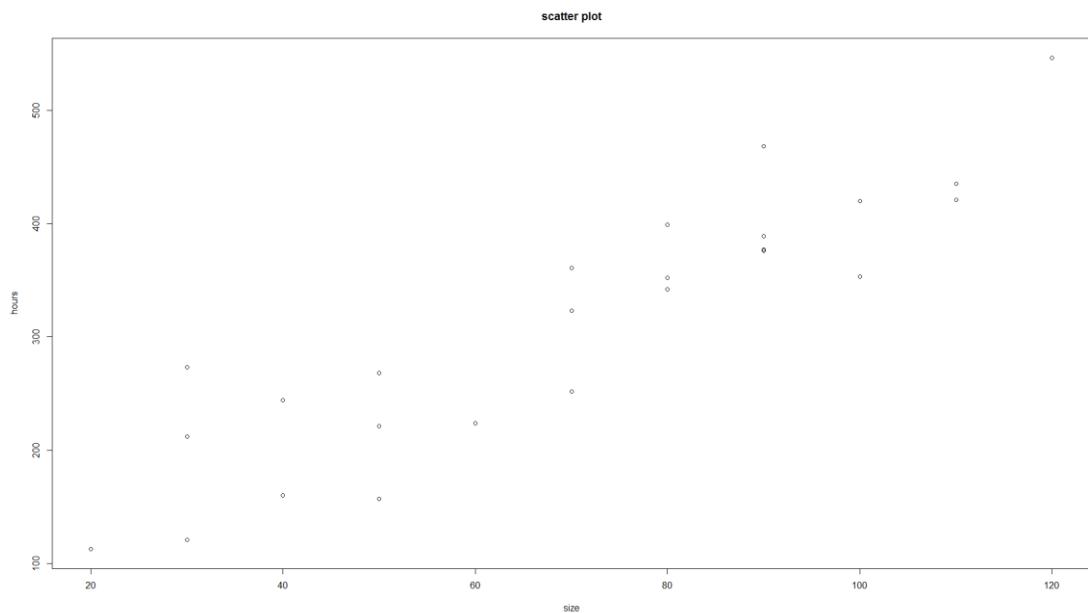


```
> #####testing constant variance#####
##Toluca Company example. To discover the relationship
##between lot size and labor hours required to produce the lot. Data
##on lot size and work hours for 25 recent production runs were utilized
```

```
tuluca<-read.table(file="C:/jenn/teaching/stat440540/data/CH1/CH01TA01.txt")
> tuluca
  V1  V2
1 80 399
2 30 121
3 50 221
4 90 376
> size<-tuluca$V1
  > hours<-tuluca$V2

plot(size,hours, main="scatter plot")
```



```
> myfit<-lm(hours~size)
> b_0 <- myfit$coef[1]
> b_1 <- myfit$coef[2]
> plot(size,hours, main="Fitted Line Plot")
> abline(b_0, b_1)
> resid<-myfit$residuals
> residsquare<-resid^2
  myfit2<-lm(residsquare~size)
```

```
anova(myfit)
#myfit<-lm(hours~size)
```

Analysis of Variance Table

Response: hours

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
size	1	252378	252378	105.88	4.449e-10 ***
Residuals	23	54825	2384		

```
> anova(myfit2) # myfit2<-lm(residsquare~size)
```

Analysis of Variance Table

Response: residsquare

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
size	1	7896142	7896142	1.0914	0.307
Residuals	23	166395896	7234604		

BP statistic = $(7896142/2)/(54825/25)^2 = 0.821$, compared with $\chi^2(0.95, 1) = 3.84$, don't reject H_0 , that the error variance is constant.

```
> bptest(hours~size,studentize=FALSE)
```

Breusch-Pagan test

data: hours ~ size
BP = 0.82092, df = 1, p-value = 0.3649

```
>
> shapiro.test(resid)
```

Shapiro-Wilk normality test

data: resid
W = 0.9789, p-value = 0.8626

p-value is 0.8626, do not reject null hypothesis of normality.

