

```
> #read in the data from a csv file
> dat1=read.table("Data Sets/Chapter 3/Examples/ex-3-1.csv", header=T, sep=",")
> str(dat1) #checking data format
```

```
'data.frame':      25 obs. of  4 variables:
 $ Observation      : int   1  2  3  4  5  6  7  8  9 10 ...
 $ Delivery.Time..y : num  16.7 11.5 12.0 14.9 13.8 ...
 $ Number.of.Cases..x1: int   7  3  3  4  6  7  2  7 30 5 ...
 $ Distance..x2..ft. : int  560 220 340 80 150 330 110 210 1460 605 ...
```

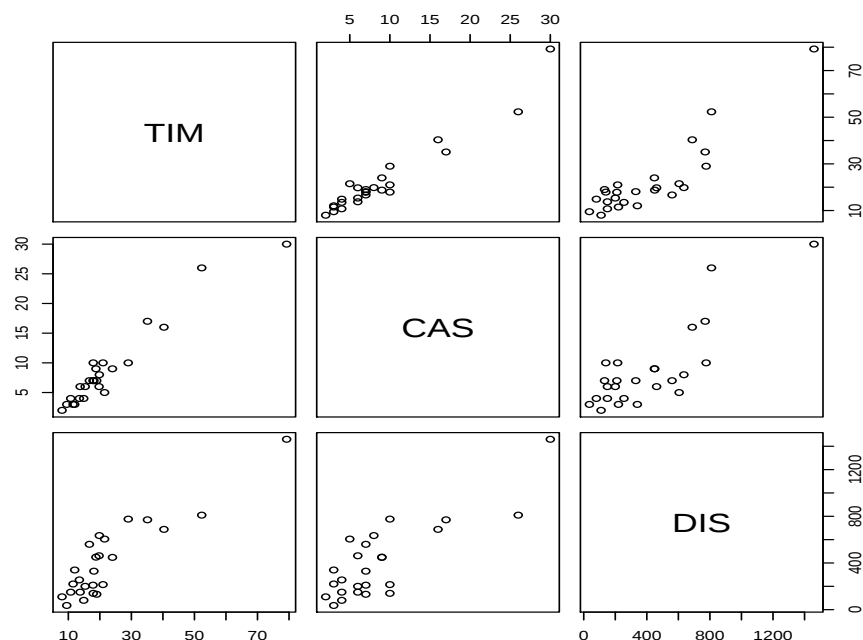
```
> TIM=dat1[,2] #assign the second column in dat1 to be variable TIM
> CAS=dat1[,3] #assign the third column in dat1 to be variable CAS
> DIS=dat1[,4] #assign the forth column to be variable DIS
> EX3=data.frame(TIM, CAS, DIS)
> head(EX3) #printout the first few row in dat1
```

```
      TIM CAS DIS
1 16.68   7 560
2 11.50   3 220
3 12.03   3 340
4 14.88   4  80
5 13.75   6 150
6 18.11   7 330
```

```
> cor(EX3)
```

```
      TIM      CAS      DIS
TIM 1.0000000 0.9646146 0.891670
CAS 0.9646146 1.0000000 0.824215
DIS 0.8916701 0.8242150 1.000000
```

```
> pairs(EX3)
```



```
> library(xtable)
> g=lm(TIM~CAS+DIS, data=EX3)
> xtable(anova(g))
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
CAS	1	5382.41	5382.41	506.62	0.0000
DIS	1	168.40	168.40	15.85	0.0006
Residuals	22	233.73	10.62		

```
> xtable(summary(g))
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	2.3412	1.0967	2.13	0.0442
CAS	1.6159	0.1707	9.46	0.0000
DIS	0.0144	0.0036	3.98	0.0006

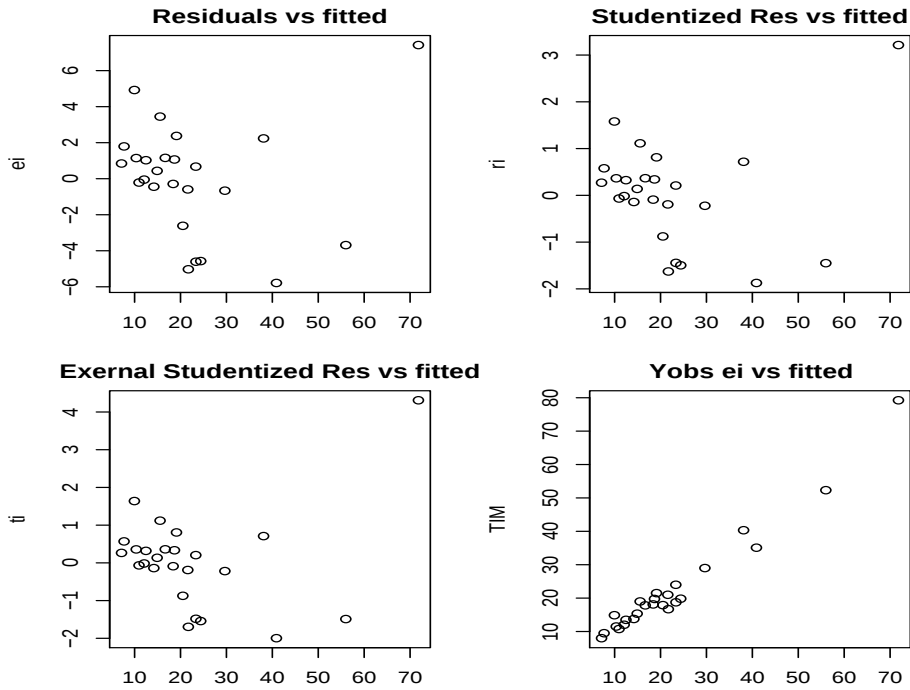
```
> yhat=fitted(g) #fitted values
> ei=resid(g) #residuals
> ri=rstandard(g) #studentized residuals: ei/(s*sqrt(1-hii))
> ti=rstudent(g) #external studentized residuals: ei/(s(i)*sqrt(1-hii))
> hii=hatvalues(g) #hii: diagnoal of H
> out=cbind(TIM, yhat, ei, ri, ti, hii)
> xtable(out) #two decimal places for printout
```

