

Time: 8.00-13.00. Limits for the credits 3, 4, 5 are 25, 30 and 35 points, respectively. The solutions should be well motivated.

Permitted aids: One page (both sides) hand-written cheat sheet for the course. Pocket calculator.

1. (8p) Suppose that we have a data set of three observations given by

$$\begin{array}{cc} y & x \\ 0 & -1 \\ 0 & 0 \\ 3 & 1 \end{array}$$

. We want to fit a simple linear regression $y_i = \beta_0 + \beta_1 x_i + e_i$.

- (a) (2p) Find the ordinary least squares estimates of β_0 and β_1 .

Solution: The OLS estimators are

$$\hat{\beta}_1 = \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\sum_{i=1}^n (x_i - \bar{x})^2},$$
$$\hat{\beta}_0 = \bar{y} - \hat{\beta}_1 \bar{x}.$$

Direct calculation shows that $\hat{\beta}_1 = 1.5$ and $\hat{\beta}_0 = 1$.

- (b) (2p) What are the fitted values for this data set?

Solution: The fitted values are -0.5 , 1.0 , and 2.5 .

- (c) (2p) Find the value of the residual sum of squares.

Solution: The residuals are 0.5 , -1.0 and 0.5 . The residual sum of squares is 1.5 .

- (d) (2p) The variance of e_i given x_i is σ^2 . Estimate σ^2 .

Solution: $\hat{\sigma}^2 = RSS / (n - 2) = 1.5$.

2. (8p) Consider the model $y_i = \mathbf{x}_i^T \boldsymbol{\beta} + e_i$, where $\text{Var}(e_i | \mathbf{x}_i) = \sigma^2 (\mathbf{x}_i^T \mathbf{x}_i + 1)$ and $\boldsymbol{\beta}$ is a $p \times 1$ vector. We also assume that observations are independent of each other.

- (a) (2p) Derive the ordinary least squares estimator.

Solution: The residual sum of squares is

$$\text{RSS}(\boldsymbol{\beta}) = (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^T (\mathbf{y} - \mathbf{X}\boldsymbol{\beta}).$$

The OLS estimator is still

$$\hat{\boldsymbol{\beta}} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}.$$

- (b) (2p) Is your ordinary least squares estimator unbiased for $\boldsymbol{\beta}$ if the model is correctly specified?

Solution:

$$\begin{aligned} E[\hat{\boldsymbol{\beta}} | \mathbf{X}] &= E[(\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y} | \mathbf{x}] \\ &= (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T E[\mathbf{y} | \mathbf{x}] \\ &= (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{X} \boldsymbol{\beta} \\ &= \boldsymbol{\beta}. \end{aligned}$$

- (c) (2p) Find the covariance matrix of your estimator.

Solution:

$$\text{Var}(\hat{\boldsymbol{\beta}} | \mathbf{X}) = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \text{Var}(\mathbf{y} | \mathbf{X}) \mathbf{X} (\mathbf{X}^T \mathbf{X})^{-1}.$$

- (d) (2p) Do the residuals have a zero sample covariance with the predicted value given \mathbf{X} , if the sample mean of the residuals is zero?

Solution:

$$\frac{1}{n} \hat{\mathbf{e}}^T \hat{\mathbf{y}} = \frac{1}{n} \mathbf{y}^T (\mathbf{I} - \mathbf{H}) \mathbf{X} (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y} = \mathbf{0}.$$

Hence, the sample covariance is zero.

