

Mini Project #2

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Contribution of each group member: 100% for Junmei Fan, 100% for Xi Cui

1. Built the multiple linear regression model to predict murder rate based on the other variables.
 - a) Fit the multiple linear regression to predict murder.rate by all variables (ie, poverty, high.school, college, single.parent, unemployed, metropolitan, region)

$$y_i = \beta_0 + \beta_1 x_{i,1} + \beta_2 x_{i,2} + \cdots + \beta_{p-1} x_{i,p-1} + \varepsilon_i$$

Testing the multiple linear model significance:

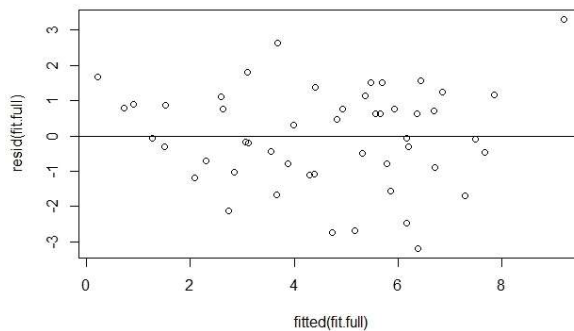
$H_0: \beta_1 = \beta_2 = \cdots = \beta_{p-1} = 0$ VS H_a : at least one β_i not equal to 0

Test statistic: $F = \frac{MS_{reg}}{MS_{err}} = 9.851 \sim F_{9,40}$, P-value = 9.287×10^{-8} .

Conclusion: As the P-value is small, we reject the H_0 and conclude that there is linear relationship between Murder rate and predictors of poverty, high school etc. The multiple linear model is reasonable.

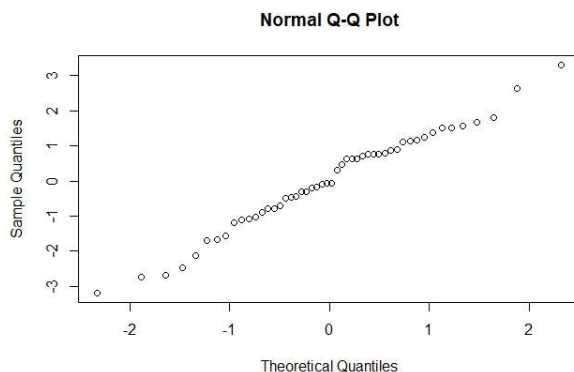
- b) Check the multiple linear regression assumptions.

- i) ε_i : Errors have mean zero and constant variance



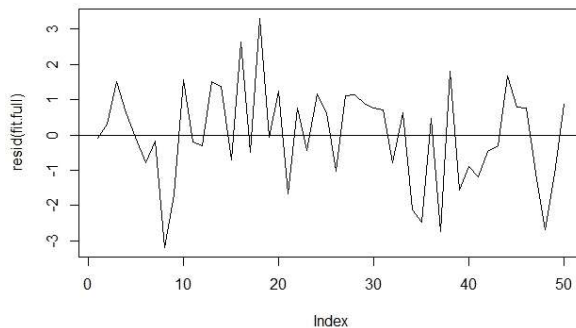
From the plot we can see the errors distribute around 0 and no pattern, which means Errors have mean zero and constant variance.

- ii) ε_i : Errors are normal distributed



The plot shows the errors are good fit of normal distribution, there is no long tail.

iii) ε_t : Errors are independent



The time series plot shows the errors are randomly to the index, which means they are independent.

c) Testing the significance of jth predictor:

$H_0: \beta_j = 0$ VS $H_a: \beta_j$ not equal to 0.

Test statistic: $t = \frac{\hat{\beta}_j - 0}{SE(\hat{\beta}_j)} \sim t_{40}$

Out put of R:

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	1.15569	11.06682	0.104	0.917352
poverty	0.07124	0.12615	0.565	0.575397
high.school	-0.12534	0.11815	-1.061	0.295116
college	0.08368	0.08238	1.016	0.315857
single.parent	0.38015	0.10559	3.600	0.000867 ***
unemployed	0.29521	0.33119	0.891	0.378059
metropolitan	0.03095	0.01536	2.015	0.050607 .
regionNortheast	-2.57007	0.76665	-3.352	0.001761 **
regionSouth	-0.12303	0.77605	-0.159	0.874832
regionWest	-0.83460	0.76033	-1.098	0.278904

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Find the important predictors.

Depending on the output of testing the significance of jth predictor, we drop the predictors with high P-values which we failed the reject the H_0 of the test and conclude the weak relation between predictors and murder rate.

i) Drop the “Poverty” and “Unemployed” and test whether the new model is more reasonable.