1. (Scaled) Residuals vs fitted values

outliner, constant variance, any patterns. transformation of y or xi weighted least squared estimator

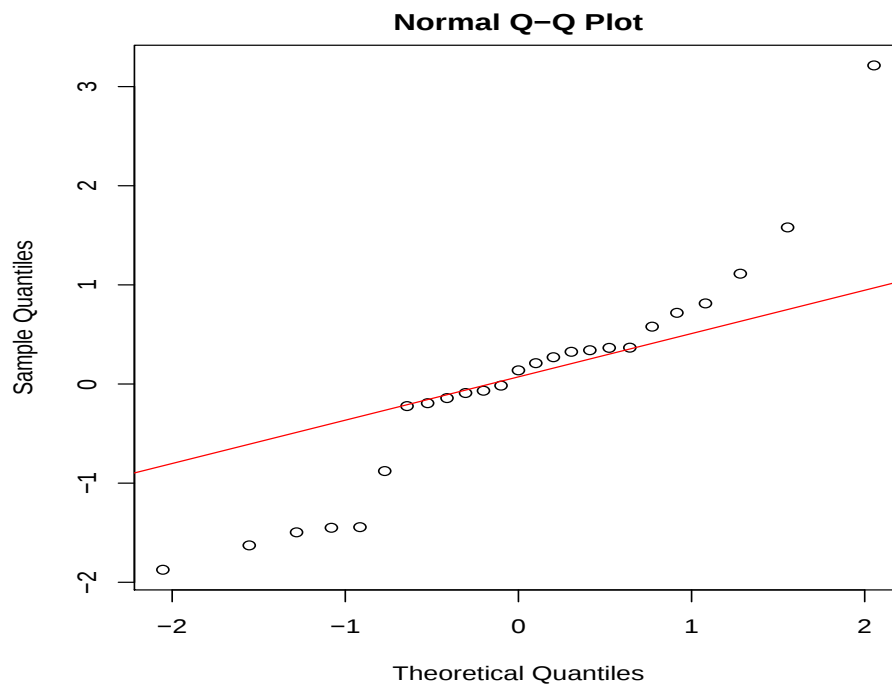
```
> par(mfrow=c(2,2),mar=c(3,4,2,2))
> plot(yhat, ei, main="Residuals vs fitted")
> plot(yhat, ri, main="Studentized Res vs fitted")
> plot(yhat, ti, main="External Studentized Res vs fitted")
> plot(yhat, TIM, main="Yobs ei vs fitted")
```

	TIM	yhat	ei	ri	ti	hii
1	16.68	21.71	-5.03	-1.63	-1.70	0.10
2	11.50	10.35	1.15	0.36	0.36	0.07
3	12.03	12.08	-0.05	-0.02	-0.02	0.10
4	14.88	9.96	4.92	1.58	1.64	0.09
5	13.75	14.19	-0.44	-0.14	-0.14	0.08
6	18.11	18.40	-0.29	-0.09	-0.09	0.04
7	8.00	7.16	0.84	0.27	0.26	0.08
8	17.83	16.67	1.16	0.37	0.36	0.06
9	79.24	71.82	7.42	3.21	4.31	0.50
10	21.50	19.12	2.38	0.81	0.81	0.20
11	40.33	38.09	2.24	0.72	0.71	0.09
12	21.00	21.59	-0.59	-0.19	-0.19	0.11
13	13.50	12.47	1.03	0.33	0.32	0.06
14	19.75	18.68	1.07	0.34	0.33	0.08
15	24.00	23.33	0.67	0.21	0.21	0.04
16	29.00	29.66	-0.66	-0.22	-0.22	0.17
17	15.35	14.91	0.44	0.14	0.13	0.06
18	19.00	15.55	3.45	1.11	1.12	0.10
19	9.50	7.71	1.79	0.58	0.57	0.10
20	35.10	40.89	-5.79	-1.87	-2.00	0.10
21	17.90	20.51	-2.61	-0.88	-0.87	0.17
22	52.32	56.01	-3.69	-1.45	-1.49	0.39
23	18.75	23.36	-4.61	-1.44	-1.48	0.04
24	19.83	24.40	-4.57	-1.50	-1.54	0.12
25	10.75	10.96	-0.21	-0.07	-0.07	0.07