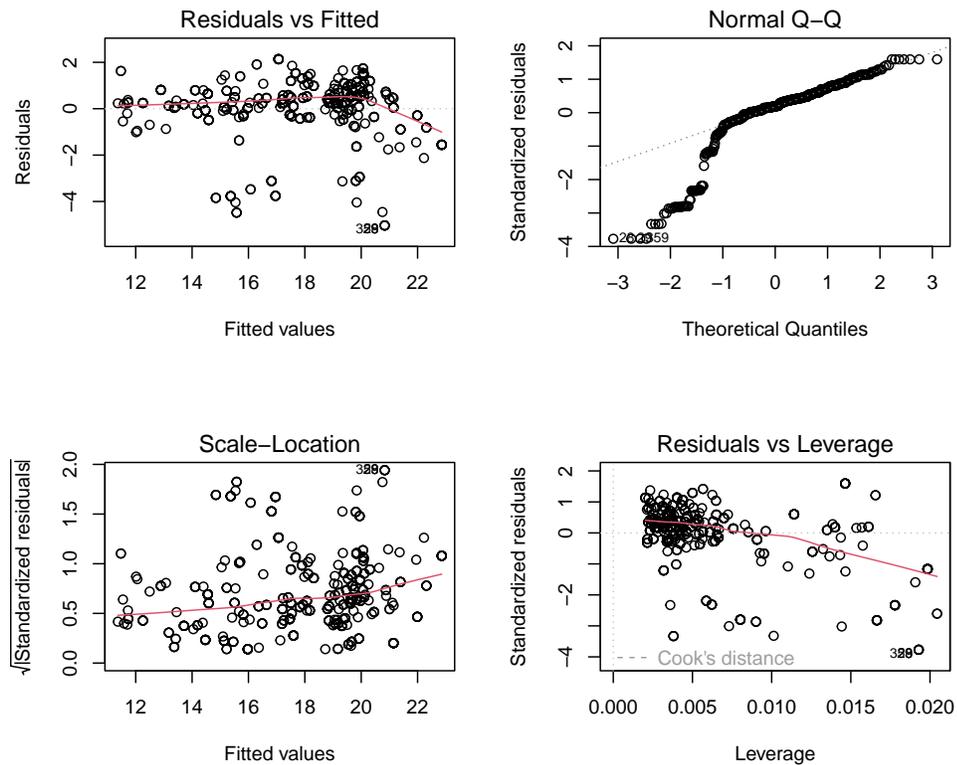
3. (12p) Suppose that we have measured the weight of water, the weight of fat, and the weight of protein of one type of meat. We have regressed the weight of protein on the weight of water and the weight of fat in R as follows.

```
LR <- lm(Protein ~ Water + Fat, data = Data)
summary(LR)

##
## Call:
## lm(formula = Protein ~ Water + Fat, data = Data)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
```

```
## -5.0331 -0.2616  0.2483  0.7249  2.1343
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 62.28170    2.98936   20.83  <2e-16 ***
## Water      -0.52985    0.03869  -13.69  <2e-16 ***
## Fat        -0.61068    0.03023  -20.20  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.348 on 497 degrees of freedom
## Multiple R-squared:  0.7827, Adjusted R-squared:  0.7818
## F-statistic:   895 on 2 and 497 DF,  p-value: < 2.2e-16
```

- (a) (1p) Does Water significantly affect Protein when controlling for Fat?
Solution: Yes, since the p-value is $< 2e - 16$.
- (b) (1p) What is the fitted regression model?
Solution: The fitted model is $\hat{y} = 62.28170 - 0.52985\text{Water} - 0.61068\text{Fat}$
- (c) (1p) How would you interpret the effect of Fat?
Solution: For a fixed level of water, one unit increase in Fat is expected to decrease Protein by 0.61068 units.
- (d) (2p) Interpret Multiple R-Squared.
Solution: 78.27% of the variation in Protein has been explained by the linear model.
- (e) (2p) Construct a 95% confidence interval for the regression coefficient for Fat. Use the notation $t_\alpha(a)$ to denote the quantile of a t distribution with a degrees of freedom.
Solution: The confidence interval is $-0.61068 \pm t_{0.975}(500 - 3)0.03023$.
- (f) (2p) Do the residuals of this model have the zero sample mean?
Solution: The residual vector is $\hat{\mathbf{e}} = \mathbf{y} - \mathbf{X}\hat{\boldsymbol{\beta}} = (\mathbf{I} - \mathbf{H})\mathbf{X}$. We can show that $\mathbf{X}^T(\mathbf{I} - \mathbf{H})\mathbf{X} = \mathbf{0}$. Hence, the sample mean is zero if the intercept is included in the model.
- (g) (2p) The residual plots of the fitted model is



What conclusions can you draw from the residual plots?

Solution: The variation of the residuals depends on the fitted values, so the homoscedasticity assumption may be violated. The QQ plot suggests that the residuals may not be normally distributed.

(h) (1p) What has been calculated by the following R code

```
predict(LR, newdata = data.frame(Water = 50, Fat = 20), interval = "p")
```

Solution: Prediction interval when water is 50 and Fat is 20.

4. (8p) Suppose that we have measured the weight of protein of one type of meat produced by five brands. We want to test whether different brands have the same weights. For each brand, 100 samples are measured.

