RTSM_Error_analysis.R

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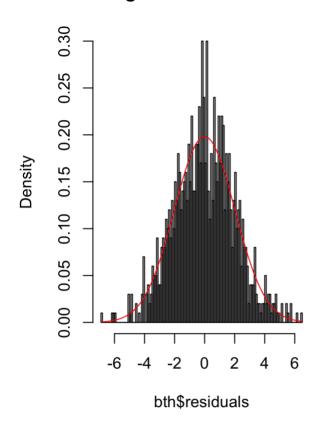
```
# Model adequacy checking
library("sde")
## Loading required package: MASS
## Loading required package: stats4
## Loading required package: fda
## Loading required package: splines
## Loading required package: Matrix
##
## Attaching package: 'fda'
## The following object is masked from 'package:graphics':
##
       matplot
## Loading required package: zoo
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
       as.Date, as.Date.numeric
## sde 2.0.15
## Companion package to the book
## 'Simulation and Inference for Stochastic Differential Equations With R Examples'
## Iacus, Springer NY, (2008)
## To check the errata corrige of the book, type vignette("sde.errata")
```

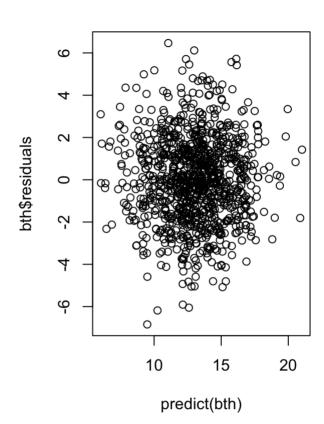
```
############
                   MODEL 1
                               ###########
k<-3 # independent variables
n<- 1000 # number of observations
X < -array(0, dim = c(n, (k+1)))
bt < -c(1,2.3, 1.5,0.05)
x1 < -rgamma(n, 2, 3)
x2 < -rbinom(n, 10, 0.7)
x3 < -rbeta(n, 0.5, 0.5)
X[,1] < -1
X[,2]<-x1
X[,3]<-x2
X[,4]<-x3
sigma<-2
eps<-rnorm(n,0,sigma)
y<-X%*%bt+eps
d<- data.frame(y,X[,2],X[,3],X[,4])</pre>
bth < -lm(y \sim X[,2] + X[,3] + X[,4], data = d)
cat("trur beta=",bt, "sigma=",sigma,'\n')
```

```
## trur beta= 1 2.3 1.5 0.05 sigma= 2
```

```
## beta_hat= 1.216331 2.280402 1.45387 0.2396212 sigma_hat= 2.016307
```

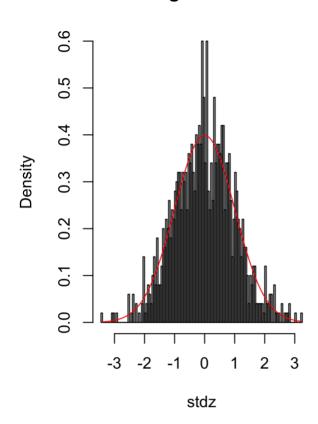
```
par(mfrow=c(1,2))
hist(bth$residuals, breaks = 100, probability = T )
s<-seq(min(bth$residuals), max(bth$residuals), length=100)
lines(dnorm(s, mean(bth$residuals), sd(bth$residuals))~s, col=2)
plot(predict(bth),bth$residuals)</pre>
```

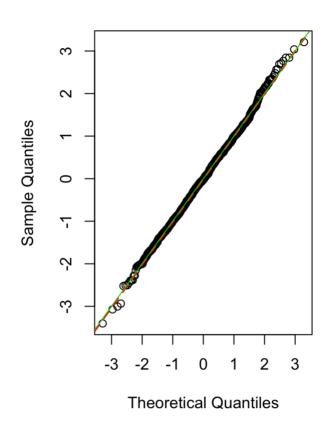




```
#qqnorm(y = bth$residuals, plot.it = T )
#qqline(y=bth$residuals, col=2, lwd=2,lty=2)

stdz<-(stdres(bth))
hist(stdz, breaks = 100, probability = T )
s<-seq(min(stdz), max(stdz), length=100)
lines(dnorm(s, 0,1)~s, col=2)
qqnorm(y = stdz, plot.it = T )
qqline(y=stdz, col=2, lwd=2,lty=2)
abline(a = 0,b = 1, col=3)</pre>
```





```
print(ks.test(x=bth$residuals, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: bth$residuals
## D = 0.16961, p-value < 2.2e-16
## alternative hypothesis: two-sided</pre>
```

```
print(ks.test(x=stdz, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: stdz
## D = 0.016154, p-value = 0.9566
## alternative hypothesis: two-sided
```

