MAST30025 (620-328) Linear Statistical Models

Semester 1 Exam, 2011

Department of Mathematics and Statistics
The University of Melbourne

Exam duration: 3 hours
Reading time: 15 minutes
This exam has 7 pages, including this page.

Authorised materials:

Scientific calculators are permitted, but not graphical calculators.

One A4 double-sided handwritten sheet of notes.

Instructions to invigilators:

The exam paper may be taken out of the examination room.

Instructions to students:

There are 6 questions. All questions should be attempted.

The number of marks for each question is indicated.

The total number of marks available is 80.

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1. [13 marks] Consider the column vectors

$$\left[\begin{array}{c}1\\2\\2\end{array}\right], \left[\begin{array}{c}2\\-2\\1\end{array}\right], \left[\begin{array}{c}2\\1\\-2\end{array}\right].$$

- (a) [1 mark] Show that these vectors are mutually orthogonal.
- (b) [1 mark] What constant c makes the following matrix orthogonal?

$$P \doteq c \left[\begin{array}{ccc} 1 & 2 & 2 \\ 2 & -2 & 1 \\ 2 & 1 & -2 \end{array} \right]$$

(c) [2 marks] Let $A = P\Lambda P^T$, where

$$\Lambda = \left[\begin{array}{ccc} 2 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 0 \end{array} \right].$$

What are the rank r(A) and trace tr(A)?

(d) [3 marks] Show that

$$A^c = P \left[\begin{array}{rrr} 1/2 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 0 \end{array} \right] P^T$$

is a conditional inverse of A.

- (e) [3 marks] Find a conditional inverse of $A^T A$.
- (f) [3 marks] Show that for your choice of $(A^TA)^c$ above, $A(A^TA)^cA^T$ is symmetric and idempotent (by direct calculation or otherwise).
- 2. [10 marks] Consider the following ANCOVA model, with a single factor and a single regression variable x,

$$y_{ij} = \mu + \tau_i + \gamma x_{ij} + \epsilon_{ij}$$

Suppose that the factor has two levels, and that for each level there are two observations. Also suppose that $\sum_{j} x_{1j} = 0$, $\sum_{j} x_{2j} = 1$, and $\sum_{i,j} x_{ij}^2 = 3$.

- (a) [1 mark] What are the parameter vector β and the design matrix X for this model?
- (b) [3 marks] Write down X^TX and hence show that it has rank $r(X^TX) = 3$ (provided the x_{ij} are not pathological).
- (c) [3 marks] Give a conditional inverse for X^TX . You may use the fact that, when it exists,

$$\begin{bmatrix} x & 0 & a \\ 0 & y & b \\ a & b & c \end{bmatrix}^{-1} = \frac{1}{cxy - a^2y - b^2x} \begin{bmatrix} cy - b^2 & ab & -ay \\ ab & cx - a^2 & -bx \\ -ay & -bx & xy \end{bmatrix}.$$

(d) [3 marks] Give a solution to the normal equations.

3. [13 marks] The following data concerns population growth in Taiwan.

```
> Taiwan <- data.frame(year = 40:46, growth = c(1.62, 1.63, 1.9,
```

> summary(model)

Call:

lm(formula = growth ~ year, data = Taiwan)

Residuals:

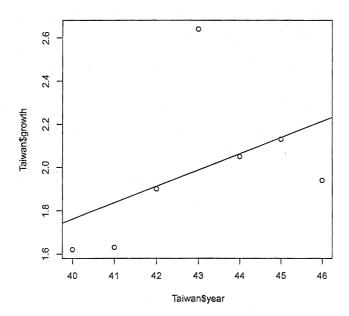
Coefficients:

Estimate Std. Error t value Pr(>|t|)
(Intercept) -1.25321 2.73157 -0.459 0.666
year 0.07536 0.06346 1.188 0.288

Residual standard error: 0.3358 on 5 degrees of freedom Multiple R-squared: 0.22, Adjusted R-squared: 0.064 F-statistic: 1.41 on 1 and 5 DF, p-value: 0.2883

> plot(Taiwan\$year, Taiwan\$growth)

> abline(coef = model\$coefficients)

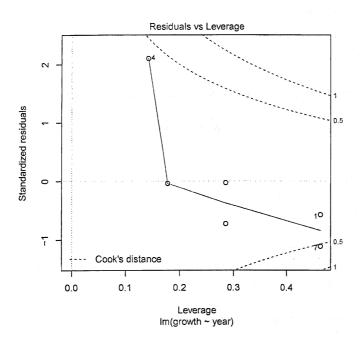


> plot(model, which = 5)

[5,]

[6,] [7,] 0.25000000 0.35714286

0.46428571



- (a) [6 marks] The second plot indicates there may be some outliers. Define standardised residuals, leverage and Cook's distance, and briefly explain how they are used for regression diagnostics.
- (b) [4 marks] Using the output above and below, calculate the Cook's distance for observation 7. (Do not just read it from the graph.)

```
> X <- matrix(nrow = 7, ncol = 2)
> X[, 1] <- 1
> X[, 2] <- Taiwan$year
> (H <- X %*% solve(t(X) %*% X) %*% t(X))
            [,1]
                          [,2]
                                      [,3]
                                                [,4]
                                                           [,5]
                                                                          [,6]
                  3.571429e-01 0.25000000 0.1428571 0.03571429 -7.142857e-02
[1,]
     0.46428571
     0.35714286 2.857143e-01 0.21428571 0.1428571 0.07142857 -6.938894e-16
                  2.142857e-01 0.17857143 0.1428571 0.10714286
     0.25000000
                  1.428571e-01 0.14285714 0.1428571 0.14285714
[4,]
     0.14285714
                  7.142857e-02 0.10714286 0.1428571 0.17857143
     0.03571429
[6,] -0.07142857 -7.216450e-16 0.07142857 0.1428571 0.21428571
                                                                 2.857143e-01
[7,] -0.17857143 -7.142857e-02 0.03571429 0.1428571 0.25000000
            [,7]
[1,] -0.17857143
[2,] -0.07142857
[3,]
     0.03571429
[4,]
     0.14285714
```

(c) [3 marks] Give a joint 95% confidence region for the intercept and slope. You may express your region as an inequality in matrix form, and note that the upper 5% point for an $F_{2,5}$ distribution is 5.79.