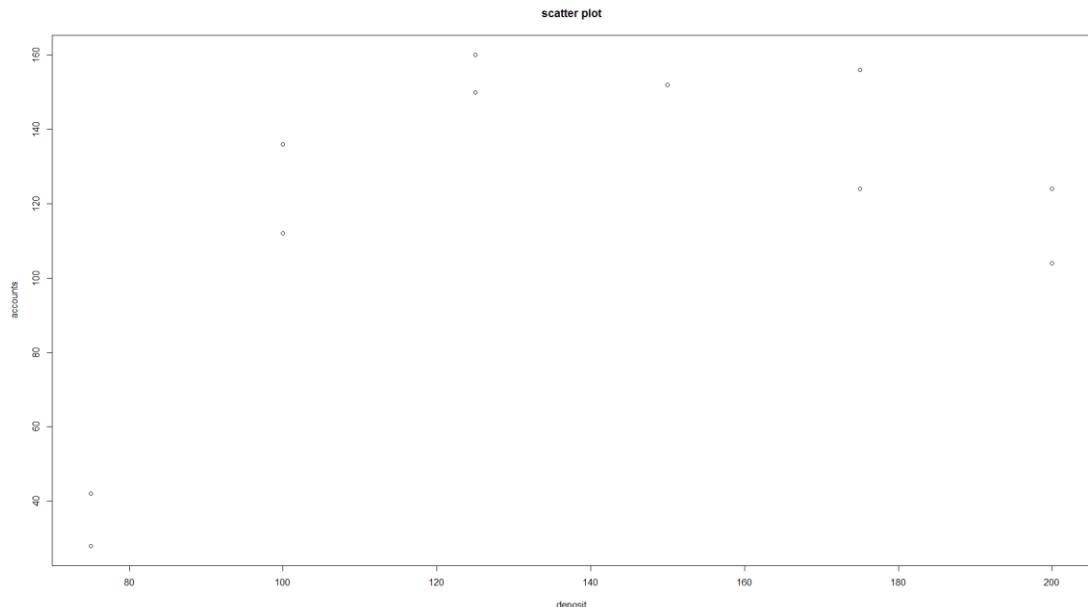
#####Fisher's lack of fit test#####

12 suburban branch offices of a commercial bank.
Holders of checking accounts were offered gifts for setting up money market accounts. Minimum deposits were required and the value of the gift was directly proportional to the minimum deposit. 6 levels were used. One bank dropped out. Linear fit is bad.

```

> bank<-read.table(file="C:/jenn/teaching/stat440540/data/CH3/CH03TA04.txt")
> bank
  V1 V2
1 125 160
2 100 112
3 200 124
4 75 28
5 150 152
6 175 156
7 75 42
8 175 124
9 125 150
10 200 104
11 100 136
> deposit<-bank$V1
> accounts<-bank$V2
plot(deposit,accounts, main="scatter plot")

```



```

> #full model
> myfit = aov(accounts~factor(deposit), data=bank)
> summary(mymfit)

```

Df	Sum Sq	Mean Sq	F value	Pr(>F)	
factor(deposit)	5	18735	3747	16.32	0.00409 **
Residuals	5	1148	230		

```

> #reduced model
> myfit2<-lm(accounts~deposit)
> summary(mymfit2)

```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	50.7225	39.3979	1.287	0.23
deposit	0.4867	0.2747	1.772	0.11

Residual standard error: 40.47 on 9 degrees of freedom
 Multiple R-squared: 0.2586, Adjusted R-squared: 0.1762
 F-statistic: 3.139 on 1 and 9 DF, p-value: 0.1102

> summary(myfit)

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
factor(deposit)	5	18735	3747	16.32	0.00409 **
Residuals	5	1148	230		

> anova(myfit2)

Analysis of Variance Table

Response: accounts

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
deposit	1	5141.3	5141.3	3.1389	0.1102
Residuals	9	14741.6	1638.0		

>

$F^* = MSLF/MSPE = [(14741.6 - 1148)/4]/(1148/5) = 3398.4/229.6 = 14.80$, let $\alpha = .01$, then

$F(.99; 4, 5) = 11.4$, $F^* = 14.8 > 11.4$, P value for the test is 0.006. we conclude $H_{\{\alpha\}}$, that the regression function is not linear.

#generalized linear test

> anova(myfit, myfit2)

Analysis of Variance Table

Model 1: accounts ~ factor(deposit)

Model 2: accounts ~ deposit

Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	5	1148			
2	9	14742	-4	-13594	14.801 0.005594 **

Reject the reduced model, conclude that the regression function is not linear.

```

##use boxcox to select transformation
library(MASS)
boxcox(plasma ~ age, lambda = seq(-2, 2, length = 10))

##find sses in TABLE 3.9

#W_I = K_1(Y_I^{\lambda} - 1), if \lambda \neq 0
#W_I = K_2 (\log_e Y_I), if \lambda = 0
#Where
#K_2 = (\prod_{I=1}^n Y_I)^{1/n}
#K_1 = 1/\lambda K_2^{\lambda - 1}

gmean <- exp(mean(log(plasma))) #geometric mean K_2 in page 135
sse <- c() #assign null vector to sse
lambda <- c() #assign null vector to lambda
i <- 1 #assign 1 to i as initial value
##construct a loop to calculate sses for each lambda
for (lam in seq(-1,1,0.1)){
  if (lam != 0){
    tY <- (plasma^lam - 1) / (lam*gmean^(lam-1))
  }
  else {
    tY <- log(plasma)*gmean
  }
  test <- anova(lm(tY~age))
  sse[i] <- test['Residuals','Sum Sq']
  lambda[i] <- lam
  i <- i+1
}
cbind(lambda,sse)

> cbind(lambda,sse)
   lambda     sse
[1,] -1.0 33.90887
[2,] -0.9 32.70442
[3,] -0.8 31.76453
[4,] -0.7 31.09066
[5,] -0.6 30.68680
[6,] -0.5 30.55961
[7,] -0.4 30.71859
[8,] -0.3 31.17631
[9,] -0.2 31.94867
[10,] -0.1 33.05520
[11,]  0.0 34.51945
[12,]  0.1 36.36939
[13,]  0.2 38.63791
[14,]  0.3 41.36342
[15,]  0.4 44.59051
[16,]  0.5 48.37072
[17,]  0.6 52.76343

```

```
[18,] 0.7 57.83686  
[19,] 0.8 63.66932  
[20,] 0.9 70.35050  
[21,] 1.0 77.98306  
>
```