Econometrics Case Project

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Solution for (a):

The likelihood ratio tests can be given by the test statistic

$$LR = -2(\log(L(b_0)) - \log(L(b_1))) \sim \chi^2(m)$$

where $L(b_0)$ is the maximum likelihood value in the restricted model, $L(b_1)$ is the maximum likelihood in the full model and m is the number of parameter restrictions in the restricted model. The critical value of the test is $\chi^2_{0.05}(2) = 5.991$,

 H_0 : the m parameter restrictions are correct

```
> # model with constant (m = 2 restrictions) vs full model - c
ritical value=5.991
> LR_full <- -2*(-152.763 - (-134.178)) ; LR_full
[1] 37.17
>
> # model with li1 (m =1 restrictions) vs full model - critica
l value = 3.841
> LR_1 <- -2*(-139.747 - (-134.178)) ; LR_1
[1] 11.138
> # model with li2 (m=1 restrictions) vs full model
> LR_2 <- -2*(-149.521 - (-134.178)) ; LR_2
[1] 30.686</pre>
```

For the model with two restrictions (li1=li2=0, m=2) we got LR = 37.17 > 5.991 so we reject the null hypothesis (at 5% significance level) and thus the full model is statistically significant.

For the model with restriction li2=0 (m=1), we got LR=11.138 > 3.841 (3.841 is the critical value for m=1) and so we reject the null hypothesis (at 5% significance level).

For the model with restriction li1=0 (m=1), we got LR=30.686>3.841, and so we reject the null hypothesis at 5% significance level.

Solution for (b):

We can use the McFadden \mathbb{R}^2 because the dependent variable (predictor) of the model is a nominal variable. The R squared is given by the formula:

$$R^2 = 1 - \frac{\log(L(b))}{\log(L(b_1))}$$

where b1 is the model with only the intercept and b is the model under consideration.

```
> Rsq_li11 <- 1 - (-134.178)/(-152.763) ; Rsq_li11
[1] 0.121659
> Rsq_li12 <- 1 - (-134.126)/(-152.763) ; Rsq_li12
[1] 0.1219994
> Rsq_li21 <- 1 - (-130.346)/(-152.763) ; Rsq_li21
[1] 0.1467436
> Rsq_li22 <- 1 - (-130.461)/(-152.763) ; Rsq_li22
[1] 0.1459908</pre>
```

The R squared for li1(-1) and li2(-1) is 0.1216.

The R squared for li1(-1) and li2(-2) is 0.1219.

The R squared for li1(-2) and li2(-1) is 0.1467.

The R squared for li1(-2) and li2(-2) is 0.1459.

We got that the highest R squared is 0.1467 which corresponds to the model with variables li1(-2) and li2(-1).

Solution for (d):

We denote $Y_t = LOGGDP$. The model for the Augmented Dickey-Fuller test is the following:

$$\Delta Y_t = \alpha + \beta t + \rho Y_{t-1} + \gamma \Delta Y_{t-1} + \omega_t$$

```
Call:
lm(formula = DYt \sim T + Yt1 + DYt_Lag1, data = df_new)
Residuals:
Min 1Q Median 3Q
-0.0113538 -0.0024430 -0.0001446 0.0027148
                                                  0.0182796
Coefficients:
               Estimate Std. Error t value Pr(>|t|)
                          3.724e-02
2.573e-05
8.073e-03
                                       2.509
2.481
-2.479
              9.344e-02
(Intercept)
              6.382e-05
Yt1
             -2.001e-02
                                       12.027
              6.090e-01
                           5.064e-02
DYt_Lag1
                 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Signif. codes:
Residual standard error: 0.004542 on 236 degrees of freedom
Multiple R-squared: 0.3976,
                                 Adjusted R-squared:
F-statistic: 51.91 on 3 and 236 DF, p-value: < 2.2e-16
```

The estimated model is $\Delta Y_t = 0.093 + 0.000 * t - 0.02 * Y_{t-1} + 0.609 \Delta Y_{t-1} + \omega_t$

The test statistic is $t_{\rho}=-2.479>-3.5\,$ so we do not reject the null hypothesis ($H_0:$ the time series Y_t is non-stationary) at 5% significance level.

Solution for (e):

For k1=k2=1 we get the following:

```
Call:
lm(formula = GrowthRate ~ GrowthRate_Lag1 + li1_Lag_k1 + li2_L
ag_k2,
    data = df_new)
Residuals:
Min 1Q Median 3Q
-0.0097644 -0.0028559 -0.0001994 0.0026151
                                              0.0158224
Coefficients:
                  Estimate Std. Error t value Pr(>|t|)
                 1.737e-03 3.197e-04
                                         5.433 1.38e-07 ***
(Intercept)
                                               < 2e-16 ***
                                         9.556
GrowthRate_Lag1 4.616e-01
                            4.830e-02
                                        -7.880 1.20e-13 ***
li1_Lag_k1
                -1.023e-03
                           1.298e-04
1i2_Lag_k2
                -1.494e-04 6.421e-05
                                       -2.326 0.0209 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
Residual standard error: 0.004106 on 236 degrees of freedom
Multiple R-squared: 0.508,
                               Adjusted R-squared: 0.5017
F-statistic: 81.22 on 3 and 236 DF, p-value: < 2.2e-16
```

R squared = 0.508 (k1=k2=1)

