows.
 - Find the prediction sum of square errors

$$\text{PRESS} = \sum_{\text{Removed Rows}} (Y_i - \hat{Y}_i)^2.$$

- Finally, average all of these PRESS values to find a single overall PRESS value.

- Diagnostics

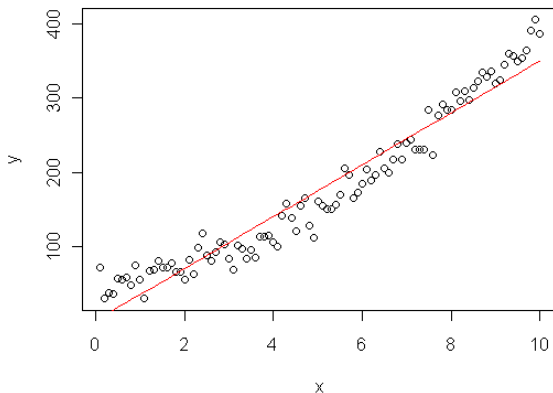
- ▶ Plot Y vs. X_j
- ▶ Plot e vs. \hat{Y} or X_j
- ▶ Compare original model to a model with higher order terms using an F -test or using overall measures of fit.

- Remedial Measures

- ▶ Transform X_j or add higher order terms.

Example Involving Curvature

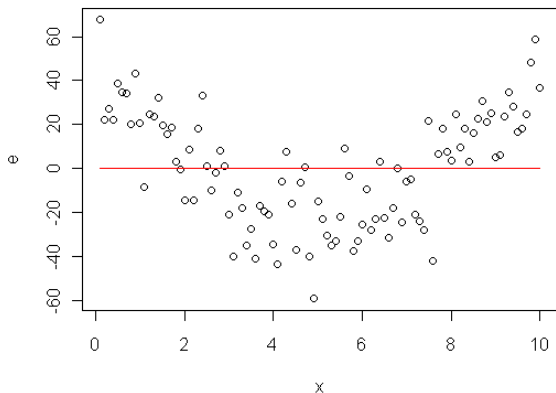
- Scatterplot of Y vs. X



- Do we need higher order terms?

Example Involving Curvature

- Scatterplot of e vs. X

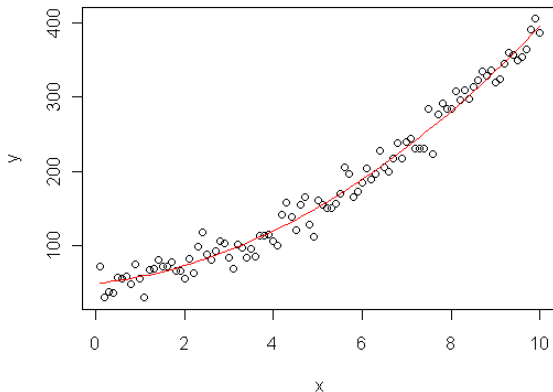


- Trend in residual plot indicates functional form is wrong.

Example Involving Curvature

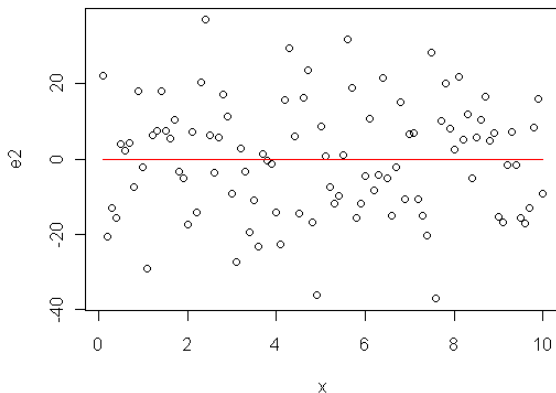
- Fitting quadratic model

$$Y_i = \beta_1 + \beta_2 X_i + \beta_3 X_i^2 + \epsilon_i$$



Example Involving Curvature

- Scatterplot of e vs. X for quadratic model



- Lack of trend in residual plot indicates functional form is right.