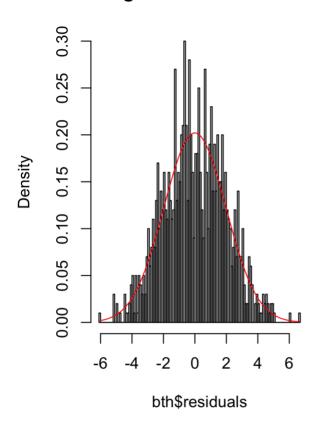
```
MODEL 2 # Normal with non zero mean
############
                                                                ###########
k<-3 # independent variables
n<- 1000 # number of observations
X < -array(0, dim = c(n, (k+1)))
bt < -c(1,2.3, 1.5,0.05)
x1 < -rgamma(n, 2, 3)
x2 < -rbinom(n, 10, 0.7)
x3 < -rbeta(n, 0.5, 0.5)
X[,1] < -1
X[,2]<-x1
X[,3]<-x2
X[,4]<-x3
sigma<-2
eps<-rnorm(n,3,sigma)
y<-X%*%bt+eps
d<- data.frame(y,X[,2],X[,3],X[,4])</pre>
bth < -lm(y \sim X[,2] + X[,3] + X[,4], data = d)
cat("trur beta=",bt, "sigma=",sigma,'\n')
```

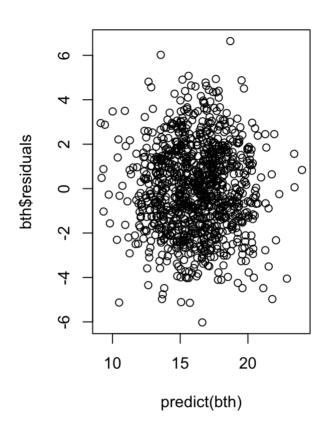
```
## trur beta= 1 2.3 1.5 0.05 sigma= 2
```

 $\verb|cat("beta_hat=",coefficients(bth), "sigma_hat=",summary(bth)$| sigma,'\n'|| \# model coefficients|| ficients||$

```
## beta_hat= 4.292785 2.394685 1.461973 0.02515 sigma_hat= 1.977013
```

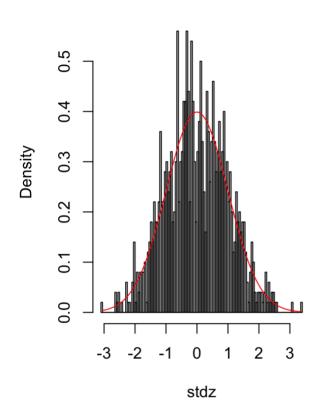
```
par(mfrow=c(1,2))
hist(bth$residuals, breaks = 100, probability = T )
s<-seq(min(bth$residuals), max(bth$residuals), length=100)
lines(dnorm(s, mean(bth$residuals), sd(bth$residuals))~s, col=2)
plot(predict(bth),bth$residuals)</pre>
```

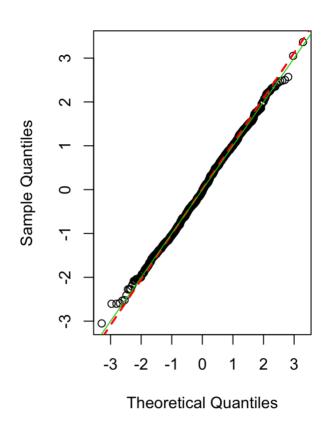




```
# qqnorm(y = bth$residuals, plot.it = T )
# qqline(y=bth$residuals, col=2, lwd=2,lty=2)

stdz<-(stdres(bth))
hist(stdz, breaks = 100, probability = T )
s<-seq(min(stdz), max(stdz), length=100)
lines(dnorm(s, 0,1)~s, col=2)
qqnorm(y = stdz, plot.it = T )
qqline(y=stdz, col=2, lwd=2,lty=2)
abline(a = 0,b = 1, col=3)</pre>
```





```
print(ks.test(x=bth$residuals, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: bth$residuals
## D = 0.17472, p-value < 2.2e-16
## alternative hypothesis: two-sided</pre>
```

```
print(ks.test(x=stdz, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: stdz
## D = 0.023654, p-value = 0.6305
## alternative hypothesis: two-sided
```

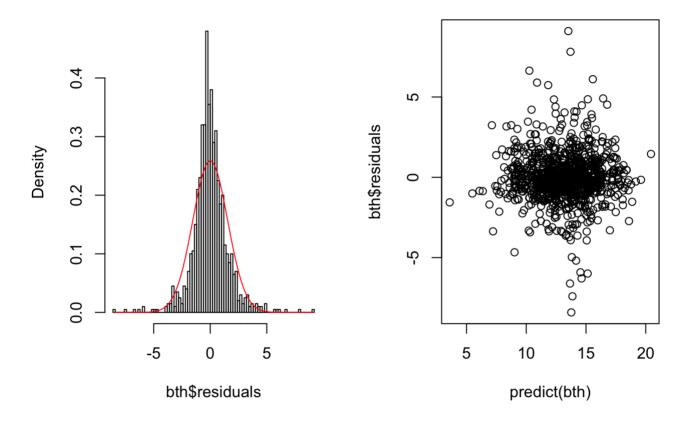
```
############
                  MODEL 3 # Heavy tailed distribution
                                                              ##########
k<-3 # independent variables
n<- 1000 # number of observations
X < -array(0, dim = c(n, (k+1)))
bt < -c(1,2.3, 1.5,0.05)
x1 < -rgamma(n, 2, 3)
x2 < -rbinom(n, 10, 0.7)
x3 < -rbeta(n, 0.5, 0.5)
X[,1] < -1
X[,2]<-x1
X[,3]<-x2
X[,4]<-x3
sigma<-2
eps<-rt(n,df = 3) # Heavy tailed distribution</pre>
y<-X%*%bt+eps
d<- data.frame(y,X[,2],X[,3],X[,4])</pre>
bth < -lm(y \sim X[,2] + X[,3] + X[,4], data = d)
cat("trur beta=",bt, "sigma=",sigma,'\n')
```

```
## trur beta= 1 2.3 1.5 0.05 sigma= 2
```

cat("beta_hat=",coefficients(bth), "sigma_hat=",summary(bth)\$sigma,'\n') # model coef
ficients

