Stat3032_homework4

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Answer for 6.1

```
m6ln <- lm(lifeExpF ~ 1, data = UN11)
m6la <- lm(lifeExpF ~ group, data = UN11)
anova(m6ln, m6la)</pre>
```

```
## Analysis of Variance Table
##
## Model 1: lifeExpF ~ 1
## Model 2: lifeExpF ~ group
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 198 20293.2
## 2 196 7730.2 2 12563 159.27 < 2.2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</pre>
```

Answer for 6.3

```
m63n <- lm(lifeExpF ~ group, data = UN11)
m63a <- lm(lifeExpF ~ group + log(ppgdp), data = UN11)
anova(m63n, m63a)</pre>
```

```
## Analysis of Variance Table
##
## Model 1: lifeExpF ~ group
## Model 2: lifeExpF ~ group + log(ppgdp)
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 196 7730.2
## 2 195 5090.4 1 2639.8 101.12 < 2.2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</pre>
```

Answer for 6.4

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Answer for 6.4.1

NH model means lifeExpF is dependent on log(ppgdp) and the intercetion term of group and log(ppgdp). in other words, $E(lifeExpF \mid log(ppgdp), group) = b0 + b1 \times log(ppgdp) + b2 \times log(ppgdp) \times group$

Answer for 6.4.2

```
m64n <- lm(lifeExpF ~ log(ppgdp) + group:log(ppgdp), data = UN11)
m64a <- lm(lifeExpF ~ group + log(ppgdp) + group:log(ppgdp), data = UN11)
anova(m64n, m64a)</pre>
```

```
## Analysis of Variance Table
##
## Model 1: lifeExpF ~ log(ppgdp) + group:log(ppgdp)
## Model 2: lifeExpF ~ group + log(ppgdp) + group:log(ppgdp)
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 195 5232.0
## 2 193 5077.7 2 154.31 2.9326 0.05564 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Answer for 6.5

Answer for 6.5.1

```
u651 <- lm(lifeExpF ~ group + log(ppgdp) + group:log(ppgdp), UN11) summary(u651)$coef
```

```
## (Intercept) 59.2136614 15.220345 3.8904284 0.0001376669
## groupother -11.1731029 15.594836 -0.7164617 0.4745723433
## groupafrica -22.9848394 15.783786 -1.4562310 0.1469536042
## log(ppgdp) 2.2425354 1.466444 1.5292337 0.1278438256
## groupother:log(ppgdp) 0.9294372 1.517667 0.6124117 0.5409862449
## groupafrica:log(ppgdp) 1.0949810 1.578460 0.6937019 0.4887032151
```

Based on the table, the p-value are all greater than 0.05(except the intercept). Therefore, there are no difference between oecd and other level.

Answer for 6.5.2

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```
UN11$group1 <- relevel(UN11$group, "other")
u652 <- lm(lifeExpF ~ group + log(ppgdp) + log(ppgdp):group1, data = UN11)
summary(u652)$coef</pre>
```

```
##
                              Estimate Std. Error
                                                     t value
                                                                  Pr(>|t|)
## (Intercept)
                            59.2136614 15.2203448 3.8904284 1.376669e-04
## groupother
                           -11.1731029 15.5948356 -0.7164617 4.745723e-01
## groupafrica
                           -22.9848394 15.7837862 -1.4562310 1.469536e-01
## log(ppgdp)
                             3.1719725  0.3909682  8.1131211  5.638388e-14
## log(ppgdp):grouploecd
                           -0.9294372 1.5176672 -0.6124117 5.409862e-01
## log(ppgdp):grouplafrica
                             0.1655438    0.7028055    0.2355471    8.140339e-01
```

Based on the table, the p-value are all less than 0.05. Therefore, group other is not equal to group africa.

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