Assignment 2

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1.

(a.)

> library(foreign)

> setwd("~/Desktop/Data/Assignment 2")

> mydata= read.dta("WAGE1.DTA")

(b.)

> tail(mydata,8)

(c.)

> mean(mydata\$wage)

[1] 5.896103

> max(mydata\$wage)

[1] 24.98

> min(mydata\$wage)

[1] 0.53

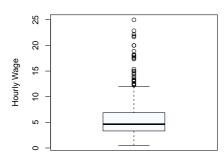
> median(mydata\$wage)

[1] 4.65

(d.)

>boxplot(mydata\$wage, main="A Simple Box Plot", ylab="Hourly Wage", col=blues9)





For the lower hinge, it is below 2.5. For the upper hinge, it is nearly 12.5.

Q1(25%) is about 3, Q2(50%) is about 5, and Q3(75%) is about 7.

And the dots above upper hinge are the outlier, which are a lot.

(e.)

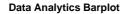
> mydata\$female[mydata\$female == 0] <- "Male"

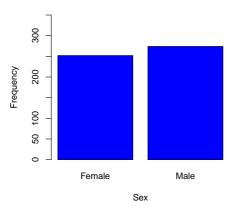
> mydata\$female[mydata\$female == 1] <- "Female"

> counts<-table(mydata\$female)

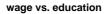
> counts

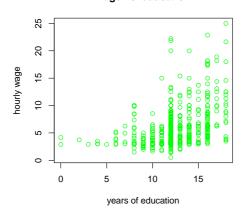
> barplot(counts, main = "Data Analytics Barplot", xlab = "Sex", ylab="Frequency", ylim=c(0,350),col="blue")





(f.)
>plot(mydata\$educ,mydata\$wage, main="wage vs. education", xlab = "years of education",
ylab = "hourly wage", col="green")





2. (a.)

> hours=mydata[,2]

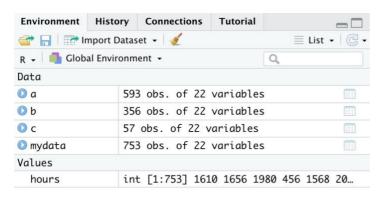
> table(hours)

hours

108 112 > (753-325)/753

[1] 0.5683931

(b.)



```
> table(mydata[,6])
```

```
5 6 7 8 9 10 11 12 13 14 15 16 17
```

> (381+44+51+14+57+46)

[1] 593

> 593/753

[1] 0.7875166

>b=mydata[mydata[,6]>=12 & mydata[,2]>0,]

We can see the observations from the upper right corner "environment", and the number is 356.

> 356/593

[1] 0.6003373

(c.)

>mydata[mydata[,6]>=16 & mydata[,11]<=16,]

> c=mydata[mydata[,6]>=16 & mydata[,11]<=16,]

We can see the observations from the upper right corner "environment", and the number is 57.

> 57/753

[1] 0.07569721

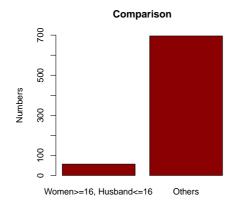
(d.)

> 753-57 (Total-"Women>=16, Husband<=16")

[1] 696,

> d<-c(57, 696)

> barplot(d, main = "Comparison", ylab = "Numbers",ylim =c(0,700), names.arg = c("Women>=16, Husband<=16","Others"), col = "darkred")



(e.)
> Im(formula= inlf~ nwifeinc+educ+kidslt6+exper+I(exper^2), data = mydata)

Call:

Im(formula = inlf ~ nwifeinc + educ + kidslt6 + exper + I(exper^2),
 data = mydata)

Coefficients:

(Intercept) nwifeinc educ kidslt6 exper I(exper^2)
-0.1288083 -0.0052223 0.0447794 -0.1696127 0.0421868 -0.0008764

> fit.marry<-lm(formula= inlf~ nwifeinc+educ+kidslt6+exper+I(exper^2), data = mydata)
> summary(fit.marry)

Call:

Im(formula = inlf ~ nwifeinc + educ + kidslt6 + exper + I(exper^2),
 data = mydata)

Residuals:

Min 1Q Median 3Q Max -0.9575 -0.4012 0.1523 0.3488 0.9978

Coefficients:

Estimate Std. Error t value Pr(>|t|) (Intercept) -0.1288083 0.0911499 -1.413 0.15803 nwifeinc -0.0052223 0.0014712 -3.550 0.00041 *** educ 0.0447794 0.0075133 5.960 3.89e-09 *** kidslt6 -0.1696127 0.0315918 -5.369 1.06e-07 ***

```
exper 0.0421868 0.0058298 7.236 1.15e-12 ***
I(exper^2) -0.0008764 0.0001867 -4.693 3.20e-06 ***
---
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Residual standard error: 0.4416 on 747 degrees of freedom Multiple R-squared: 0.2115, Adjusted R-squared: 0.2062

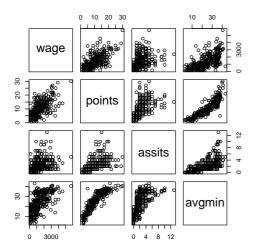
F-statistic: 40.07 on 5 and 747 DF, p-value: < 2.2e-16

 $\label{eq:control_inf} $$\inf_{-0.1288083} + (-0.0052223) *nwifeinc+0.0447794 *educ+(-0.1696127) *kidslt6+0.0421868 *exper+(-0.0008764) *I(exper^2) $$> (-0.1288083) + (-0.0052223) *20 + 0.0447794 *12 + (-0.1696127) *3 + 0.0421868 *6 + (-0.0008764) *36 $$ [1] 0.0168308 $$ inlf_hat=0.0168308$

3. (a.)

> Wage_Points_Assists_Avgmin=NBA_Salary[,c(2,11,13,15)]

> pairs(Wage_Points_Assists_Avgmin)



We could see that if we randomly pick two of the variables, the relationship between two of them would mostly be positive.

(b.)

> Wage_Points=Im(formula = wage ~ points, data = NBA_Salary)

> Im(Wage_Points)

```
Call:
```

Im(formula = Wage_Points)

Coefficients:

(Intercept) points 278.1 111.7

> summary(Wage_Points)

Call:

Im(formula = wage ~ points, data = NBA_Salary)

Residuals:

Min 1Q Median 3Q Max -1923.10 -463.10 -96.44 385.23 2728.56

Coefficients:

Estimate Std. Error t value Pr(>|t|)

(Intercept) 278.102 92.694 3.00 0.00295 ** points 111.667 7.841 14.24 < 2e-16 ***

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

Residual standard error: 755.1 on 267 degrees of freedom

Multiple R-squared: 0.4317, Adjusted R-squared: 0.4296

F-statistic: 202.8 on 1 and 267 DF, p-value: < 2.2e-16

The value of vmultiple R square is 0.4317, and for adjusted R square is 0.4296, which means that for variable "points", it could explain nearly 43% of the dependent variable "wage".

And normally for R square around 50%, we could consider as good.

(c.)

> confint(Wage Points, level=0.9)

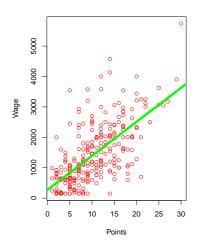
5 % 95 %

(Intercept) 125.10334 431.1012

points 98.72424 124.6092

90% confidence interval is between 98.72424 and 124.6092, meaning there is 90% of chances that our true value will lie within this range. Another interpretation is that if we have 100 samples, 90 samples' confidence interval will include the true value, and the other 10 samples will not.

(d.)
> plot(NBA_Salary\$points,NBA_Salary\$wage, xlab = "Points",ylab = "Wage" ,col="red")
> abline(Wage_Points, col="green" , lwd=5)



(e.)
> E=Im(wage~ points+avgmin+forward+center+exper+black, data=NBA_Salary)
> E

Call:

Im(formula = wage ~ points + avgmin + forward + center + exper +
 black, data = NBA_Salary)

Coefficients:

(Intercept) points avgmin forward center exper black -503.35 83.11 16.70 259.73 606.63 77.65 82.57

(f.)

- $> Include_guard=Im(wage^{\sim}\ points+avgmin+forward+center+exper+black+guard,\\ data=NBA_Salary)$
- > Include_guard

Call:

Im(formula = wage ~ points + avgmin + forward + center + exper +
 black + guard, data = NBA_Salary)

Coefficients:

(Intercept)	points	avgmin	forward	center	exper	black	guard
-503.35	83.11	16.70	259.73	606.63	77.65	82.57	NA

We can add variable "guard" in the regression, however, the coefficient will turn out to be "NA". And the reason is that "guard" is linearly dependent on "forward" and "center", the linear equation ought to be (1= forward+center+guard), you could only be one of them and we already include "forward" and "center" in our regression. Another example would be dummy variable "gender", if female=1 and male=0, when we run the regression on gender, we would only put either female or male but not both, the reason is that you could only be either one of them, thus female and male are linearly dependent.

(g.)

> summary(E)

Call:

```
Im(formula = wage ~ points + avgmin + forward + center + exper +
    black, data = NBA_Salary)
```

Residuals:

Min 1Q Median 3Q Max -1664.61 -382.68 -56.48 354.36 2830.20

Coefficients:

Estimate Std. Error t value Pr(>|t|)

(Intercept) -5	03.354	156.014 -3.	226 0.0	0141 **
points	83.115	15.119	5.498 9	.14e-08 ***
avgmin	16.695	9.306	1.794	0.07396 .
forward	259.732	89.941	2.888	0.00420 **
center	606.627	121.699	4.985 1	.13e-06 ***
exper	77.654	12.437	6.244 1	L.71e-09 ***
black	82.569	106.185	0.778	0.43751

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

Residual standard error: 669.5 on 262 degrees of freedom Multiple R-squared: 0.5616, Adjusted R-squared: 0.5515

F-statistic: 55.93 on 6 and 262 DF, p-value: < 2.2e-16

From the above summary of E,we could discover that variable "black" is not statistically significant, so it is unlikly to say that there is racial salary discrimination.

```
(h.)
```

> step(E, direction = "both")

Start: AIC=3507.43

wage ~ points + avgmin + forward + center + exper + black

Df Sum of Sq RSS AIC

- black 1 271037 117712809 3506.1

<none> 117441772 3507.4

- avgmin 1 1442751 118884523 3508.7

- forward 1 3738139 121179910 3513.9

- center 1 11137562 128579334 3529.8

- points 1 13547301 130989072 3534.8

- exper 1 17475480 134917252 3542.7

Step: AIC=3506.05

wage ~ points + avgmin + forward + center + exper

Df Sum of Sq RSS AIC

<none> 117712809 3506.1

+ black 1 271037 117441772 3507.4

- avgmin 1 1503284 119216093 3507.5

- forward 1 3805357 121518165 3512.6

- center 1 10866537 128579346 3527.8

- points 1 13565974 131278783 3533.4

- exper 1 17406020 135118828 3541.2

Call:

Im(formula = wage ~ points + avgmin + forward + center + exper,
 data = NBA_Salary)

Coefficients:

(Intercept)	points	avgmin	forward	center	exper
-442.75	83.17	17.02	261.93	591.96	77.49

From the AIC result of (wage ~ points + avgmin + forward + center + exper + black), we could know that it is better to remove variable "black" as it is shown on the first row. And from the AIC result of (wage ~ points + avgmin + forward + center + exper), we know that it is better not to remove any variable. And this result is consistent with the summary of E from question (e.). They both show that "black" is not an important variable.

(i.)> H=Im(wage ~ points + avgmin + forward + center + exper, data = NBA_Salary)> summary(H)

Coefficients:

Estimate Std. Error t value Pr(>|t|) (Intercept) -442.754 135.054 -3.278 0.00118 ** points 83.171 15.107 5.505 8.75e-08 *** 1.833 0.06798. avgmin 17.024 9.289 89.829 2.916 0.00385 ** forward 261.928 center 591.959 120.138 4.927 1.48e-06 *** 12.426 6.236 1.78e-09 *** 77.488 exper

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

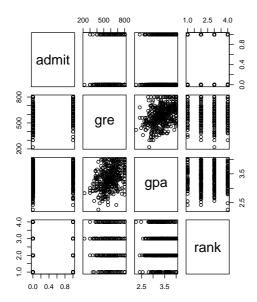