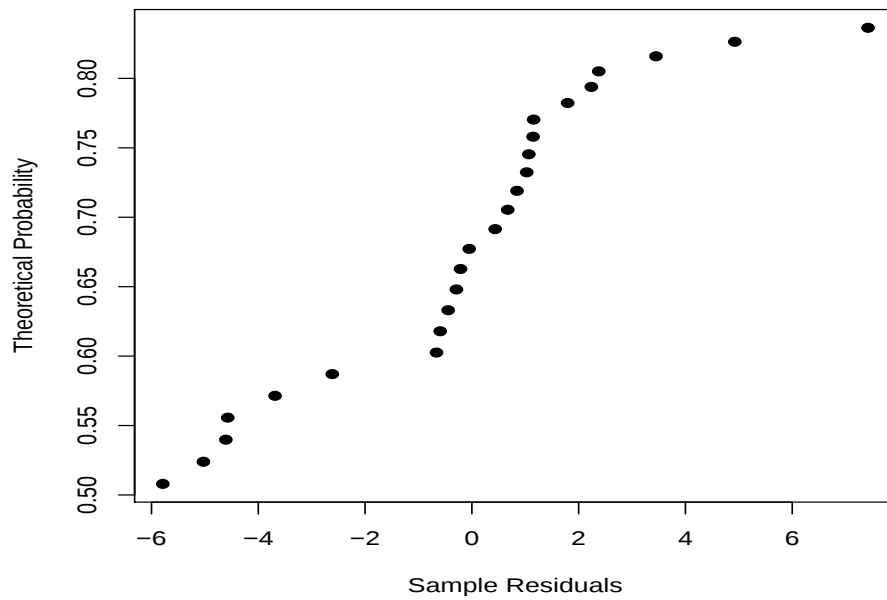


2. Normal Probability Plots: checking if errors follow a normal distribution
Differences between ε_i and e_i ?

```
> par(mfrow=c(1,1),mar=c(5,4,2,2))
> qqnorm(ri);qqline(ri,col=2)
```



```
> my.ppnorm=function(r) {
+   n=length(r)
+   order.res=sort(r)
+   i=1:n
+   pi=pnorm((i-1/2)/n)
+   plot(order.res, pi, xlab = "Sample Residuals", ylab = "Theoretical Probability", pch=16)
+ }
> my.ppnorm(ei)
```



example

Some normal probability plots:

```
> par(mfrow=c(1,1))
> curve(dnorm(x), -4,4)
> curve(dcauchy(x), -4,4, add=TRUE, col=2)
> curve(dt(x,2), -4,4, add=TRUE, col=3)
> curve(dt(x,4), -4,4, add=TRUE, col=3, lty=3)
> curve(dexp(x),0,4, add=TRUE, col=4)
> legend("topright", legend=c("Normal", "Cauchy", "t", "Exp"), lty=1, col=c(1:4))
```

```
> par(mfrow=c(3,3), mar=c(2,2,1,1))
> for (i in 1:9) { e=3+rnorm(50); qqnorm(e, main="");qqline(e,col=2) }
```

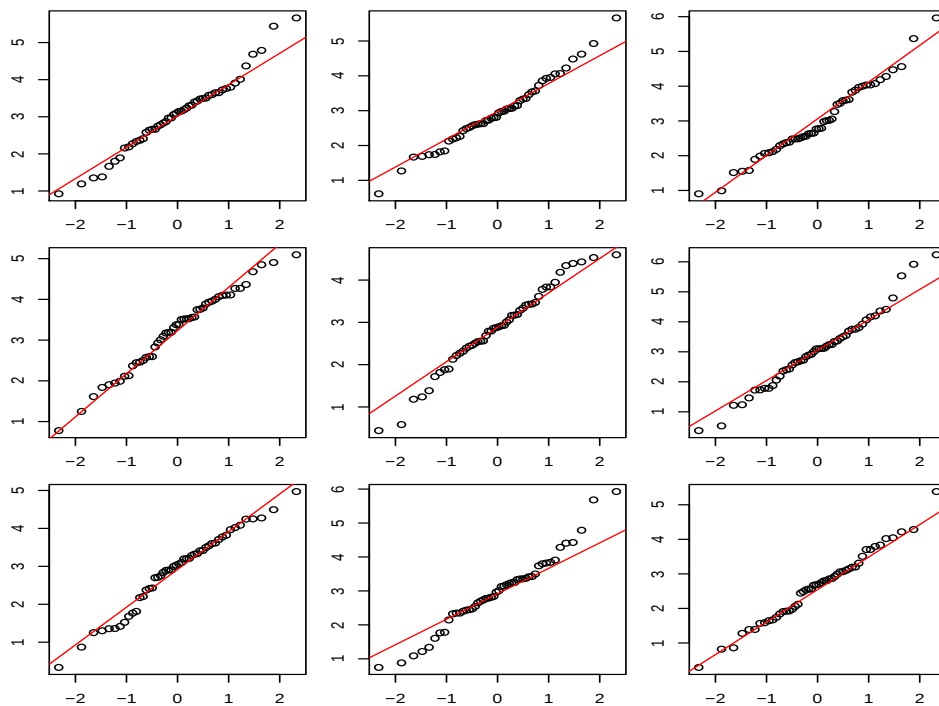
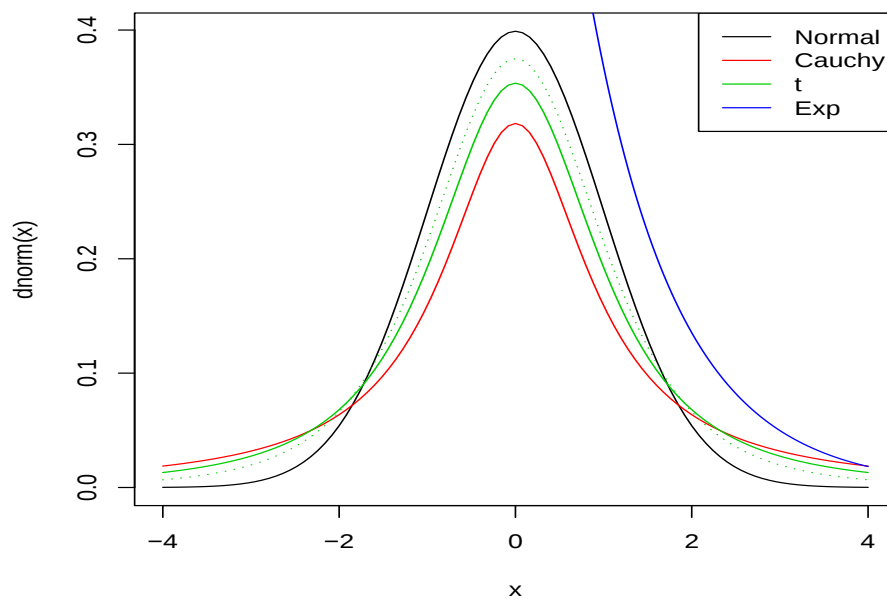