- 4. [22 marks] Suppose that $y \sim MVN(\mu, I_n)$, and that A is an $n \times n$ symmetric idempotent matrix of rank k.
 - (a) [6 marks] Show that the rank equals the trace r(A) = tr(A).
 - (b) [6 marks] Show that $\mathbf{y}^T A \mathbf{y} \sim \chi^2_{k,\mu^T A \mu/2}$.
 - (c) [10 marks] Suppose that B is symmetric and AB = 0. Show that $\mathbf{y}^T A \mathbf{y}$ and $\mathbf{y}^T B \mathbf{y}$ are independent.

If in addition B is idempotent of rank h, what is the distribution of $\mathbf{y}^T A \mathbf{y} + \mathbf{y}^T B \mathbf{y}$?

5. [11 marks] Four tropical feeds were fed to baby chicks. The gains in weight (in grams) after two weeks were:

```
Feed A
           42
                 68
                       85
           42
                 97
                       81
Feed B
                             95
                                   61
                                       103
Feed C
          61
               112
                       30
                             89
                                   63
         169
               137
                     169
Feed D
                           111
                                  154
```

```
> chicks <- data.frame(gain = c(42, 68, 85, 42, 97, 81, 95, 61,
```

- + 103, 61, 112, 30, 89, 63, 169, 137, 169, 111, 154), feed = rep(c("A",
- + "B", "C", "D"), c(3, 6, 5, 5)))
- > options(contrasts = c("contr.treatment", "contr.poly"))
- > model <- lm(gain ~ feed, data = chicks)
- > summary(model)

Call:

lm(formula = gain ~ feed, data = chicks)

Residuals:

```
Min 1Q Median 3Q Max -41.00 -14.92 3.00 19.00 41.00
```

Coefficients:

```
Estimate Std. Error t value Pr(>|t|)
               65.00
                           14.94
                                    4.351 0.000571 ***
(Intercept)
                                    0.811 0.430247
feedB
                14.83
                           18.30
feedC
                 6.00
                           18.90
                                    0.317 0.755251
               83.00
                           18.90
                                    4.392 0.000525 ***
feedD
```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 25.88 on 15 degrees of freedom

Multiple R-squared: 0.6758, Adjusted R-squared: 0.6109

F-statistic: 10.42 on 3 and 15 DF, p-value: 0.0005872

- (a) [6 marks] Give formulae for the *F*-statistic and *p*-value given on the last line of output above. Take care to define all the terms you use. What do you conclude from this *p*-value?
- (b) [2 marks] Estimate the mean weight gain for a chick on feed D.
- (c) [3 marks] Can you estimate the difference between the mean weight gain for a chick on feed D, and the mean weight gain for a chick on feed A, B or C? (That is, the difference between τ_4 and $(\tau_1 + \tau_2 + \tau_3)/3$.) If so, explain why, and then give the estimate. If not, explain why not.

6. [11 marks] In an experiment to understand what makes a good cheese, a variety of cheeses were selected and subjected to a taste test by a panel of experts, who gave each cheese a numerical rating. The levels of acetic acid, hydrogen sulphide, and lactic acid were then measured for each cheese.

Here is an analysis of the data in R.

```
> cheese <- read.table("cheese.csv", sep = ",", header = T)
```

- > cheese\$ln_acetic <- log(cheese\$acetic)</pre>
- > cheese\$ln_H2S <- log(cheese\$H2S)</pre>
- > pairs(~taste + ln_acetic + ln_H2S + lactic, data = cheese)
- > full_model <- lm(taste ~ ln_acetic + ln_H2S + lactic, data = cheese)</pre>
- > summary(full_model)

Call:

lm(formula = taste ~ ln_acetic + ln_H2S + lactic, data = cheese)

Residuals:

```
Min 1Q Median 3Q Max -17.390 -6.611 -1.008 4.907 25.448
```

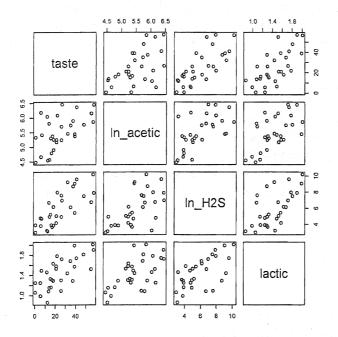
Coefficients:

Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1

Residual standard error: 10.13 on 26 degrees of freedom

Multiple R-squared: 0.6518, Adjusted R-squared: 0.6116

F-statistic: 16.22 on 3 and 26 DF, p-value: 3.81e-06



- (a) [1 mark] Why were the variables acetic and H2S transformed?
- (b) [1 mark] Write down the fitted model.
- (c) [7 marks] Suppose that a new cheese has the following measured values

$$acetic = 200, H2S = 2000, lactic = 1.5$$

For these values a 95% confidence interval for the mean taste is (25.75, 38.45). Give a 95% prediction interval for the taste of this cheese. (Note that the upper 2.5% point for a t_{26} distribution is 2.056.)

(d) [2 marks] If you were to perform one step of backward elimination, which variable would you remove, if any, and why?

End of examination



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