Residual standard error: 669 on 263 degrees of freedom

Multiple R-squared: 0.5606, Adjusted R-squared: 0.5522

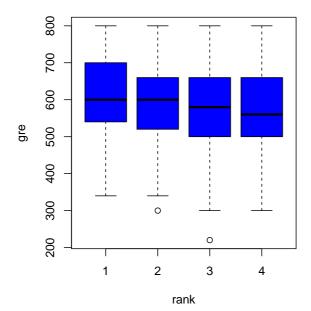
F-statistic: 67.1 on 5 and 263 DF, p-value: < 2.2e-16

In (e.), its R-square is 0.5616, adjusted R-square is 0.5515, as for (h.)its R-square is 0.5606, adjusted R-square is 0.5522. R-square decreases as we remove one of the variables, but it does not capture the quality of how we improve the model. For the quality of model improvement, we should put our focus on adjusted R-square, since adjusted R-square increases, we could know that removing "black" will make our model better, and this result is consistent with AIC result.

- 4.
- (a.)
- > library(foreign)
- > setwd("~/Desktop/Data/Assignment 2")
- > mydata= read.csv("admission.csv")
- > pairs(mydata)



(b.)
> boxplot(gre ~ rank, data = mydata, col="Blue")



(c.)

> rank_1<-mydata[mydata[,4]==1,]

61 are from ranked 1 institution, we can know that by looking at the upper right hand side.

> rank_1_rejected<- mydata[mydata[,4]==1 & mydata[,1]==0,]

28 are from ranked 1 institution and rejected, we can know that by looking at the upper right hand side.

> 28/61

[1] 0.4590164

Nearly 46% of them are rejected.

(d.)

> rank_4<-mydata[mydata[,4]==4,]

67 are from ranked 4 institution, we can know that by looking at the upper right hand side.

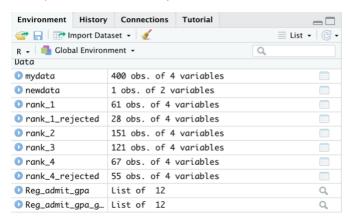
> rank_4_rejected<-mydata[mydata[,4]==4 & mydata[,1]==0,]

55 are from ranked 4 institution, we can know that by looking at the upper right hand side.

> 55/67

[1] 0.8208955

Nearly 82% of them are rejected.



(e.)

> Reg admit gpa<- Im(admit~gpa, data = mydata)

> Reg_admit_gpa

Call:

Im(formula = admit ~ gpa, data = mydata)

Coefficients:

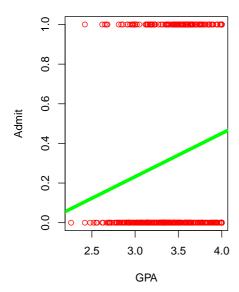
(Intercept) gpa -0.4224 0.2183

The coefficient is 0.2183, which means if there is an 1 unit increase in GPA, on average,

the chances of getting admitted will increase by 21.83%.

(f.)

- > plot(mydata\$gpa,mydata\$admit, xlab = "GPA",ylab = "Admit",col="red")
- > abline(Reg_admit_gpa, col="green", lwd=5)



We can see that for "admit"=1, people with lower GPA are unlikely to get admitted.

However, we can still see that people with higher GPA will somehow also be rejected.

From the regression line, we can see there is a positive relation between "admit" and "GPA".

(g.)

- > Reg_admit_gpa_gre<-lm(admit~ gpa+ gre, data=mydata)
- > Reg_admit_gpa_gre

Call:

Im(formula = admit ~ gpa + gre, data = mydata)

Coefficients:

- > newdata= data.frame(gpa=3.6, gre=700)
- > predict(Reg_admit_gpa_gre, newdata, interval="prediction", level=0.95)

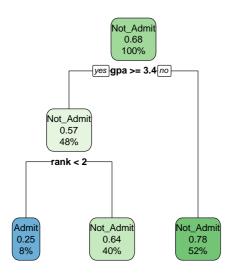
1 0.4115465 -0.4871885 1.310282

The probability of being accepted is around 41.15%.