Figure 1: $\text{Normal}(\mu = 3, \text{sd}=1)$



3. Plot of Residuals in Time sequence: autocorrelations

```
> par(mfrow=c(3,3), mar=c(2,2,1,1))
> for (i in 1:9) { e=exp(rnorm(50)); qqnorm(e, main="");qqline(e,col=2) }
```

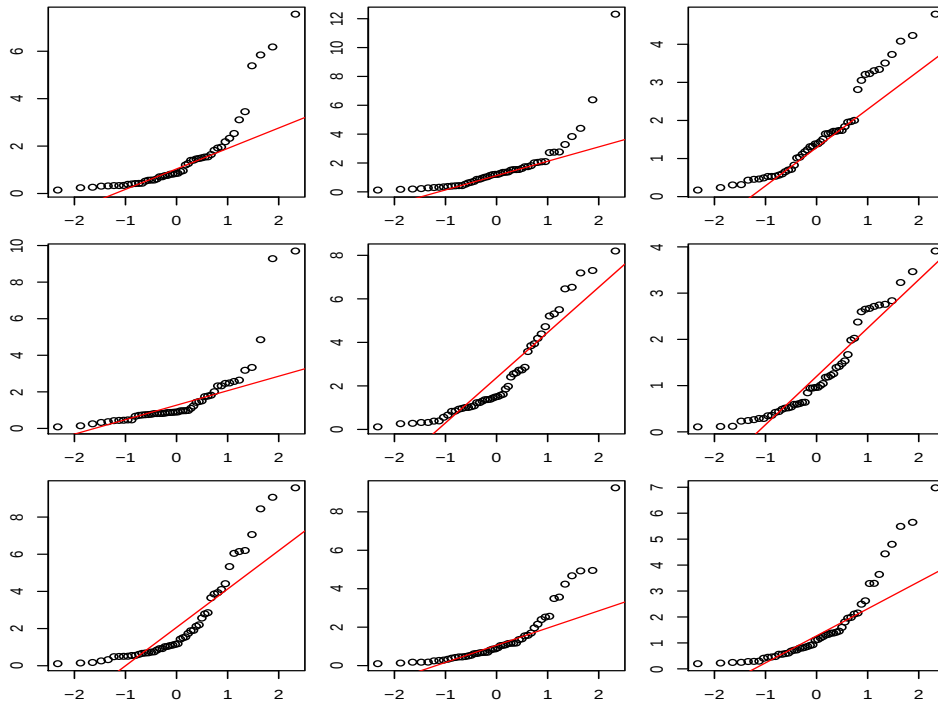


Figure 2: Exponential: right skew

```
> par(mfrow=c(3,3), mar=c(2,2,1,1))
> for (i in 1:9) { e=3+rcauchy(50); qqnorm(e, main="");qqline(e,col=2) }
```

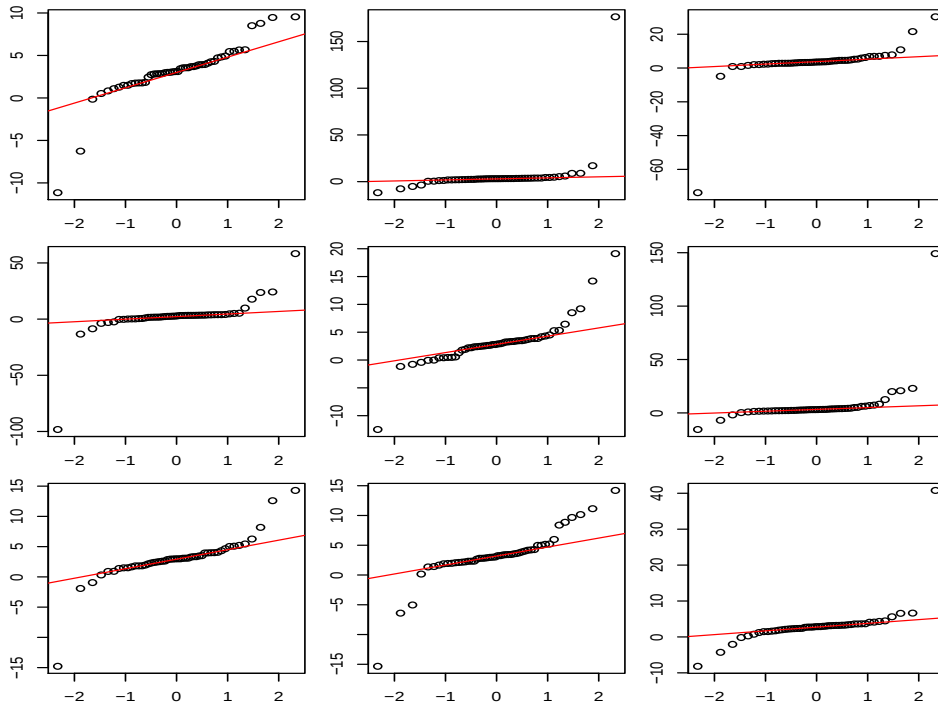


Figure 3: Cauchy: heavy tailed

```
> par(mfrow=c(3,3), mar=c(2,2,1,1))
> for (i in 1:9) { e=runif(50,-1,1); qqnorm(e, main="");qqline(e,col=2) }
```