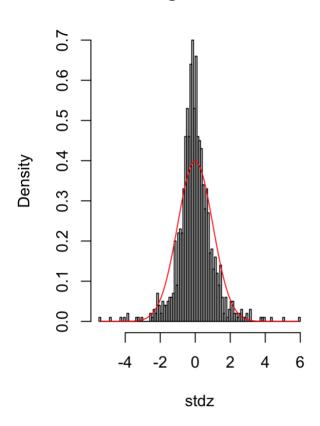
```
## beta_hat= 1.215887 2.22745 1.488538 -0.004132992 sigma_hat= 1.544559
```

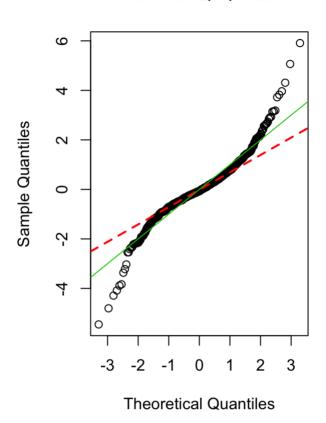
```
par(mfrow=c(1,2))
hist(bth$residuals, breaks = 100, probability = T )
s<-seq(min(bth$residuals), max(bth$residuals), length=100)
lines(dnorm(s, mean(bth$residuals),sd(bth$residuals))~s, col=2)
plot(predict(bth),bth$residuals)</pre>
```



```
#qqnorm(y = bth$residuals, plot.it = T )
#qqline(y=bth$residuals, col=2, lwd=2,lty=2)

stdz<-(stdres(bth))
hist(stdz, breaks = 100, probability = T )
s<-seq(min(stdz), max(stdz), length=100)
lines(dnorm(s, 0,1)~s, col=2)
qqnorm(y = stdz, plot.it = T )
qqline(y=stdz, col=2, lwd=2,lty=2)
abline(a = 0,b = 1, col=3)</pre>
```





```
print(ks.test(x=bth$residuals, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: bth$residuals
## D = 0.059392, p-value = 0.001727
## alternative hypothesis: two-sided
```

```
print(ks.test(x=stdz, y="pnorm",alternative = c("two.sided")))
```

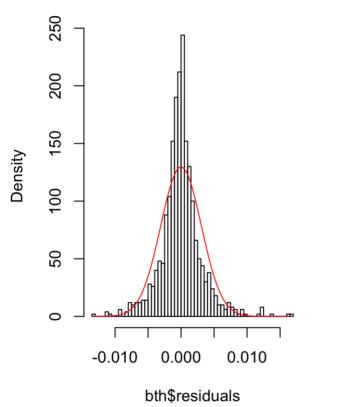
```
##
## One-sample Kolmogorov-Smirnov test
##
## data: stdz
## D = 0.080359, p-value = 4.92e-06
## alternative hypothesis: two-sided
```

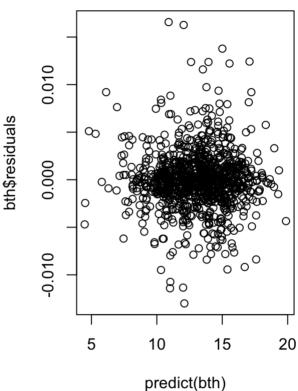
```
############
                  MODEL 4 # thin tailed distribution
                                                            ###########
k<-3 # independent variables
n<- 1000 # number of observations
X < -array(0, dim = c(n, (k+1)))
bt < -c(1,2.3, 1.5,0.05)
x1 < -rgamma(n, 2, 3)
x2 < -rbinom(n, 10, 0.7)
x3 < -rbeta(n, 0.5, 0.5)
X[,1] < -1
X[,2]<-x1
X[,3]<-x2
X[,4]<-x3
sigma<-2
eps<-rexp(n,rate =500)*(rbinom(n,1,0.5)-0.5)*2 # thin tailed distribution
y<-X%*%bt+eps
d<- data.frame(y,X[,2],X[,3],X[,4])</pre>
bth < -lm(y \sim X[,2] + X[,3] + X[,4], data = d)
cat("trur beta=",bt, "sigma=",sigma,'\n')
```

```
## trur beta= 1 2.3 1.5 0.05 sigma= 2
```

```
## beta_hat= 0.9997061 2.300246 1.500031 0.04999736 sigma_hat= 0.003082326
```