For k1=1 and k2=2:

```
Call:
lm(formula = GrowthRate ~ GrowthRate_Lag1 + li1_Lag_k1 + li2_L
ag_k2,
    data = df_new
Residuals:
                          Median
                   10
-0.009785 -0.002792 -0.000221 0.002526 0.015915
Coefficients:
                   Estimate Std. Error t value Pr(>|t|) 1.754e-03 3.188e-04 5.501 9.79e-08
(Intercept)
                                              5.501 9.79e-08 ***
GrowthRate_Lag1 4.578e-01 4.845e-02
                                              9.449 < 2e-16 ***
                               1.296e-04
                                             -7.860 1.37e-13 ***
lil Lag kl
                  -1.018e-03
                  -1.471e-04
                                6.416e-05
                                                       0.0227 *
1i2_Lag_k2
                                            -2.293
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.004107 on 236 degrees of freedom
Multiple R-squared: 0.5077, Adjusted R-squared: 0.5
F-statistic: 81.12 on 3 and 236 DF, p-value: < 2.2e-16
                                   Adjusted R-squared: 0.5014
```

R squared = 0.5077 (for k1=1 k2=2)

For k1=2, and k2=1 we get the following results:

```
Call:
lm(formula = GrowthRate ~ GrowthRate_Lag1 + li1_Lag_k1 + li2_L
ag_k2,
    data = df_new
Residuals:
                  1Q
                        Median
      Min
                                                 Max
-0.009053 -0.002661 -0.000229 0.002342 0.016491
Coefficients:
                   Estimate Std. Error t value Pr(>|t|)
                             3.376e-04
                                           5.611 5.60e-08 ***
(Intercept)
                  1.894e-03
                                           7.546 9.70e-13 ***
GrowthRate_Lag1 4.109e-01
                              5.445e-02
li1_Lag_k1
                                          -6.674 1.76e-10 ***
                 -9.871e-04
                              1.479e-04
                 -1.585e-04 6.675e-05
1i2_Lag_k2
                                          -2.374 0.0184 *
                 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Signif. codes:
Residual standard error: 0.004233 on 236 degrees of freedom
Multiple R-squared: 0.4772, Adjusted R-squared: 0.4 F-statistic: 71.8 on 3 and 236 DF, p-value: < 2.2e-16
                                 Adjusted R-squared: 0.4705
```

The R squared is 0.4772 (for k1=2, k2=1).

And finally for k1=k2=2:

```
call:
lm(formula = GrowthRate ~ GrowthRate_Lag1 + li1_Lag_k1 + li2_L
ag_k2,
    data = df_new
Residuals:
       Min
                   1Q
                          Median
                                                   Max
-0.0091209 -0.002615\overline{5} -0.0002767 0.0023941 0.0161471
Coefficients:
                  Estimate Std. Error t value Pr(>|t|)
                 1.911e-03 3.369e-04
                                        5.673 4.10e-08 ***
(Intercept)
                4.069e-01 5.466e-02
                                       7.444 1.82e-12 ***
GrowthRate_Lag1
                                       -6.662 1.88e-10 ***
li1_Lag_k1
                -9.833e-04
                           1.476e-04
                -1.579e-04 6.667e-05
                                              0.0187 *
1i2_Lag_k2
                                       -2.368
                0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Signif. codes:
Residual standard error: 0.004233 on 236 degrees of freedom
Multiple R-squared: 0.4771,
                              Adjusted R-squared: 0.4705
F-statistic: 71.78 on 3 and 236 DF, p-value: < 2.2e-16
```

R squared = 0.4771 (k1=k2=2)

From the above we conclude that the model with largest R squared (equal to 0.508) is the model for k1=k2=1. The model can be written as:

```
GrowthRate_t = \alpha + \rho GrowthRate_{t-1} + \beta_1 li1_{t-k_1} + \beta_2 li2_{t-k_2} + \varepsilon_t
```

where $\alpha = 0.001737$, $\rho = 0.4616$, $\beta_1 = -0.001023$ and $\beta_2 = 0.000149$.

Solution for (f):

We take the residuals from the model (with k1=k2=1)

$$GrowthRate_t = \alpha + \rho GrowthRate_{t-1} + \beta_1 li1_{t-1} + \beta_2 li2_{t-1} + \varepsilon_t$$

and regress them (OLS) on the following model (first order serial correlation):

$$e_t = \alpha + \rho GrowthRate_{t-1} + \beta_1 li1_{t-1} + \beta_2 li2_{t-1} + \gamma_1 e_{t-1} + \omega_t$$

```
Call:
lm(formula = e ~ GrowthRate_Lag1 + li1_Lag_k1 + li2_Lag_k2 +
     e_Lag1, data = df_new)
Residuals:
Min 1Q Median 3Q
-0.0096882 -0.0028198 -0.0002075 0.0026143
Coefficients:
                       Estimate Std. Error t value Pr(>|t|)
(Intercept)
                                    3.838e-04
GrowthRate_Lag1
                                    7.915e-02
                      2.914e-02
  1_Lag_k1
                      2.768e-06
                                    6.496e-05
1i2_Lag_k2
e_Lag1
                     -4.999e-02
Residual standard error: 0.004121 on 234 degrees of freedom
(1 observation deleted due to missingness)
Multiple R-squared: 0.001002, Adjusted R-squared: -0.01607
F-statistic: 0.05868 on 4 and 234 DF, p-value: 0.9936
```

We got that the R squared is equal to 0.001002. Now we calculate the test statistic of the Breusch-Godfrey test, that is $BG = nR^2$ and compare it with the critical value $\chi^2_{0.05}(1) = 3.841$ (at 5% significance level)

$$BG = nR^2 = 239 * 0.001 = 0.239 < 3.841$$

and so we do not need to adjust the model further.