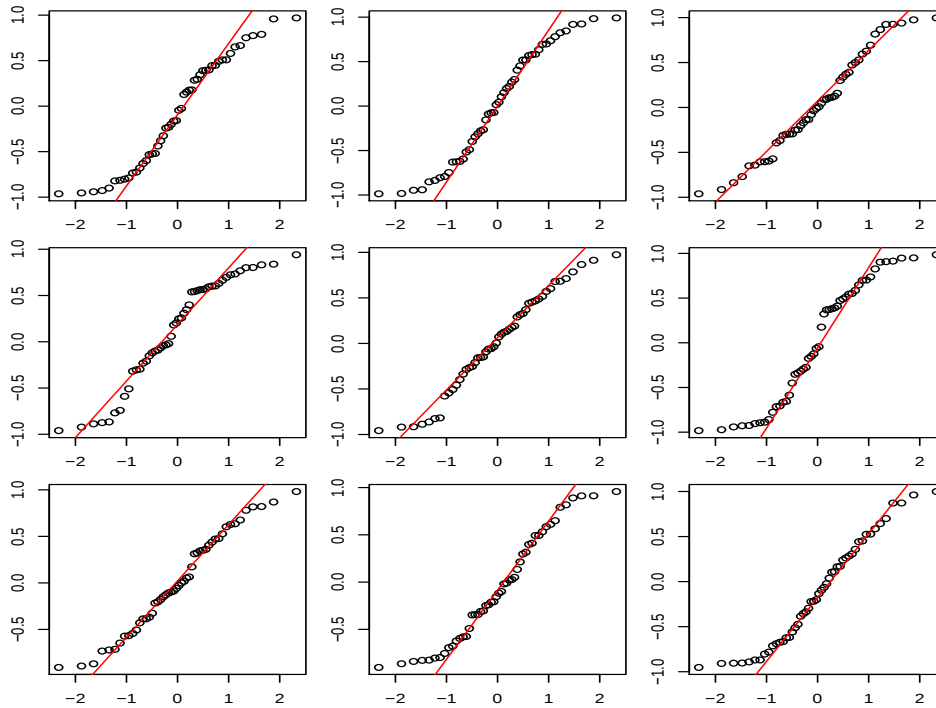


Figure 4: Unifor

4. Partial regression and partial residual plots We can look at plots of e_i against x_i . The drawback is that the other regressors are also in the model, and they may not show the correct marginal effect of a regressor.

- (a) partial regression or added-variable plots can help to isolate the effect of x_i on y .

The relationship between the slope of the partial regression line and the (partial) regression coefficient of x_i in the multiple regression model.

```
> coef(lm(e~d))
```

```
(Intercept)          d
9.981487e-18 1.615907e+00
```

```
> coef(g)
```

```
(Intercept)      CAS      DIS
2.34123115  1.61590721  0.01438483
```

- (b) partial residual plots: $e + \hat{\beta}_j x_j$ against x_j

```
> cr.plot(g, CAS)
```

```

> e=resid(lm(TIM~DIS))
> d=resid(lm(CAS~DIS))
> plot(d, e, xlab="CAS", ylab="TIM Residuals")
> abline(0, coef(g)['CAS'])