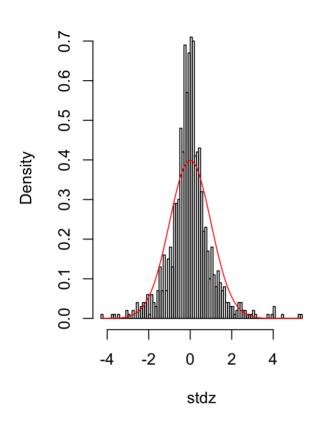
```
par(mfrow=c(1,2))
hist(bth$residuals, breaks = 100, probability = T )
s<-seq(min(bth$residuals), max(bth$residuals), length=100)
lines(dnorm(s, mean(bth$residuals),sd(bth$residuals))~s, col=2)
plot(predict(bth),bth$residuals)</pre>
```

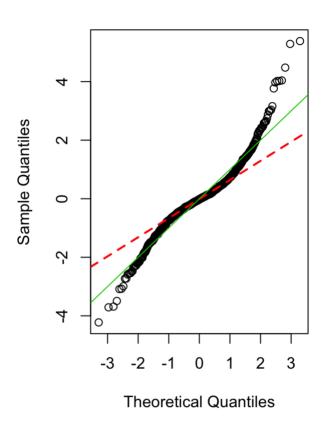




```
#qqnorm(y = bth$residuals, plot.it = T )
#qqline(y=bth$residuals, col=2, lwd=2,lty=2)

stdz<-(stdres(bth))
hist(stdz, breaks = 100, probability = T )
s<-seq(min(stdz), max(stdz), length=100)
lines(dnorm(s, 0,1)~s, col=2)
qqnorm(y = stdz, plot.it = T )
qqline(y=stdz, col=2, lwd=2,lty=2)
abline(a = 0,b = 1, col=3)</pre>
```





```
print(ks.test(x=bth$residuals, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: bth$residuals
## D = 0.49481, p-value < 2.2e-16
## alternative hypothesis: two-sided</pre>
```

```
print(ks.test(x=stdz, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: stdz
## D = 0.090302, p-value = 1.653e-07
## alternative hypothesis: two-sided
```

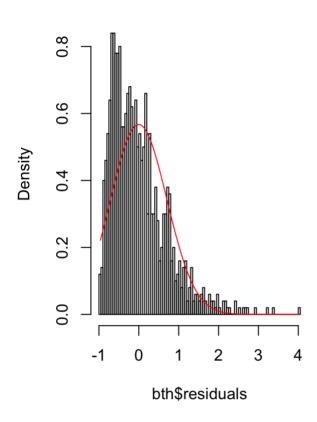
```
############
                  MODEL 5 # Asymmtric distribution #########
k<-3 # independent variables
n<- 1000 # number of observations
X < -array(0, dim = c(n, (k+1)))
bt < -c(1,2.3, 1.5,0.05)
x1 < -rgamma(n, 2, 3)
x2 < -rbinom(n, 10, 0.7)
x3 < -rbeta(n, 0.5, 0.5)
X[,1] < -1
X[,2]<-x1
X[,3]<-x2
X[,4]<-x3
sigma<-2
eps<-rgamma(n,shape = 2, rate = sigma) # positively skewed</pre>
y<-X%*%bt+eps
d<- data.frame(y,X[,2],X[,3],X[,4])</pre>
bth < -lm(y \sim X[,2] + X[,3] + X[,4], data = d)
cat("trur beta=",bt, "sigma=",sigma,'\n')
```

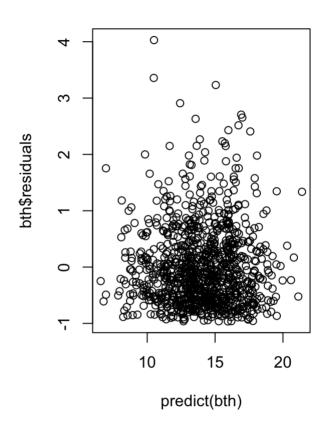
```
## trur beta= 1 2.3 1.5 0.05 sigma= 2
```

cat("beta_hat=",coefficients(bth), "sigma_hat=",summary(bth)\$sigma,'\n') # model coef
ficients

```
## beta_hat= 1.922114 2.284543 1.511539 0.02955286 sigma_hat= 0.7044242
```

```
par(mfrow=c(1,2))
hist(bth$residuals, breaks = 100, probability = T )
s<-seq(min(bth$residuals), max(bth$residuals), length=100)
lines(dnorm(s, mean(bth$residuals),sd(bth$residuals))~s, col=2)
plot(predict(bth),bth$residuals)</pre>
```



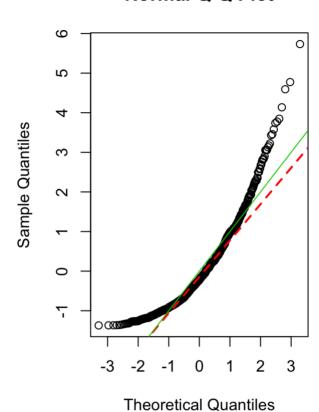


```
#qqnorm(y = bth$residuals, plot.it = T )
#qqline(y=bth$residuals, col=2, lwd=2,lty=2)

stdz<-(stdres(bth))
hist(stdz, breaks = 100, probability = T )
s<-seq(min(stdz), max(stdz), length=100)
lines(dnorm(s, 0,1)~s, col=2)
qqnorm(y = stdz, plot.it = T )
qqline(y=stdz, col=2, lwd=2,lty=2)
abline(a = 0,b = 1, col=3)</pre>
```

