SET 8

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```
model_1 <- lm(formula = lifeExpF ~ I(log(ppgdp)) + pctUrban + fertility,</pre>
data = UN11)
summary(model_1)
##
## Call:
## lm(formula = lifeExpF ~ I(log(ppgdp)) + pctUrban + fertility,
       data = UN11)
##
## Residuals:
       Min
                  1Q
                      Median
                                    3Q
                                            Max
## -22.6165 -1.6683
                       0.5406
                                2.6425 11.2263
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                 64.21764
                             3.84281 16.711 < 2e-16 ***
                                       5.079 8.83e-07 ***
## I(log(ppgdp)) 2.17842
                             0.42888
## pctUrban
                  0.02116
                             0.02353
                                       0.900
                                                0.369
## fertility
                 -4.19652
                             0.39396 -10.652 < 2e-16 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Residual standard error: 5.145 on 195 degrees of freedom
## Multiple R-squared: 0.7456, Adjusted R-squared: 0.7417
## F-statistic: 190.5 on 3 and 195 DF, p-value: < 2.2e-16
```

Question 1(b)

Conclusion for the Overall F-Test

From the R output, the **overall F-test** evaluates whether at least one of the regression coefficients (excluding the intercept) is significantly different from zero.

- **F-statistic**: 190.5
- Degrees of Freedom (DF): 3, 195
- **p-value**: $< 2.2 \times 10\text{-}16$
- Significance Level (α): 0.01

Conclusion: The p-value for the F-test ($< 2.2 \times 10\text{-}16$) is much smaller than the significance level of 0.01. We reject the null hypothesis, concluding that the model as a whole is statistically significant.

Reasoning: At least one predictor in the model significantly contributes to explaining the variability in life expectancy.

```
reduced_model <- lm(lifeExpF ~ fertility, data = UN11)</pre>
# Perform the nested F-test
anova_result <- anova(reduced_model, model_1)</pre>
anova_result
## Analysis of Variance Table
## Model 1: lifeExpF ~ fertility
## Model 2: lifeExpF ~ I(log(ppgdp)) + pctUrban + fertility
     Res.Df
               RSS Df Sum of Sq
                                       F
                                            Pr(>F)
## 1
        197 6528.4
## 2
        195 5161.7
                          1366.6 25.814 1.133e-10 ***
```

Testing Whether the Parameters for log(ppgdp) and pctUrban Are Both Zero

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

1. Hypotheses

- Null Hypothesis (H_0): The parameters for log(ppgdp) and pctUrban are both 0. These variables do not contribute to explaining the variability in lifeExpF.
- Alternative Hypothesis (H_a) : At least one of the parameters for log(ppgdp) or pctUrban is not 0. Including these variables significantly improves the model.
- 2. Test Statistic The test statistic is the F-statistic from the ANOVA output:

$$F = 25.814$$

- **Degrees of Freedom**: Numerator (df_1) : 2 (number of predictors added: log(ppgdp) and pctUrban). Denominator (df_2) : 195 (residual degrees of freedom from the full model).
- **3. Significance Level** The significance level (α) is 0.05.
- **4. p-value** From the ANOVA output:

$$p = 1.133 \times 10^{-10}$$

5. Decision Compare the p-value to the significance level ($\alpha = 0.05$):

$$p = 1.133 \times 10^{-10} < 0.05$$

Since the p-value is much smaller than 0.05, we reject the null hypothesis.

6. Conclusion At the 0.05 significance level, there is sufficient evidence to conclude that the parameters for log(ppgdp) and/or pctUrban are not 0. Adding these predictors significantly improves the model's ability to explain lifeExpF.