```
5.
```

(a.)

- > library(foreign)
- > library(rpart)
- > setwd("~/Desktop/Data/Assignment 2")
- > Admission= read.csv("admission.csv")
- > Admission\$admit[Admission\$admit==0]<- "Not_Admit"
- > Admission\$admit[Admission\$admit==1]<- "Admit"
- > rtree<-rpart(admit~., data= Admission, minsplit=20, cp=0.05)
- > rpart.plot(rtree)



From this decision tree, we could see the first node is GPA, 48% of the samples are above 3.4, 52% are below 3.4, and the chances of not getting admitted is 78%. For those who had GPA above 3.4, and rank <2, admission rate is 75%. For those who had GPA above 3.4, and rank >2, the chances of not getting admitted would be 64%.

Since we set (cp=0.05 and minsplit =20), GRE seems to be not that important compare to other variables, so there is no node for GRE. But if we set cp=0.01, then we would see the node of GRE.

(b.)

> newdata= data.frame(gpa=3.6, gre=580, rank=2)

> predict(rtree, newdata)

Admit Not_Admit

1 0.3625 0.6375

> predict(rtree, newdata, type = "class")

1

Not_Admit

Based on the above prediction, it is likely that he or she will not be admitted.

6.

(a.)

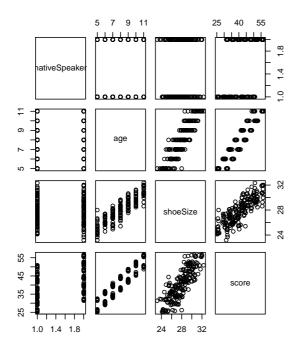
Both Gini index and entropy measure impurity of the nodes. And their goal is to group same observations together and looking for the most purity. The point that we are looking for is where it has the smallest Gini index or the largest entropy. Gini Index has values inside the interval [0, 0.5] whereas the interval of the Entropy is [0, 1] (b.)

- > install.packages("party")
- > library(party)
- > data("readingSkills")
- > summary(readingSkills)

nativeSpeaker	age		shoeSize			score	
no :100	Min.	: 5.000	Min.	:23.17	Min.	:25.26	
yes:100	1st Qu.:	6.000	1st Qu.:	26.23	1st Qu.:	33.94	
	Median : 8.000		Median :27.85		Median :40.33		
	Mean	: 7.925	Mea	n :27.	87 M	ean :40.66	
	3rd Qu.: 9.250		3rd Qu.:29.49 3rd (3rd Qu	ı.:47.57	
	Max.	:11.000	Max.	:32.3	3 Ma	x. :56.71	
(c.)							

(c.)

> pairs(readingSkills)



From the scatter plot, we could see the relationship between either two variables, but we

can't tell which is independent variable and which is dependent variable. So we can't draw causal inference.

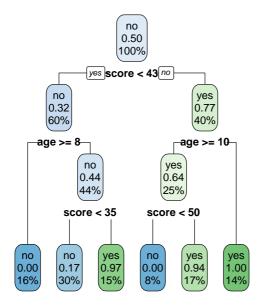
(d.)

- > library(rpart)
- > library(rpart.plot)

(e.)

> rtree<-rpart(nativeSpeaker~ shoeSize+age+score, data= readingSkills, minsplit=20, cp=0.05)

> rpart.plot(rtree)



The first node is score, then we could see that it uses age>=8 and age>=10 to be the nodes and again, it used score to divide the group. And we can see there is no such a node for shoe size, it seems shoesize isn't significant compared to the other variables.

7.
$$G(Hale) = 1 - (\frac{1}{3})^2 - (\frac{1}{3})^2 = \frac{4}{9}$$
 $G(Fenale) = 1 - (\frac{1}{3})^2 - (\frac{1}{3})^2 = \frac{4}{9}$
 $G(Suburban) = 1 - (\frac{1}{3})^2 - (\frac{1}{3})^2 = \frac{4}{9}$
 $G(Urban) = 1 - (\frac{1}{3})^2 - (\frac{1}{3})^2 = \frac{12}{49}$
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 $G(Secondary) = 1 - (\frac{1}$

yes, they do have the same answer since they are just using different ways to calculate the level of impurity.

I build the data in the excel and save it as "marry.csv"

- > library(foreign)
- > setwd("~/Downloads")
- > read.csv("marry.csv")
- > marry\$marry<-as.factor(marry\$marry)
- > marry\$gender<-as.factor(marry\$gender)
- > marry\$place<-as.factor(marry\$place)
- > marry\$education<-as.factor(marry\$education)
- > str(marry)

'data.frame': 12 obs. of 4 variables:

\$ marry : Factor w/ 2 levels "married", "single": 1 1 1 1 2 2 2 2 1 1 ...

\$ gender : Factor w/ 2 levels "female", "male": 1 1 1 1 1 1 2 2 2 2 ...

\$ education: Factor w/ 3 levels "degree", "no formal education", ...: 3 2 2 3 1 1 1 3 2 1 ...

\$ place : Factor w/ 2 levels "suburban", "urban": 1 1 1 2 2 2 1 1 2 2 ...

- > library(rpart)
- > library(rpart.plot)
- > rtree_gini<-rpart(marry~place+gender+education, data= marry, minsplit=0, cp=0.01, parms
- = list(split = "gini"))
- > rpart.plot(rtree_gini)

rtree_gini uses gini index to do decision tree, and its first node is education.

- > rtree_information<-rpart(marry~place+gender+education, data= marry, minsplit=0, cp=0.01, parms = list(split = "information"))
- > rpart.plot(rtree_information)

rtree_information uses information or entropy to do decision tree, and its first node is also education.