Density 0.0 0.1 0.2 0.3 0.4 0.5 0.6

stdz



```
print(ks.test(x=bth$residuals, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: bth$residuals
## D = 0.17285, p-value < 2.2e-16
## alternative hypothesis: two-sided</pre>
```

```
print(ks.test(x=stdz, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: stdz
## D = 0.09405, p-value = 4.15e-08
## alternative hypothesis: two-sided
```

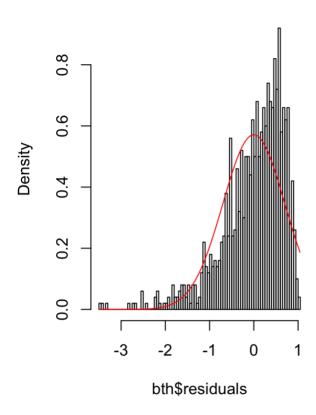
```
############
                  MODEL 6 # Asymmtric distribution #########
k<-3 # independent variables
n<- 1000 # number of observations
X < -array(0, dim = c(n, (k+1)))
bt < -c(1,2.3, 1.5,0.05)
x1 < -rgamma(n, 2, 3)
x2 < -rbinom(n, 10, 0.7)
x3 < -rbeta(n, 0.5, 0.5)
X[,1] < -1
X[,2]<-x1
X[,3]<-x2
X[,4]<-x3
sigma<-2
eps<- rgamma(n,shape = 2, rate = sigma)
y<-X%*%bt-eps # negatively skewed
d<- data.frame(y,X[,2],X[,3],X[,4])</pre>
bth < -lm(y \sim X[,2] + X[,3] + X[,4], data = d)
cat("trur beta=",bt, "sigma=",sigma,'\n')
```

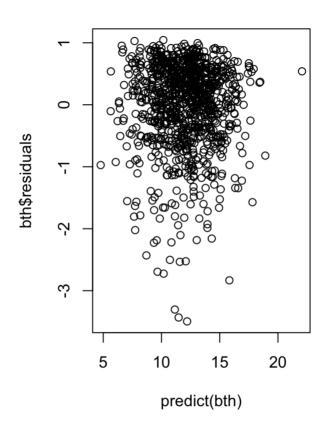
```
## trur beta= 1 2.3 1.5 0.05 sigma= 2
```

cat("beta_hat=",coefficients(bth), "sigma_hat=",summary(bth)\$sigma,'\n') # model coef
ficients

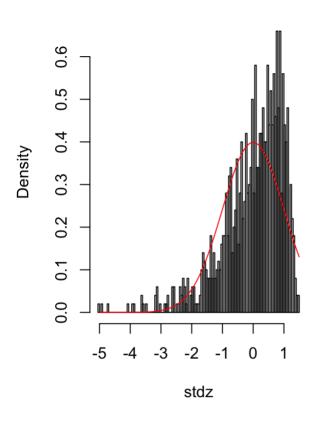
```
## beta_hat= -0.1056956 2.35867 1.519239 -0.04961164 sigma_hat= 0.6994423
```

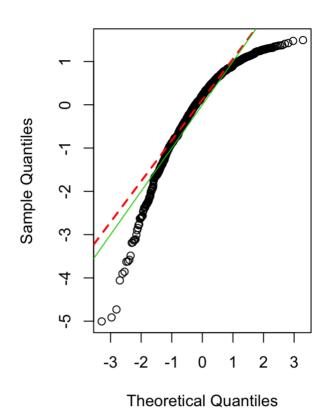
```
par(mfrow=c(1,2))
hist(bth$residuals, breaks = 100, probability = T )
s<-seq(min(bth$residuals), max(bth$residuals), length=100)
lines(dnorm(s, mean(bth$residuals),sd(bth$residuals))~s, col=2)
plot(predict(bth),bth$residuals)</pre>
```





```
#qqnorm(y = bth$residuals, plot.it = T )
#qqline(y=bth$residuals, col=2, lwd=2,lty=2)
stdz<-(stdres(bth))
hist(stdz, breaks = 100, probability = T )
s<-seq(min(stdz), max(stdz), length=100)
lines(dnorm(s, 0,1)~s, col=2)
qqnorm(y = stdz, plot.it = T )
qqline(y=stdz, col=2, lwd=2,lty=2)
abline(a = 0,b = 1, col=3)</pre>
```





```
print(ks.test(x=bth$residuals, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: bth$residuals
## D = 0.16872, p-value < 2.2e-16
## alternative hypothesis: two-sided</pre>
```

```
print(ks.test(x=stdz, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: stdz
## D = 0.092255, p-value = 8.101e-08
## alternative hypothesis: two-sided
```