- True Model:

$$Y_i = 50 + 5X_i + 3X_i^2 + \epsilon_i$$

- Model 1:

$$Y_i = \beta_1 + \beta_2 X_i + \epsilon_i$$

- Model 2:

$$Y_i = \beta_1 + \beta_2 X_i + \beta_3 X_i^2 + \epsilon_i$$

Two Models

- True Model:

$$Y_i = 50 + 5X_i + 3X_i^2 + \epsilon_i$$

```
Call:
lm(formula = y ~ x)

Residuals:
    Min       1Q   Median       3Q      Max
-59.411 -21.309   0.998  20.750  67.615

Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept)   0.7928     5.2482   0.151   0.88
x             34.8783     0.9022  38.657 <2e-16 ***
---

```

Model 1

```
Call:
lm(formula = y ~ x + I(x^2))

Residuals:
    Min       1Q   Median       3Q      Max
-37.0814 -11.7763   0.8271   9.0966  36.9520

Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept)  49.2248     4.6405  10.608 < 2e-16 ***
x             6.3889     2.1208   3.012  0.00331 **
I(x^2)       2.8207     0.2034  13.865 < 2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Model 2

Comparing the Models with an F -test

- Model 1:

$$Y_i = \beta_1 + \beta_2 X_i + \epsilon_i$$

- Model 2:

$$Y_i = \beta_1 + \beta_2 X_i + \beta_3 X_i^2 + \epsilon_i$$

-

$$\text{ftest}(\text{model1}, \text{model2}) = 9.59 \times 10^{-25}$$

- So, we reject Model 1 in favor of Model 2.

Comparing the Models with Measures of Overall Fit

- R^2 (higher is better)
 - ▶ Model 1: 0.9385
 - ▶ Model 2: 0.9794
- Adjusted R^2 (higher is better)
 - ▶ Model 1: 0.9378
 - ▶ Model 2: 0.9789
- AIC (lower is better)
 - ▶ Model 1: 653.94
 - ▶ Model 2: 546.67
- Leave One Out PRESS (lower is better)
 - ▶ Model 1: 69572.47
 - ▶ Model 2: 23633.27
- SSE ($\|e\|^2$)
 - ▶ Model 1: 66474.03
 - ▶ Model 2: 22293.14

Interaction Terms

- Consider the regression model

$$Y_i = \beta_1 X_{i1} + \cdots + \beta_p X_{ip} + \epsilon_i$$

- An *interaction term* is a term of the form

$$\beta_{j_1 j_2} X_{ij_1} X_{ij_2}$$

Example

- Consider the regression model

$$\text{BloodPressure}_i = \beta_0 + \beta_1 \text{Gender}_i + \beta_2 \text{Cholesterol}_i + \epsilon_i$$

- Here is the same model with an added interaction term for Gender and Cholesterol

$$\begin{aligned} \text{BloodPressure}_i = & \beta_0 + \beta_1 \text{Gender}_i + \beta_2 \text{Cholesterol}_i \\ & + \beta_{12} \text{Gender}_i \text{Cholesterol}_i + \epsilon_i \end{aligned}$$

- Diagnostics
 - ▶ Plot e vs. interaction term.
 - ▶ Compare original model to a model with the interaction term using an F -test or using overall measures of fit.
- Remedial Measures
 - ▶ Include the interaction term if necessary.

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General Guidelines

- Data should be screened for errors.
- Rule of thumb: Sample size should be about 6 to 10 times as large as the number of variables in the pool of potential variables.
- Variables may need to be eliminated if they
 - ▶ are not clinically relevant
 - ▶ have large measurement errors
 - ▶ duplicate other variables
- Clinical considerations should be taken into account. Subject matter experts should be consulted.

Model Building Overview

- Data cleaning/checking
- Split Data into a training sample and a validation sample (this step is not necessary if it is possible to generate new data).
 - ▶ Univariate Analyses
 - ★ Quantitative variables can be checked for curvature.
 - ★ Appropriate categories can be considered for categorical variables.
 - ▶ Variable Selection
 - ▶ Diagnostics and Remedial Measures
- Model Validation: Can be done by comparing model to
 - ▶ New data
 - ▶ Data from the validation sample

Variable Selection Methods

- Manually
- Stepwise Method