Partial F-test to compare the two models:

Model 1: $\text{murder.rate} \sim \text{poverty} + \text{high.school} + \text{college} + \text{single.parent} + \text{unemployed} + \text{metropolitan} + \text{region}$

Model 2: murder.rate ~ single.parent + region + metropolitan + high.school + college

$H_0: \beta_{poverty} = \beta_{unemployed} = 0$ VS H_a : at least one β not equal to 0.

Test statistic: $F = \frac{MS_{\text{Sex}}}{MS_{\text{err(full)}}} = 0.6476$, P – value = 0.5287

Conclusion: As the P-value is greater than 5%, we failed to reject the H_0 , and conclude that the “Poverty” and “Unemployed” can be dropped.

ii) Test whether the “High school” and “College” could be dropped:

Model 1: murder.rate ~ single.parent + region + metropolitan + high.school + college

Model 2: murder.rate ~ single.parent + region + metropolitan + high.school

$H_0: \beta_{college} = 0$ VS $H_a: \beta_{college}$ not equal to 0.

Test statistic: $F = \frac{MS_{\text{Sex}}}{MS_{\text{err(full)}}} = 0.9436$, P – value = 0.3369

Conclusion: As the P-value is greater than 5%, we failed to reject the H_0 , and conclude that the “College” can be dropped.

Model 1: murder.rate ~ single.parent + region + metropolitan + high.school

Model 2: murder.rate ~ single.parent + region + metropolitan

$H_0: \beta_{high\ school} = 0$ VS $H_a: \beta_{high\ school}$ not equal to 0.

Test statistic: $F = \frac{MS_{\text{Sex}}}{MS_{\text{err(full)}}} = 2.5724$, P – value = 0.1161

Conclusion: As the P-value is greater than 5%, we failed to reject the H_0 , and conclude that the “High school” can be dropped.

iii) We also test if we can drop more predictors, and both the test of partial F-test for “single parent”, “region”, and “metropolitan” are failed to reject H_0 , so we should keep these three predictors.

R Output:

Model 1: murder.rate ~ single.parent + metropolitan

Model 2: murder.rate ~ single.parent + region + metropolitan

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	47	152.21				
2	44	107.39	3	44.824	6.122	0.001425 **

Model 1: murder.rate ~ single.parent + region

Model 2: murder.rate ~ single.parent + region + metropolitan

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	45	132.91				
2	44	107.39	1	25.528	10.46	0.002317 **

Model 1: murder.rate ~ metropolitan + region

Model 2: murder.rate ~ single.parent + region + metropolitan

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	45	175.70				
2	44	107.39	1	68.316	27.991	3.672e-06 ***

Above all we keep the “single parent”, “region”, and “metropolitan” as the predictors for “murder rate”.

- d) Explore the interaction of “single parent”, “region”, and “metropolitan”.

Partial F-test for comparing the two models:

Interaction between “single parent” and “metropolitan”

Model 1: `murder.rate ~ single.parent + region + metropolitan`

Model 2: `murder.rate ~ single.parent + region + metropolitan + single.parent:metropolitan`

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	44	107.387				
2	43	98.882	1	8.5051	3.6985	0.0611

Conclusion: The p-value is greater than 0.05, we failed to reject the hypothesis that the coefficient of new term “interaction of single parent and metropolitan” is “0”, and conclude that there is no interaction between these two predictors.

And we get the same conclusion for inspection of “single parent” and “region”, “region” and “metropolitan”.

R Output show as follow:

Model 1: `murder.rate ~ single.parent + region + metropolitan`

Model 2: `murder.rate ~ single.parent + region + metropolitan + single.parent:region`

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	44	107.39				
2	41	102.24	3	5.1466	0.6879	0.5646

Model 1: `murder.rate ~ single.parent + region + metropolitan`

Model 2: `murder.rate ~ single.parent + region + metropolitan + metropolitan:region`

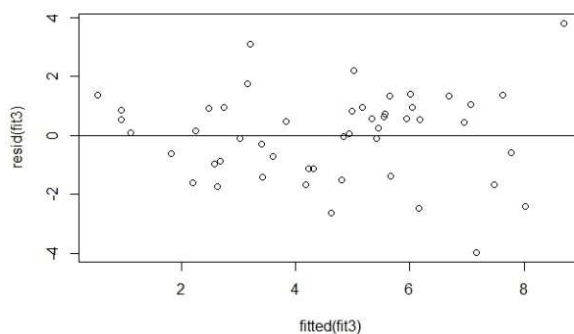
	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	44	107.39				
2	41	103.60	3	3.7879	0.4997	0.6846

- e) Final Multiple Linear Model:

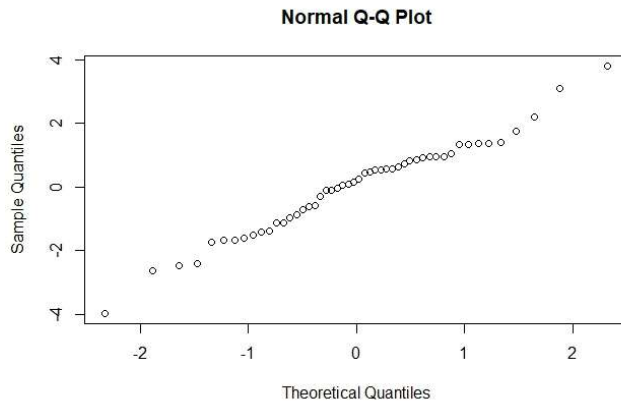
$$\text{Murder.rate} = -8.44469 + 0.47472 \text{ single.parent} + 0.03627 \text{ metropolitan} - 2.29258 \text{ regionNortheast} + 0.51237 \text{ regionSouth} - 0.24384 \text{ regionWest}$$

Check the multiple linear regression assumptions.

- i) ϵ_i : Errors have mean zero and constant variance

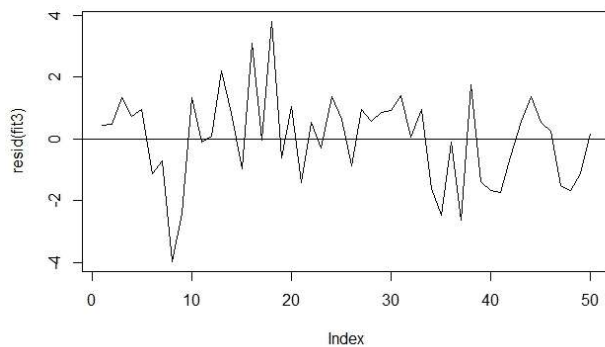


ii) ε_t : Errors are normal distributed



The plot shows the errors are good fit of normal distribution, there is no long tail.

iii) ε_t : Errors are independent



The time series plot shows the errors are randomly to the index, which means they are independent.

The final model holds all assumptions, we don't need do transformation for this model.

2. Predict murder rate:

Mean of single.parent = 22.97; Mean of metropolitan = 67.73; Most frequent region = "south"

Murder.rate = 5.428477

R_CODE:

```
#Data preparing
data<-read.table("C:/Users/xicui/Desktop/crime.csv", header = TRUE, sep=",")

#summery of mean statistics
summary(data)
str(data)
#Factorlize the 'region'
data$region<-as.factor(data$region)
```

```

#Full modle multile linear regression
fit.full<-
lm(murder.rate~poverty+high.school+college+single.parent+unemployed+metropolitan+region,data=
data)
summary.lm(fit.full)

# Residual plot
plot(fitted(fit.full), resid(fit.full))
abline(h = 0)
# QQ plot
qqnorm(resid(fit.full))
# Time series plot of residuals
plot(resid(fit.full), type="l")
abline(h=0)

#Finding proper predictors with partial F-test
fit1<-lm(murder.rate~single.parent+region+metropolitan+high.school+college,data)
summary.lm(fit1)
anova(fit1,fit.full)
fit2<-lm(murder.rate~single.parent+region+metropolitan+high.school,data)
summary.lm(fit2)
anova(fit1,fit2)
fit3<-lm(murder.rate~single.parent+region+metropolitan,data)
summary.lm(fit3)
anova(fit2,fit3)
#use the fit to check if we could drop more predictors
fit<-lm(murder.rate~metropolitan+region,data)
anova(fit,fit3)

#Check if there is interaction between predictors
fit4<-lm(murder.rate~single.parent+region+metropolitan+single.parent:metropolitan,data)
anova(fit3,fit4)
fit5<-lm(murder.rate~single.parent+region+metropolitan+single.parent:region,data)
anova(fit3,fit5)
fit6<-lm(murder.rate~single.parent+region+metropolitan+metropolitan:region,data)
anova(fit3,fit6)

#fit3 as the final model

#library(MASS)
#step <- stepAIC(fit.full, direction="both")

```

```
#step$anova  
#transformation  
#library(MASS)  
#bc<-boxcox(fit2, lambda = seq(-2, 2, 1/10))  
#(lambda <- bc$x[which.max(bc$y)])
```

```
# Residual plot  
plot(fitted(fit3), resid(fit3))  
abline(h = 0)  
# QQ plot  
qqnorm(resid(fit3))  
# Time series plot of residuals  
plot(resid(fit3), type="l")  
abline(h=0)
```

```
#Q2.predict the murder.rate  
predict(fit3, newdata=data.frame(single.parent=mean(data$single.parent),metropolitan =  
mean(data$metropolitan),region ='South'))
```