

## **TEXAS A&M UNIVERSITY**

DEPARTMENT OF STATISTICS

# STAT 608 - Regression Analysis Homework VII

Salih Kilicli

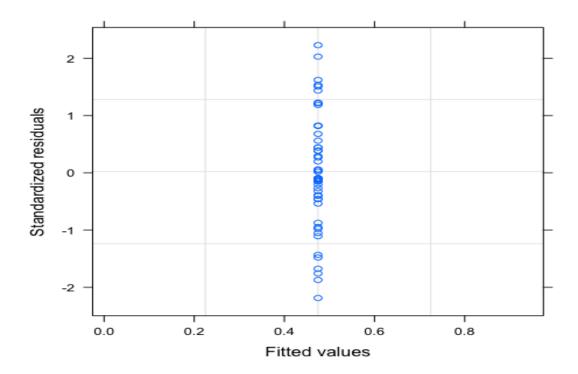
July 20, 2019

#### Question 1:

Solution: (1.1) Testing Null Hypothesis ignoring the possibility of serial correlation, we fail to reject it since the p-value of the model is given by p=0.128>0.05. The summary of the model is given below:

(1.2) Testing Null Hypothesis after checking for correlation we reject it since the p-value of the model is given by p = 0.007 < 0.05. The summary and plot of the model is given below:

```
Generalized least squares fit by maximum likelihood
 Model: x \sim 1
 Data: Q1
      AIC
            BIC logLik
 207.9588 213.6949 -100.9794
Correlation Structure: AR(1)
Formula: ~day
Parameter estimate(s):
      Phi
-0.5279798
Coefficients:
               Value Std.Error t-value p-value
(Intercept) 0.4746875 0.1710985 2.774352 0.0078
Standardized residuals:
                              Med
                                            Q3
-2.18754096 -0.45944270 -0.09817887 0.53185051 2.22976090
Residual standard error: 2.139934
Degrees of freedom: 50 total; 49 residual
```



## Question 2:

## Solution:

(2.1) The plot of the series of D-values is given below.



Since 1 year consists of approximately 52 weeks. 2 years is 104 weeks, 3 years is 156 weeks, 4 years is 208 weeks and 5 years is 260 weeks etc. The plot therefore shows a periodicity of approximately 1 year.

(2.2) First, ignoring the auto-correlation I fitted the model, and the summary is given below. Coefficient estimates without AR(1) adjustment are:

$$\hat{\beta}_0 = 0.10455, \ \hat{\beta}_1 = 0.09822, \ \hat{\beta}_2 = 1.39739$$

.

```
Generalized least squares fit by maximum likelihood
 Model: D \sim x1 + x2
 Data: Q1
             BIC logLik
      AIC
 283.4914 295.8781 -136.7457
Correlation Structure: ARMA(1,0)
 Formula: ~Week
 Parameter estimate(s):
Phi1
Coefficients:
               Value Std.Error t-value p-value
(Intercept) 0.1045455 0.1241412 0.842150 0.4021
×1
   0.0982249 0.1755622 0.559488 0.5773
           1.3973928 0.1755622 7.959532 0.0000
Correlation:
  (Intr) x1
×1 0
x2 0
Standardized residuals:
                         Med
            Q1
      Min
                                                     Max
-3.25078410 -0.62027157 0.07894603 0.72972181 1.67325607
Residual standard error: 1.144525
Degrees of freedom: 88 total; 85 residual
```

After adjustment of AR(1), the new fitted model gives parameters estimate  $\phi=0.3996349$ . The summary of AR(1) model given below:

(2.3) Looking at the summary and the plot given below, the model looks valid, since Std residuals vs Fitted Values shows a random pattern and the p-values aren't so big.

Generalized least squares fit by maximum likelihood Model: D ~ x1 + x2
Data: Q2
AIC BIC logLik
268.3022 280.6889 -129.1511

Correlation Structure: AR(1)
Formula: ~1:88
Parameter estimate(s):
Phi
0.3996349

#### Coefficients:

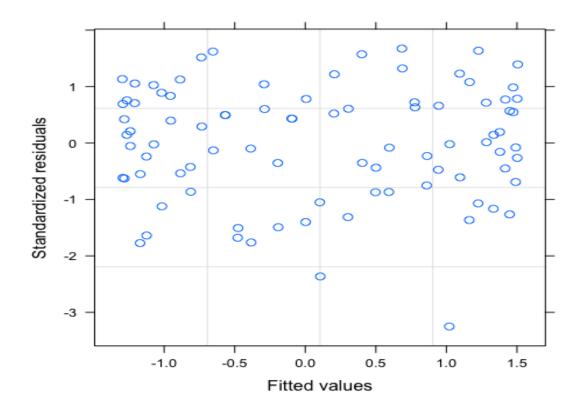
Value Std.Error t-value p-value (Intercept) 0.0918267 0.1881097 0.488155 0.6267 x1 0.1005348 0.2511485 0.400300 0.6899 x2 1.3746588 0.2479961 5.543067 0.0000

Correlation: (Intr) x1 x1 -0.003 x2 -0.019 -0.005

Standardized residuals:

Min Q1 Med Q3 Max -3.23031329 -0.62898246 0.08931067 0.74240666 1.68990576

Residual standard error: 1.144196 Degrees of freedom: 88 total; 85 residual



- Question 3: Exercise 8.3.1, page 294, in the textbook. (Note: "Traditional" linear regression methodology won't get you anywhere with this question, but a logistic regression approach will.)
  - Solution: (3.1) The author used the traditional least squares model regression on a binomial response variable. The validity of this model is therefore threatened by the fact that the mean of y, E[y] = p(x), therefore the variance of the response variable y, Var(y) = p(x)(1-p(x)) is not constant (depends on x, predictor variable) and is unknown.
    - (3.2) Fitting a logistic regression model to data we see that there is a strong evidence of a relationship between Y and x, since the estimates have highly significant coefficients. The summary of the model is given below:

```
Call:
glm(formula = cbind(playoff, noplayoff) \sim x, family = binomial)
Deviance Residuals:
  Min 1Q Median 3Q
                               Max
-2.4876 -2.0968 -0.4703 1.0666 5.3057
Coefficients:
        Estimate Std. Error z value Pr(>|z|)
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' '1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 124.10 on 29 degrees of freedom
Residual deviance: 116.22 on 28 degrees of freedom
AIC: 170.33
Number of Fisher Scoring iterations: 4
```

Question 4:

Solution: 
$$4.1 \log \left( \frac{p(3)}{1 - p(3)} \right) = -2.643 + 0.674 \times 3 = -0.621$$
 implies;

$$\left(\frac{p(3)}{1-p(3)}\right) = exp(-0.621) = 0.5374068$$

Solving the equation given above for p(3) we get:

$$p(3) = \frac{0.5374068}{1 + 0.5374068} = 0.3495541 \approx 0.35 = \frac{35}{100}$$

Therefore for 200 insect exposed to the dosage level x=3,  $200 \times p(3) = 70$  insects are expected to die. Consequently, 130 of them will survive.

4.2 
$$\frac{p(x)}{1-p(x)} = e^{-2.643+0.674(x+1)} = e^{-2.643+0674x}e^{0.674} = e^{-2.643+0674x}\phi$$

Therefore,  $\phi=e^{\hat{eta}_1}=e^{0.674}=1.96207$  is the estimated factor.

4.3 A 90% confidence interval for  $\beta_1$  is given by:

$$\hat{\beta}_1 \pm z_{0.95} se(\hat{\beta}_1) = 0.674 \pm 1.644854(0.039) = (0.6098507, 0.7381493)$$

Therefore, a 90% confidence interval for  $\phi$  can be found as:

$$(e^{0.6098507}, e^{0.7381493}) = (1.840157, 2.09206)$$

#### Question 5:

Solution: (5.1)  $V_i = arcsin(\sqrt{z_i}), \quad z_i = \frac{y_i}{m_i}$  where  $E[y_i] = m_i\theta_i, \quad Var(y_i) = m_i\theta_i(1-\theta_i).$  Therefore we have:

$$E[z_i] = \theta_i, \quad Var(z_i) = \frac{\theta_i(1 - \theta_i)}{m_i}$$

Now, from section 3.3.1 we have: (Note that,  $dV_i/dz_i = (2\sqrt{z_i(1-z_i)})^{-1}$ )

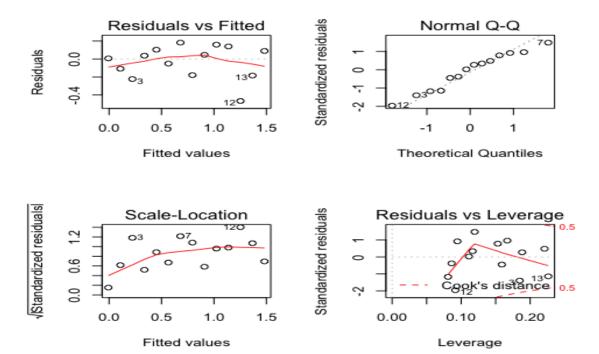
$$Var(V_i) \approx \left[ \frac{dV_i}{dz_i} (E[z_i]) \right]^2 Var(z_i)$$

$$\approx \left[ \frac{1}{2\sqrt{E[z_i](1 - E[z_i])}} \right]^2 Var(z_i)$$

$$\approx \frac{1}{4\theta_i(1 - \theta_i)} \frac{\theta_i(1 - \theta_i)}{m_i}$$

$$\approx \frac{1}{4m_i}$$

- (5.2)  $\hat{\gamma_0} = -1.72348$ ,  $\hat{\gamma_1} = 0.11447$ ,  $S^2 = (0.2967)^2 = 0.8804$
- (5.3) All of the estimates are highly significant,  $R^2=0.8563$ , the F-statistic of the model is 71.5 with a highly significant p=2.122e-06 value. o, the model is valid; however, the Residual vs Fitted and scale-location plots show slightly increasing variance and non-linear relationship, so model definitely can be improved by other means.



(5.4) 90% prediction interval for  $V_i$ \* is:

$$V_i* = sin^{-1}(\sqrt{z_i*}) \in (0.145305, 1.215334)$$
 fitted on 0.683195.

Therefore, 90% prediction interval for  $z_i*$  can be found as:

$$z_i * = sin^2(V_i *) \in \left(sin^2(0.145305), sin^2(1.215334)\right)$$
  
  $\in (0.0203241, 0.8788794)$ 

### Question 6:

Solution: (a) Using logistic regression we get the parameter estimates as below:

$$\hat{\beta}_0 = 15.07, \ \hat{\beta}_1 = -113.63, \ \hat{\beta}_2 = 255.29, \ \hat{\beta}_3 = -183.14$$

(b) There are two ways to answer this question. First, simply looking at Wald p-value of  $\hat{\beta}_3$  given in the summary 0.00316 < 0.1 we see that the coefficient is significant therefore we reject the Null Hypothesis. Alternatively, using the difference of deviances and calculating p-value from  $\chi^2$  distribution with 1 dof we get:

$$P\{G_{H_0}^2 - G_{H_A}^2 > 11.68\} = 0.0006333219 < 0.1$$

Therefore, we reject the Null Hypothesis.