```
d=data.frame(Y=Y,X=X)
bigmodel=lm(Y~., data=d)
stepmodel=step(bigmodel)
```

- Best Subsets Method

```
library(bestglm)
Xy=as.data.frame(cbind(x1,x2,x3,x4,x5,x6,Y))
bestglm(Xy,family=gaussian,IC="AIC")
bestglm(Xy,family=gaussian,IC="CV",t=10)
```

- Combination: Use stepwise to narrow the list of variables and then apply best subsets to the remaining variables.

Model Validation

- Models are validated by assessing their performance on a new data set.
- The new data set can actually be newly collected data or can be the validation sample that was set aside at the beginning of model building.
- Diagnostics should be used to determine if the fitted model is consistent with the new data.
- The Mean Squared Prediction Error (MSPE) should be determined
 - ▶ Let $Y_i, i = 1, \dots, n^*$ be the new data set.
 - ▶ For each i , use the model fitted to the training data to predict Y_i .
 - ▶ Call the predicted value \hat{Y}_i .
 - ▶ The MSPE is

$$MSPE = \frac{\sum_{i=1}^{n^*} (Y_i - \hat{Y}_i)^2}{n^*}.$$

Example with Stepwise/Best Subsets Methods

```
X=matrix(runif(5000),100,50)
```

```
X0=cbind(rep(1,100),X[,1:3])
```

```
beta=c(50,5,10,30)
```

```
epsilon=rnorm(100,0,1)
```

```
Y=X0%*%beta+epsilon
```

True Model:

$$Y_i = 50 + 5X_{i1} + 10X_{i2} + 30X_{i3} + \epsilon_i, \text{ for } i = 1, \dots, 100.$$

Variables X_{i4}, \dots, X_{i50} are just noise.

True Model:

$$Y_i = 50 + 5X_{i1} + 10X_{i2} + 30X_{i3} + \epsilon_i, \text{ for } i = 1, \dots, 100.$$

```
x1=X0[,2]
```

```
x2=X0[,3]
```

```
x3=X0[,4]
```

```
model=lm(Y~x1+x2+x3)
```

```
summary(model)
```

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	50.2288	0.3242	154.95	<2e-16	***
x1	4.1198	0.3828	10.76	<2e-16	***
x2	9.9273	0.3749	26.48	<2e-16	***
x3	30.2708	0.3708	81.63	<2e-16	***

```
d=data.frame(Y=Y,X=X)
bigmodel=lm(Y~.,data=d)

stepmodel=step(bigmodel)
summary(stepmodel)
```

```
Coefficients:
Estimate Std. Error t value Pr(>|t|)
(Intercept) 47.6880    0.8455  56.404 < 2e-16 ***
x.1          3.8129    0.3426  11.129 < 2e-16 ***
x.2         10.1911    0.3503  29.095 < 2e-16 ***
x.3         30.6207    0.3688  83.021 < 2e-16 ***
x.4          0.6246    0.3961   1.577  0.11891
x.6          0.7788    0.3333   2.337  0.02203 *
x.14         0.6239    0.3617   1.725  0.08848 .
x.15         0.9363    0.3304   2.834  0.00584 **
x.16        -0.5682    0.3289  -1.727  0.08807 .
x.21        -0.6973    0.3601  -1.936  0.05644 .
x.23         0.6019    0.3580   1.681  0.09671 .
x.27         0.4442    0.3254   1.365  0.17614
x.28        -0.6381    0.3559  -1.793  0.07684 .
x.30         0.9224    0.3210   2.874  0.00522 **
x.35        -0.8150    0.3805  -2.142  0.03532 *
x.37         0.9491    0.3569   2.659  0.00949 **
x.41         0.9769    0.3812   2.563  0.01231 *
x.42         0.9704    0.3344   2.902  0.00481 **
x.43        -1.0106    0.3639  -2.777  0.00686 **
x.47        -0.4945    0.3286  -1.505  0.13642
x.48         0.5735    0.3797   1.511  0.13496
x.50         0.6709    0.3698   1.814  0.07352 .
```