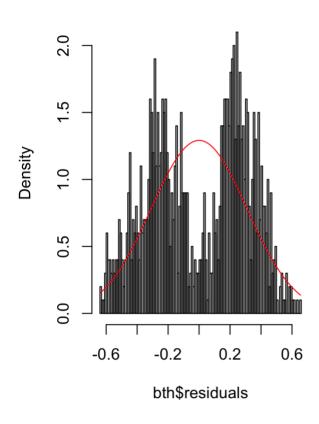
```
############
                  MODEL 7 # dependent normal #########
k<-3 # independent variables
n<- 1000 # number of observations
X < -array(0, dim = c(n, (k+1)))
bt < -c(1,2.3, 1.5,0.05)
x1 < -rgamma(n, 2, 3)
x2 < -rbinom(n, 10, 0.7)
x3 < -rbeta(n, 0.5, 0.5)
X[,1] < -1
X[,2]<-x1
X[,3]<-x2
X[,4]<-x3
sigma<-2
eps < -BM(x = 0.1, t0 = 0, T = 1, N = (n-1)) # dependent normal Brownian motion
y<-X%*%bt+eps
d<- data.frame(y,X[,2],X[,3],X[,4])</pre>
bth < -lm(y \sim X[,2] + X[,3] + X[,4], data = d)
cat("trur beta=",bt, "sigma=",sigma,'\n')
```

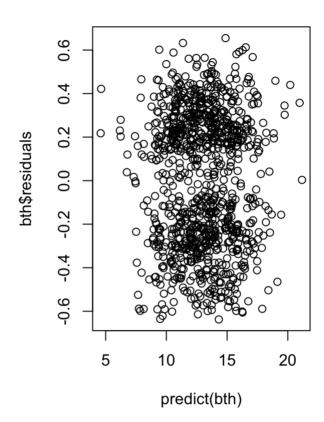
```
## trur beta= 1 2.3 1.5 0.05 sigma= 2
```

cat("beta_hat=",coefficients(bth), "sigma_hat=",summary(bth)\$sigma,'\n') # model coef
ficients

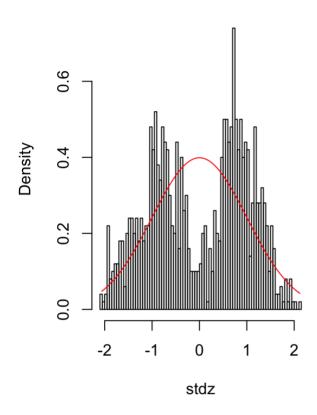
```
## beta_hat= 1.17556 2.281161 1.498571 0.09746791 sigma_hat= 0.3093547
```

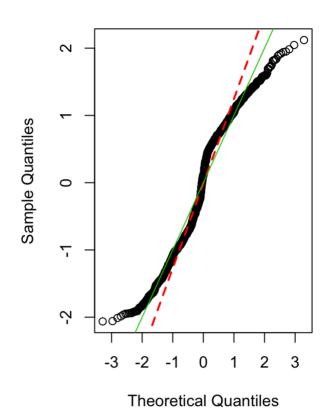
```
par(mfrow=c(1,2))
hist(bth$residuals, breaks = 100, probability = T )
s<-seq(min(bth$residuals), max(bth$residuals), length=100)
lines(dnorm(s, mean(bth$residuals),sd(bth$residuals))~s, col=2)
plot(predict(bth),bth$residuals)</pre>
```





```
#qqnorm(y = bth$residuals, plot.it = T )
#qqline(y=bth$residuals, col=2, lwd=2,lty=2)
stdz<-(stdres(bth))
hist(stdz, breaks = 100, probability = T )
s<-seq(min(stdz), max(stdz), length=100)
lines(dnorm(s, 0,1)~s, col=2)
qqnorm(y = stdz, plot.it = T )
qqline(y=stdz, col=2, lwd=2,lty=2)
abline(a = 0,b = 1, col=3)</pre>
```





```
print(ks.test(x=bth$residuals, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: bth$residuals
## D = 0.29114, p-value < 2.2e-16
## alternative hypothesis: two-sided</pre>
```

```
print(ks.test(x=stdz, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: stdz
## D = 0.12014, p-value = 5.805e-13
## alternative hypothesis: two-sided
```

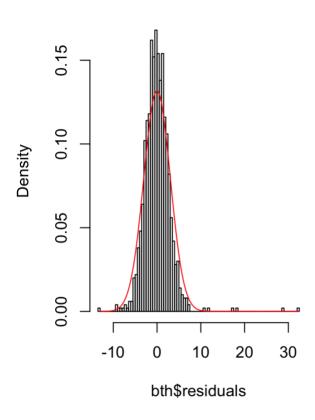
```
############
                  MODEL 8 # wrong model 1 #########
k<-3 # independent variables
n<- 1000 # number of observations
X < -array(0, dim = c(n, (k+1)))
bt < -c(1,2.3, 1.5,0.05)
x1 < -rgamma(n, 2, 3)
x2 < -rbinom(n, 10, 0.7)
x3 < -rbeta(n, 0.5, 0.5)
X[,1] < -1
X[,2]<-x1
X[,3]<-x2
X[,4]<-100*x3*x1^2
sigma<-2
eps<-rnorm(n,0,sigma)
y<-X%*%bt+eps
X[,4]<-x3
d<- data.frame(y,X[,2],X[,3],X[,4])</pre>
bth < -lm(y \sim X[,2] + X[,3] + X[,4], data = d)
cat("trur beta=",bt, "sigma=",sigma,'\n')
```

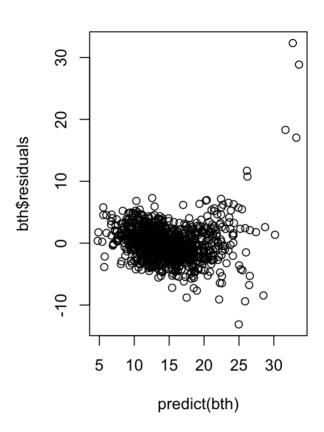
```
## trur beta= 1 2.3 1.5 0.05 sigma= 2
```