```

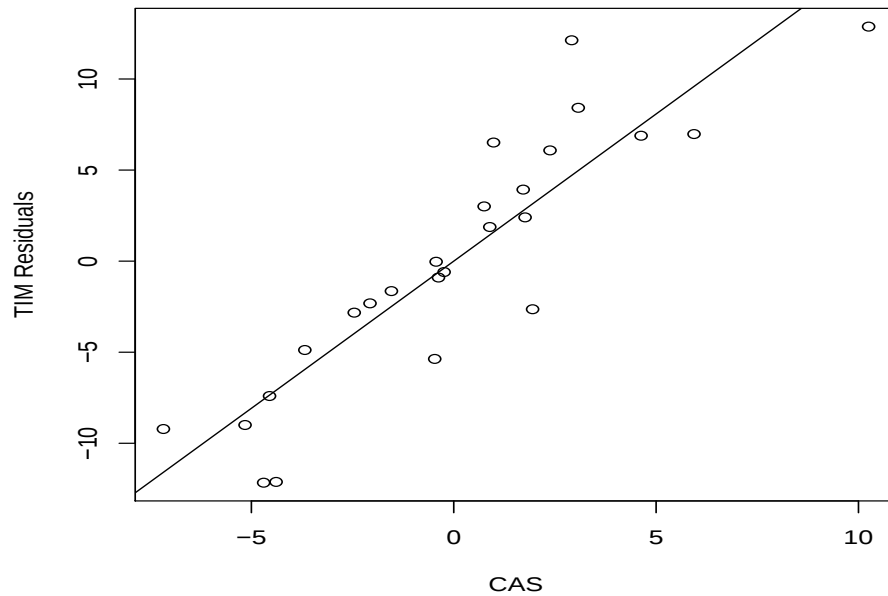


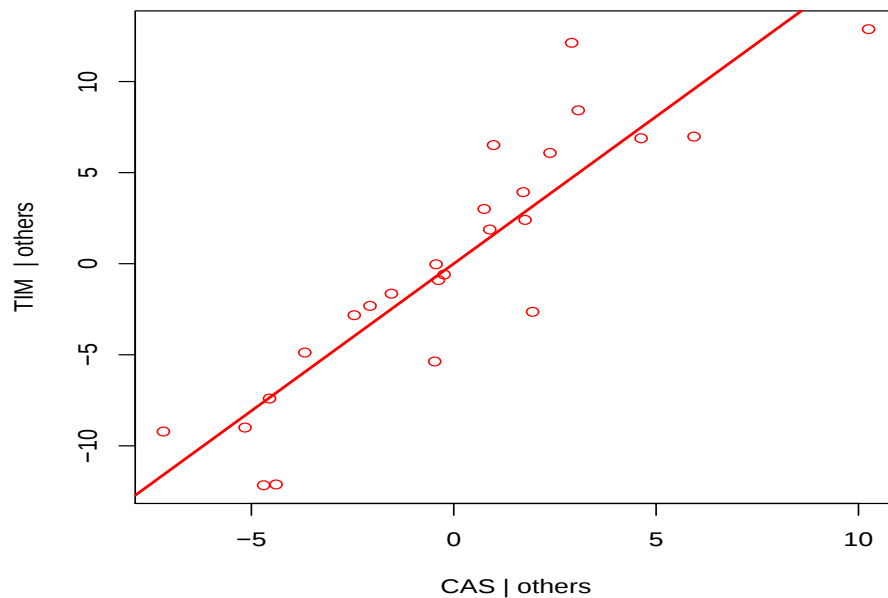
Figure 5: 4.7 Partial regression plot

```

> library(car) #include car library
> av.plot(g, CAS, identify.points=F)

```

Added-Variable Plot



```
> b=coef(g)['CAS']
> plot(CAS, resid(g)+b*CAS, xlab="CAS", ylab="RES+COMPONENT")
```

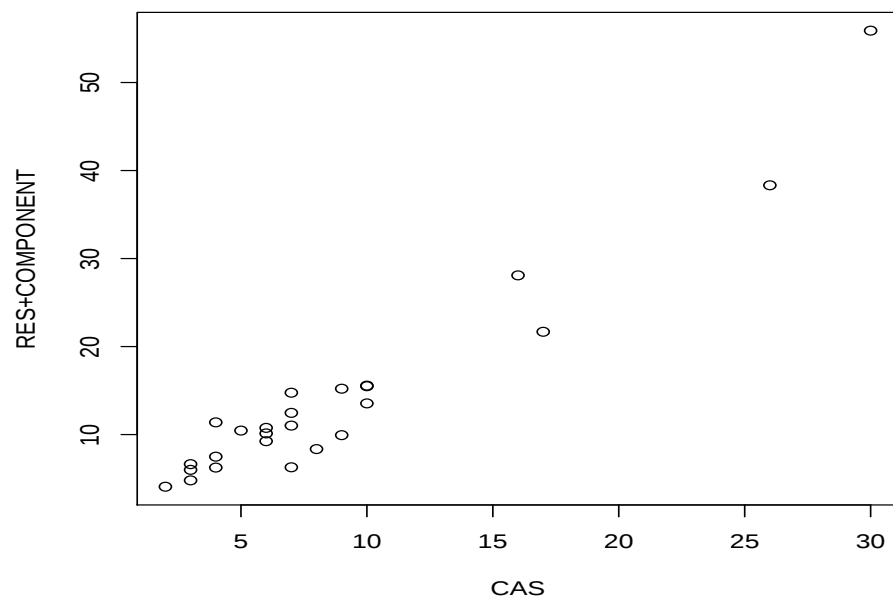
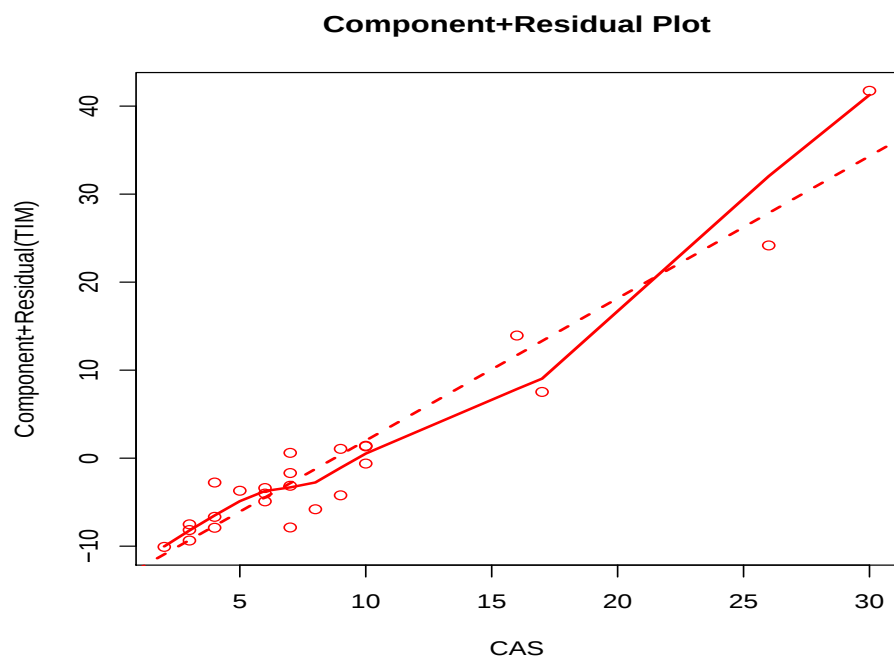