```
Xy=as.data.frame(cbind(X[,c(1,2,3,4,6,14,15,16,21,23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43,47,48,50)],Y))
```

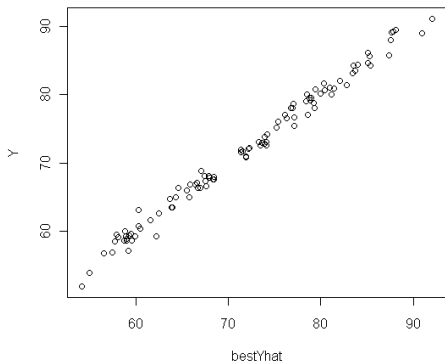
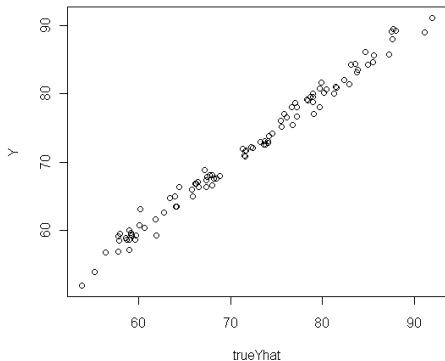
```
bestmodel=bestglm(Xy,IC="CV",family=gaussian,t=10)  
summary(bestmodel$BestModel)
```

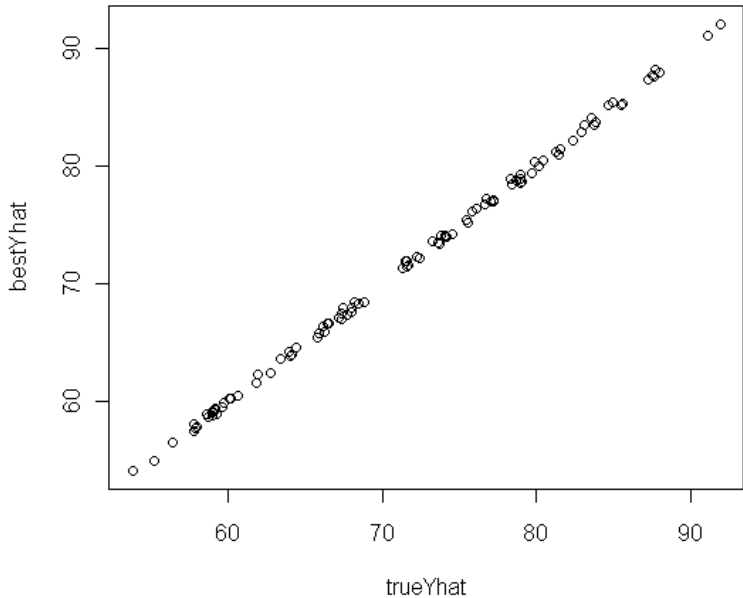
Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	49.7149	0.3706	134.153	<2e-16	***
v1	4.0777	0.3719	10.963	<2e-16	***
v2	9.8833	0.3643	27.128	<2e-16	***
v3	30.4825	0.3689	82.634	<2e-16	***
v15	0.9209	0.3508	2.625	0.0101	*

```
truemodel=lm(Y~x1+x2+x3)
trueYhat=predict(truemodel)
```

```
bestYhat=predict(bestmodel$BestModel)
```





Kutner et. al. (2005). *Applied Linear Statistical Models, 5th ed.*
McGraw-Hill/Irwin, New York, N.Y.