 $\verb|cat("beta_hat=",coefficients(bth), "sigma_hat=",summary(bth)$| sigma,'\n'|| \# model coefficients|| ficients||$

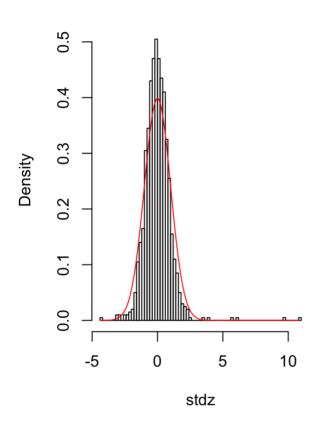
```
## beta_hat= -3.014627 7.26379 1.588446 3.406742 sigma_hat= 3.032979
```

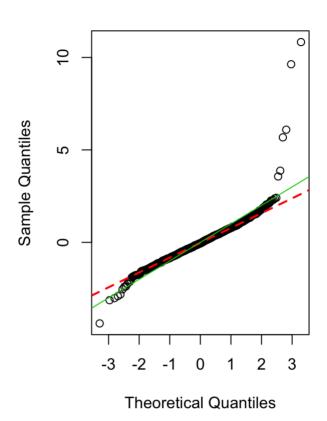
```
par(mfrow=c(1,2))
hist(bth$residuals, breaks = 100, probability = T )
s<-seq(min(bth$residuals), max(bth$residuals), length=100)
lines(dnorm(s, mean(bth$residuals), sd(bth$residuals))~s, col=2)
plot(predict(bth),bth$residuals)</pre>
```





```
#qqnorm(y = bth$residuals, plot.it = T )
#qqline(y=bth$residuals, col=2, lwd=2,lty=2)
stdz<-(stdres(bth))
hist(stdz, breaks = 100, probability = T )
s<-seq(min(stdz), max(stdz), length=100)
lines(dnorm(s, 0,1)~s, col=2)
qqnorm(y = stdz, plot.it = T )
qqline(y=stdz, col=2, lwd=2,lty=2)
abline(a = 0,b = 1, col=3)</pre>
```





```
print(ks.test(x=bth$residuals, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: bth$residuals
## D = 0.2173, p-value < 2.2e-16
## alternative hypothesis: two-sided</pre>
```

```
print(ks.test(x=stdz, y="pnorm",alternative = c("two.sided")))
```

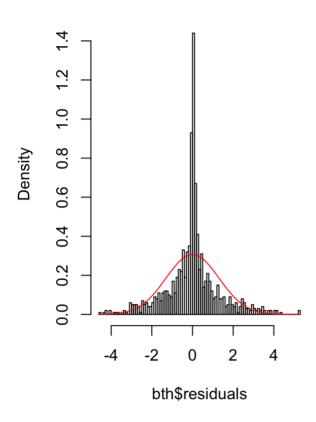
```
##
## One-sample Kolmogorov-Smirnov test
##
## data: stdz
## D = 0.06397, p-value = 0.000558
## alternative hypothesis: two-sided
```

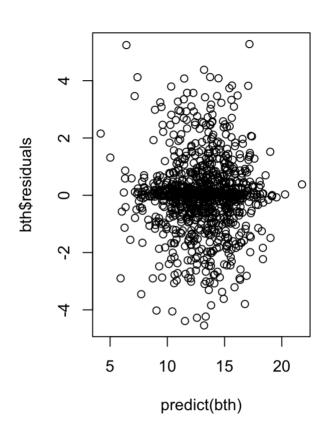
```
############
                  MODEL 9 # wrong model 2 #########
k<-3 # independent variables
n<- 1000 # number of observations
X < -array(0, dim = c(n, (k+1)))
bt < -c(1,2.3, 1.5,0.05)
x1 < -rgamma(n, 2, 3)
x2 < -rbinom(n, 10, 0.7)
x3 < -rbeta(n, 0.5, 0.5)
X[,1] < -1
X[,2]<-x1
X[,3]<-x2
X[,4]<-x3
sigma<-2
eps<-rnorm(n,0,sigma)
y<-X%*%bt+eps*x3
                    # error depends on x values
d<- data.frame(y,X[,2],X[,3],X[,4])</pre>
bth < -lm(y \sim X[,2] + X[,3] + X[,4], data = d)
cat("trur beta=",bt, "sigma=",sigma,'\n')
```

```
## trur beta= 1 2.3 1.5 0.05 sigma= 2
```

```
## beta_hat= 0.8565554 2.244991 1.521265 0.1050314 sigma_hat= 1.301697
```

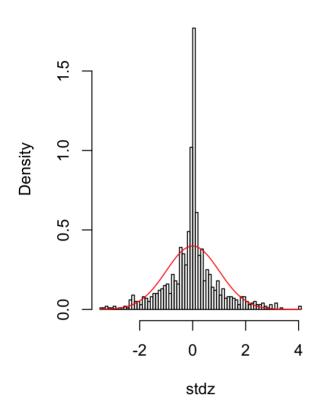
```
par(mfrow=c(1,2))
hist(bth$residuals, breaks = 100, probability = T )
s<-seq(min(bth$residuals), max(bth$residuals), length=100)
lines(dnorm(s, mean(bth$residuals), sd(bth$residuals))~s, col=2)
plot(predict(bth),bth$residuals)</pre>
```