(a) (2p) A statistician has done the following analysis in R.

```
LR <- lm(Protein ~ Brand, data = Data)
anova(LR)

## Analysis of Variance Table
##
## Response: Protein
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## Brand      4    57.0  14.2480   1.7207 0.1441
## Residuals 495 4098.8   8.2805
```

Do different brands have the same weight?

Solution: Yes, since the effect of Brand is not significant.

- (b) (2p) Another statistician has done the following analysis in R.

```
LR <- lm(Protein ~ Brand, data = Data)
summary(LR)

##
## Call:
## lm(formula = Protein ~ Brand, data = Data)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -7.3050 -2.0130  0.9445  2.3068  4.4020
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  18.3050     0.2878  63.612  <2e-16 ***
## Brand2       -1.0070     0.4070  -2.474   0.0137 *
## Brand3       -0.2920     0.4070  -0.718   0.4734
## Brand4       -0.2260     0.4070  -0.555   0.5789
## Brand5       -0.3470     0.4070  -0.853   0.3942
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.878 on 495 degrees of freedom
## Multiple R-squared:  0.01371, Adjusted R-squared:  0.005744
## F-statistic: 1.721 on 4 and 495 DF, p-value: 0.1441
```

Based on the analysis of the second statistician, do different brands have the same weight?

Solution: The p-value of the F test is above 0.05, hence we cannot reject H_0 all brands have the same weight.

- (c) (2p) What is the estimated difference in the weight of Protein between Brand 3 and Brand 5?

Solution: Estimated difference is $-0.2920 - (-0.3470)$.

- (d) (2p) Suppose that the statistician wants to test the statement: the weight of Protein of Brand 2 is twice the weight of Protein of Brand 3. Explain how you can perform such test. Note: You don't need to compute the value of your test statistics.

Solution: An F test for linear combination can be tested. The linear combination is

$$\beta_0 + \beta_1 = 2(\beta_0 + \beta_2)$$

or equivalently

$$[1 \quad -1 \quad 2 \quad 0 \quad 0] \begin{bmatrix} \beta_0 \\ \beta_1 \\ \beta_2 \\ \beta_3 \\ \beta_4 \end{bmatrix} = 0.$$

We will fit a model with the above linear restriction and compute the F value using The F test is

$$F = \frac{(\text{RSS}_0 - \text{RSS}_1) / (p - p_0)}{\text{RSS}_1 / (n - p)} \sim F(p - p_0, n - p).$$

where $n - p = 495$, and $p - p_0 = 1$.

5. (4p) We have a data set from a study of income dynamics of married women in the US. The variables in the data set are lfp (labor-force participation; a factor with levels: no; yes), k5 (number of children 5 years old or younger), age (age of the woman in years), and inc (family income exclusive of wife's income). A has been fitted as follows.

```
Logit <- glm(lfp ~ k5 + age + inc, family = binomial(link = "logit"),
             data = Mroz)
summary(Logit)

##
## Call:
## glm(formula = lfp ~ k5 + age + inc, family = binomial(link = "logit"),
##      data = Mroz)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.867  -1.184   0.731   1.003   1.970
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  3.394398   0.515576   6.584 4.59e-11 ***
## k5          -1.313316   0.187535  -7.003 2.50e-12 ***
## age         -0.056855   0.010991  -5.173 2.31e-07 ***
## inc         -0.018751   0.006889  -2.722 0.00649 **
```

```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 1029.75  on 752  degrees of freedom
## Residual deviance:  956.75  on 749  degrees of freedom
## AIC: 964.75
##
## Number of Fisher Scoring iterations: 4
```

- (a) (2p) What is the estimated probability of labor-force participation for a 30 years old women with 0 child 5 years old or younger, and the family income exclusive of wife's income is 10? It suffices to present the formula without presenting the final number.

Solution: Let $\eta = 3.394398 - 0.056855 \cdot 30 - 0.018751 \cdot 10$. The estimated probability is

$$\frac{\exp(\eta)}{1 + \exp(\eta)}$$

- (b) (2p) A second model has been fitted as follows.

```
LR <- lm(lfp ~ k5 + age + inc, data = Mroz)
```

Which model is more plausible? Motivate your answer.

Solution: The logistic regression is more plausible since a linear model does not regulate the predicted values to be proper probabilities.