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HW5 AMS 394

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Problem (1):
(1)
> stocks <-
read.table("http://www.ams.sunysb.edu/~xing/statfinbook/_BookData/Chap03/d_logret_6stocks.txt",
header=T)
> fit <- Im(stocks[,2] \sim stocks[,6])
> summary(fit)
Call:
Im(formula = stocks[, 2] ~ stocks[, 6])
Residuals:
   Min
           1Q Median
                            3Q
                                  Max
-0.049930 -0.013003 -0.000505 0.017353 0.049231
Coefficients:
       Estimate Std. Error t value Pr(>|t|)
(Intercept) -0.005325  0.002756 -1.932  0.05794.
stocks[, 6] 0.354649 0.119729 2.962 0.00433 **
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.02178 on 62 degrees of freedom
Multiple R-squared: 0.124, Adjusted R-squared: 0.1098
F-statistic: 8.774 on 1 and 62 DF, p-value: 0.004328
```

<u>Response:</u> The coefficient of correlation is .354649 and the coefficient of the intercept is -.005325.

```
(2)
> anova(fit)
Analysis of Variance Table
Response: stocks[, 2]
      Df Sum Sq Mean Sq F value Pr(>F)
stocks[, 6] 1 0.0041609 0.0041609 8.774 0.004328 **
Residuals 62 0.0294022 0.0004742
Response: The probability is lower than .01, thus we can reject the null hypothesis and conclude
that the regression effects are significant with over 99% confidence.
(3)
> Pfizer = stocks[,2]
> Exxon = stocks[,6]
> Citigroup = stocks[,4]
> data = data.frame(cbind(Pfizer,Exxon,Citigroup))
> stacked = stack(data)
> anova(lm(values ~ ind, stacked))
Analysis of Variance Table
Response: values
      Df Sum Sq Mean Sq F value Pr(>F)
ind
        2 0.001934 0.00096712 1.4351 0.2407
Residuals 189 0.127366 0.00067390
(4)
     1-sample proportions test with continuity correction
data: sum(Citigroup > 0) out of 64, null probability 0.5
```

X-squared = 1.2656, df = 1, p-value = 0.2606

```
alternative hypothesis: true p is not equal to 0.5
95 percent confidence interval:
0.4484671 0.6983808
sample estimates:
    p
0.578125
```

Response: The p value is greater than .05, so we cannot reject the null hypothesis that the proportion of Citigroup's returns that are greater than 0 is not significantly greater than .5.

```
Problem (2):
(1)
> data(juul)
> juul <- juul[,4:5]
> juul <- juul[complete.cases(juul),]
> one <- juul[juul[,"tanner"] == 1,]
> two <- juul[juul[,"tanner"] == 2,]
> three <- juul[juul[,"tanner"] == 3,]
> four <- juul[juul[,"tanner"] == 4,]
> five <- juul[juul[,"tanner"] == 5,]
> stacked = do.call("rbind", list(one,two,three,four,five))
> stacked$tanner <- as.factor(stacked$tanner)
> anova(lm(stacked[,1] \sim stacked[,2]))
Analysis of Variance Table
Response: stacked[, 1]
        Df Sum Sq Mean Sq F value Pr(>F)
stacked[, 2] 4 12696217 3174054 228.35 < 2.2e-16 ***
Residuals 787 10939116 13900
Signif. codes: 0 "*** 0.001 "** 0.01 "* 0.05 ". 0.1 " 1
```

<u>Response:</u> Since the p value is less than .01, we can say with more than 99% confidence that the five levels of tanner give significantly different results.

```
(2)
> mean(one[,1])
[1] 207.4727
> mean(two[,1])
[1] 352.6714
> mean(three[,1])
[1] 483.2222
> mean(four[,1])
[1] 513.0172
> mean(five[,1])
[1] 465.3344
(3)
> one <- one[,1]
> two <- two[,1]
> three <- three[,1]
> four <- four[,1]
> five <- five[,1]
> shapiro.test(one)
    Shapiro-Wilk normality test
data: one
W = 0.96947, p-value = 3.764e-06
> shapiro.test(two)
```

Shapiro-Wilk normality test

data: two

W = 0.9606, p-value = 0.02704

> shapiro.test(three)

Shapiro-Wilk normality test

data: three

W = 0.96348, p-value = 0.1657

> shapiro.test(four)

Shapiro-Wilk normality test

data: four

W = 0.94686, p-value = 0.01309

> shapiro.test(five)

Shapiro-Wilk normality test

data: five

W = 0.97828, p-value = 0.0001284

<u>Response</u>: Only one tanner level has normally distributed data, so non parametric tests will be used to compare all pairs of values.

> wilcox.test(one,two)

Wilcoxon rank sum test with continuity correction

data: one and two

W = 3550.5, p-value < 2.2e-16

alternative hypothesis: true location shift is not equal to 0

> wilcox.test(one,three)

Wilcoxon rank sum test with continuity correction

data: one and three

W = 712, p-value < 2.2e-16

alternative hypothesis: true location shift is not equal to 0

> wilcox.test(one,four)

Wilcoxon rank sum test with continuity correction

data: one and four

W = 300.5, p-value < 2.2e-16

alternative hypothesis: true location shift is not equal to 0

> wilcox.test(one,five)

Wilcoxon rank sum test with continuity correction

data: one and five

W = 5006.5, p-value < 2.2e-16

alternative hypothesis: true location shift is not equal to 0

<u>Response:</u> Tanner level one is significantly different from all other tanner levels, demonstrated by every pair wilcox test with one showing a p value less than .01.

> wilcox.test(two,three)

Wilcoxon rank sum test with continuity correction

data: two and three

W = 783, p-value = 5.733e-06

alternative hypothesis: true location shift is not equal to 0

> wilcox.test(two,four)

Wilcoxon rank sum test with continuity correction

data: two and four

W = 693, p-value = 1.579e-10

alternative hypothesis: true location shift is not equal to 0

> wilcox.test(two,five)

Wilcoxon rank sum test with continuity correction

data: two and five

W = 5702, p-value = 7.593e-10

alternative hypothesis: true location shift is not equal to 0

<u>Response:</u> Tanner level two is significantly different from all other tanner levels, demonstrated by every pair wilcox test with one showing a p value less than .01.

> wilcox.test(three,four)

Wilcoxon rank sum test with continuity correction

data: three and four

W = 1084, p-value = 0.1426

alternative hypothesis: true location shift is not equal to 0

> wilcox.test(three,five)

Wilcoxon rank sum test with continuity correction

data: three and five

W = 7332.5, p-value = 0.5295

alternative hypothesis: true location shift is not equal to 0

<u>Response:</u> Tanner level three is NOT significantly different from tanner levels four and five, demonstrated by those pairs wilcox tests showing a p value greater than .05.

> wilcox.test(four,five)

Wilcoxon rank sum test with continuity correction

data: four and five

W = 10996, p-value = 0.005231

alternative hypothesis: true location shift is not equal to 0

<u>Response:</u> Tanner level four is significantly different than tanner level five, as demonstrated by the p value being lower than .01.

Problem (3):

```
> data(survey)
> tbl = table(survey$Smoke, survey$Exer)
> chisq.test(tbl)
```

Pearson's Chi-squared test

data: tbl

X-squared = 5.4885, df = 6, p-value = 0.4828

Warning message:

In chisq.test(tbl): Chi-squared approximation may be incorrect

<u>Response</u>: Since the p value is greater than .05, we cannot reject the null hypothesis that the smoking habit is independent of the exercise level.