Figure 6: partial residual plot



example: Assume the true model is $y = 2 + \beta_1 x_1^2 + \beta_2 x_2 + \beta_3 x_3 + \varepsilon$ where $\varepsilon \sim N(0, sd = 0.1)$
`runif(20, -1, 1)` : draw 20 random sample for a uniform distribution within (-1, 1)

`rnorm(20, sd=0.1)` : draw 20 random sample for a normal distribution with mean 0 and standard deviation 0.1.

```
> x1=runif(20,-1,1)
> x2=runif(20,-1,1)
> x3=runif(20,-1,1)
> y=2+x2+x3+I(x1^2)+rnorm(20, sd=0.1)
> cor(cbind(y, x1, x2, x3))
```

	y	x1	x2	x3
y	1.0000000	0.32486084	0.5331992	0.61703192
x1	0.3248608	1.00000000	0.3617582	0.04026376
x2	0.5331992	0.36175820	1.0000000	-0.27535328
x3	0.6170319	0.04026376	-0.2753533	1.00000000

Fit a linear model $lm(y \sim x1 + x2 + x3)$.

```
> g=lm(y~x1+x2+x3)
> summary(g)
```

Call:

```
lm(formula = y ~ x1 + x2 + x3)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-0.36572	-0.15625	0.01433	0.16058	0.35339

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	2.31108	0.06083	37.993	< 2e-16
x1	0.02580	0.10524	0.245	0.81
x2	1.13142	0.12292	9.205	8.59e-08
x3	1.07440	0.09957	10.790	9.44e-09

Residual standard error: 0.2368 on 16 degrees of freedom

Multiple R-squared: 0.916, Adjusted R-squared: 0.9002

F-statistic: 58.12 on 3 and 16 DF, p-value: 7.997e-09

```
> anova(g)
```

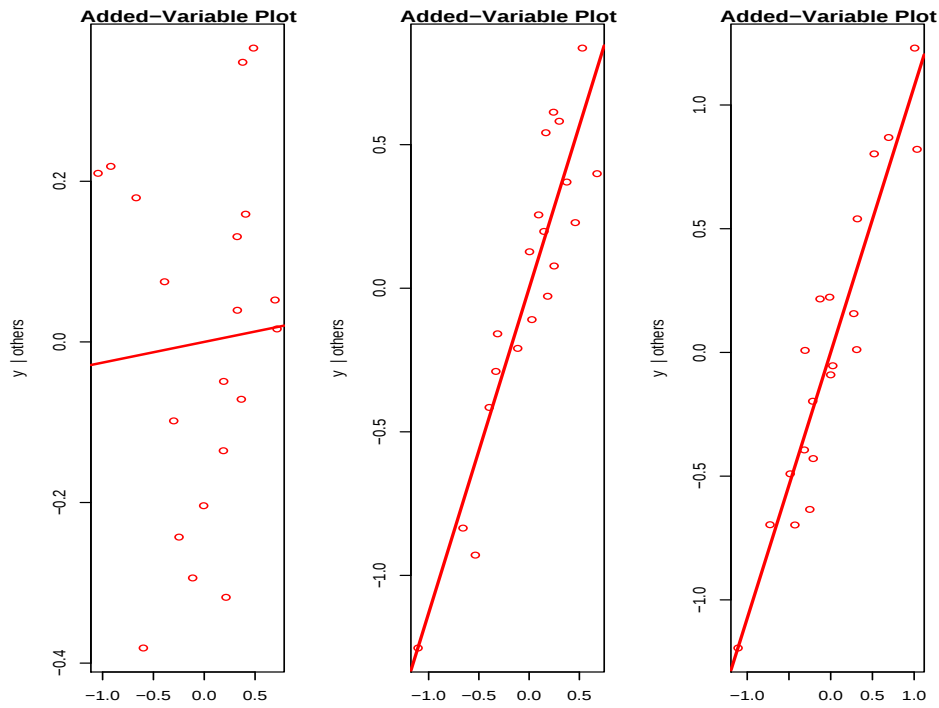
Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
x1	1	1.1270	1.1270	20.090	0.0003772
x2	1	2.1230	2.1230	37.846	1.392e-05
x3	1	6.5313	6.5313	116.430	9.438e-09
Residuals	16	0.8975	0.0561		

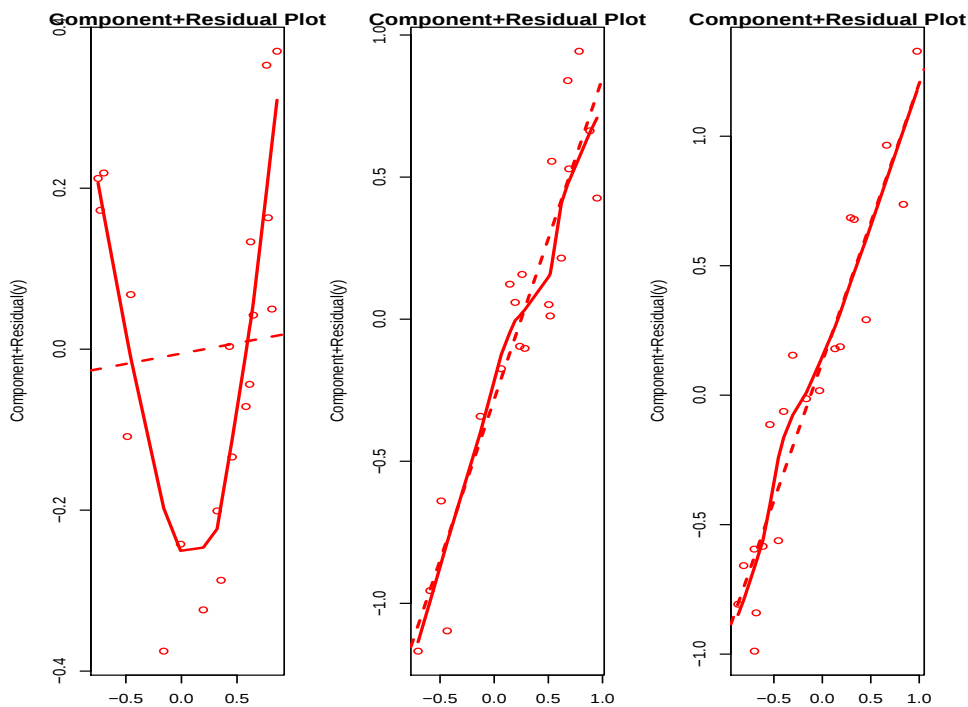
added-variable plots (partial regression plots) are better for outlier detection