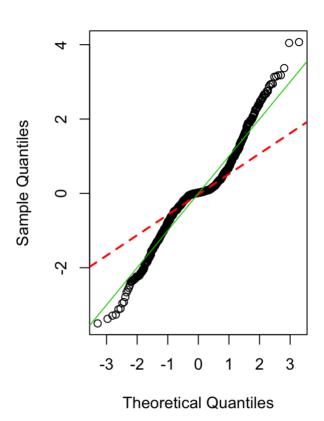




```
#qqnorm(y = bth$residuals, plot.it = T )
#qqline(y=bth$residuals, col=2, lwd=2,lty=2)

stdz<-(stdres(bth))
hist(stdz, breaks = 100, probability = T )
s<-seq(min(stdz), max(stdz), length=100)
lines(dnorm(s, 0,1)~s, col=2)
qqnorm(y = stdz, plot.it = T )
qqline(y=stdz, col=2, lwd=2,lty=2)
abline(a = 0,b = 1, col=3)</pre>
```





```
print(ks.test(x=bth$residuals, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: bth$residuals
## D = 0.10221, p-value = 1.688e-09
## alternative hypothesis: two-sided
```

```
print(ks.test(x=stdz, y="pnorm",alternative = c("two.sided")))
```

```
##
## One-sample Kolmogorov-Smirnov test
##
## data: stdz
## D = 0.12544, p-value = 4.297e-14
## alternative hypothesis: two-sided
```

```
# Motor Trend Car Road Tests
# Description
# The data was extracted from the 1974 Motor Trend US magazine,
# and comprises fuel consumption and 10 aspects of automobile design
# and performance for 32 automobiles (1973-74 models).
# A data frame with 32 observations on 11 variables.
# [, 1] mpg
              Miles/(US) gallon
# [, 2] cyl
              Number of cylinders
# [, 3] disp
                Displacement (cu.in.)
# [, 4] hp
              Gross horsepower
# [, 5] drat
                Rear axle ratio
# [, 6] wt
              Weight (1000 lbs)
# [, 7] qsec
                1/4 mile time
# [, 8] vs
              V/S
# [, 9] am
              Transmission (0 = automatic, 1 = manual)
# [,10] gear
                Number of forward gears
# [,11] carb
                Number of carburetors
# Source
# Henderson and Velleman (1981), Building multiple regression models interactively. B
iometrics, 37, 391-411.
rsqurd < -array(0, dim = c(10))
adjrsqurd < -array(0, dim = c(10))
fit<-lm(mpg~wt, data = mtcars)</pre>
rsqurd[1]<-summary(fit)$r.squared
adjrsqurd[1]<-summary(fit)$adj.r.squared
fit<-lm(mpg~wt+am, data = mtcars)</pre>
rsqurd[2]<-summary(fit)$r.squared
adjrsqurd[2]<-summary(fit)$adj.r.squared
fit<-lm(mpg~wt+am+gsec, data = mtcars)</pre>
rsqurd[3]<-summary(fit)$r.squared
adjrsqurd[3]<-summary(fit)$adj.r.squared
fit<-lm(mpg~wt+am+gsec+hp, data = mtcars)</pre>
rsqurd[4]<-summary(fit)$r.squared
adjrsqurd[4]<-summary(fit)$adj.r.squared
fit<-lm(mpg~wt+am+qsec+hp+disp, data = mtcars)</pre>
rsqurd[5]<-summary(fit)$r.squared
adjrsqurd[5]<-summary(fit)$adj.r.squared
fit<-lm(mpg~wt+am+qsec+hp+disp+drat, data = mtcars)</pre>
rsqurd[6]<-summary(fit)$r.squared</pre>
adjrsqurd[6]<-summary(fit)$adj.r.squared
fit<-lm(mpg~wt+am+qsec+hp+disp+drat+gear, data = mtcars)</pre>
rsqurd[7]<-summary(fit)$r.squared</pre>
adjrsqurd[7]<-summary(fit)$adj.r.squared
```

```
fit<-lm(mpg~wt+am+qsec+hp+disp+drat+gear+carb, data = mtcars)
rsqurd[8]<-summary(fit)$r.squared
adjrsqurd[8]<-summary(fit)$adj.r.squared

fit<-lm(mpg~wt+am+qsec+hp+disp+drat+gear+carb+vs, data = mtcars)
rsqurd[9]<-summary(fit)$r.squared
adjrsqurd[9]<-summary(fit)$adj.r.squared

fit<-lm(mpg~wt+am+qsec+hp+disp+drat+gear+carb+vs+cyl, data = mtcars)
rsqurd[10]<-summary(fit)$r.squared
adjrsqurd[10]<-summary(fit)$r.squared

par(mfrow=c(1,1))
plot(rsqurd, type='b', ylim=c(0.7,0.88), main="R^2 (black) & adj R^2 (red)")
lines(adjrsqurd,col=2,type = 'b')</pre>
```

R^2 (black) & adj R^2 (red)