```
> par(mfrow=c(1,3),mar=c(3,4,1,2))
> av.plot(g, x1, identify.points=FALSE)
> av.plot(g, x2, identify.points=FALSE)
> av.plot(g, x3, identify.points=FALSE)
```



partial residual plots are better for nonlinearity detection

```
> par(mfrow=c(1,3),mar=c(3,4,1,2))
> cr.plot(g, x1) #partial regression plot
> cr.plot(g, x2)
> cr.plot(g, x3)
```



5. PRESS statistic

```

> sum(ei^2)    #Residual Sum of squares: SSres
[1] 233.7317

> press.ei=ei/(1-hii)
> sum(press.ei^2) #PRESS
[1] 459.0393

> n=length(TIM) #number of obs
> (SST0=(n-1)*var(TIM)) #SST0=Syy
[1] 5784.543

> 1-sum(press.ei^2)/SST0 #prediction R2
[1] 0.9206438

```

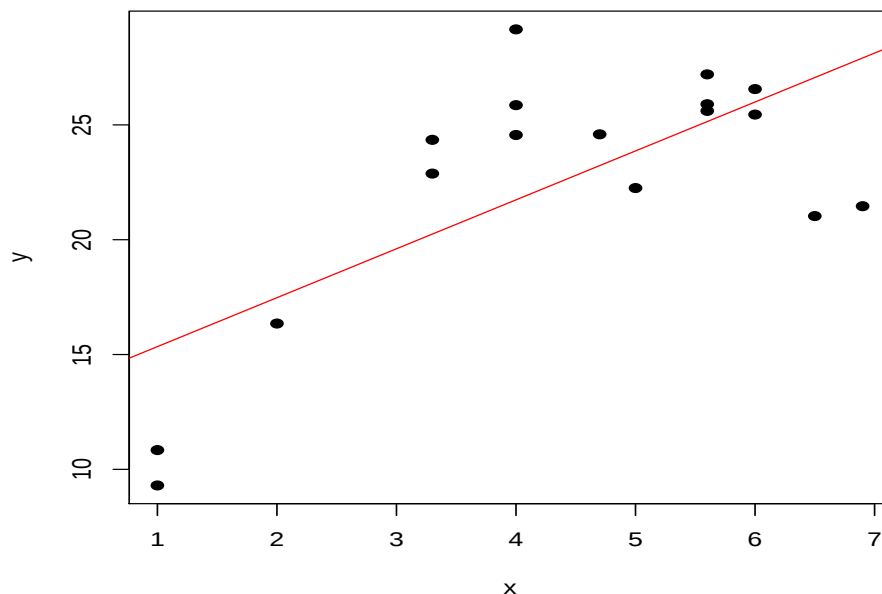
Example 4.8 testing for lack of fit

```

> par(mfrow=c(1,1))
> x=c(1,1,2, 3.3, 3.3, 4,4,4,4.7,5,5.6,5.6,5.6,6,6,6.5,6.9)
> y=c(10.84,9.3,16.35, 22.88, 24.35, 24.56, 25.86, 29.16,24.59,22.25,25.9, 27.2,
+ 25.61,25.45,26.56, 21.03, 21.46)
> print(rbind(x,y))
      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10] [,11] [,12]
x  1.00  1.0  2.00  3.30  3.30  4.00  4.00  4.00  4.70  5.00  5.6  5.6
y 10.84  9.3 16.35 22.88 24.35 24.56 25.86 29.16 24.59 22.25 25.9 27.2
      [,13] [,14] [,15] [,16] [,17]
x  5.60  6.00  6.00  6.50  6.90
y 25.61 25.45 26.56 21.03 21.46

> plot(x,y, pch=16)
> g=lm(y~x)
> abline(coef(g), col=2)

```



```
> xtable(summary(g)) #test for the significance of regression?
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	13.2139	2.6649	4.96	0.0002
x	2.1304	0.5645	3.77	0.0018

```
> xtable(anova(g))
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
x	1	237.48	237.48	14.24	0.0018
Residuals	15	250.13	16.68		

Use the function factor to encode a vector x as a factor X .

```
> X=factor(x) #re-assigne x as a factor X
```

```
> levels(X) #level for the factor X
```

```
[1] "1" "2" "3.3" "4" "4.7" "5" "5.6" "6" "6.5" "6.9"
```

```
> length(levels(X))
```

```
[1] 10
```

```
> f=lm(y~X)
```

```
> xtable(anova(f))
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
X	9	472.05	52.45	23.59	0.0002
Residuals	7	15.56	2.22		

Test for lack of fit

```
> anova(g,f)
```

Analysis of Variance Table

Model 1: $y \sim x$

Model 2: $y \sim X$

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	15	250.134				
2	7	15.563	8	234.571	13.188	0.001389