

MATHEMATICAL TRIPOS Part III

Thursday 30 May 2002 9 to 12

PAPER 36

APPLIED STATISTICS

*Attempt **FOUR** questions*

*There are **five** questions in total*

The questions carry equal weight

You may not start to read the questions
printed on the subsequent pages until
instructed to do so by the Invigilator.

1 (i) Define Ω as the linear model

$$\Omega : Y = \mu 1 + X\beta + \epsilon$$

where Y is an n -dimensional observation vector, 1 is the n -dimensional unit vector, μ and β are unknown parameters, X is a given $n \times p$ matrix of rank p , with $X^T 1 = \mathbf{0}$, and the components of ϵ are $\epsilon_1, \dots, \epsilon_n$, distributed as $NID(0, \sigma^2)$, with σ^2 unknown. Define further

$$X\beta = X_1\beta_1 + X_2\beta_2,$$

where X is partitioned as $(X_1 : X_2)$, and β is similarly partitioned as $\beta = \begin{pmatrix} \beta_1 \\ \beta_2 \end{pmatrix}$.

How would you test the hypothesis $\omega : \beta = 0$ against Ω ? How would you test the hypothesis $\omega_1 : \beta_1 = 0$ against Ω ? What does it mean to say that β_1, β_2 are orthogonal? (Standard theorems need not be proved but should be carefully quoted.)

(ii) Discuss carefully the S-Plus5 output for the data given below. How might you extend the analysis given?

From The Independent,

November 21, 2001, with the headline

'Supermarkets to defy bar on cheap designer goods'.

How prices compare: prices given in UK pounds.

Item	UK	Sweden	France	Germany	US
Levi 501 jeans	46.16	47.63	42.11	46.06	27.10
Dockers K1 khakis	58.00	54.08	47.22	46.20	32.22
Timberland women's boots	111.00	104.12	89.43	93.36	75.42
DieselKultar men's jeans	60.00	43.35	43.50	44.48	NA
Timberland cargo pants	53.33	48.58	43.54	58.66	31.70
Gap men's sweater	34.50	NA	26.93	27.26	28.76
Ralph Lauren polo shirt	49.99	42.04	36.41	40.26	32.48
H&M cardigan	19.99	17.31	18.17	15.28	NA

```
> p _ scan("pdata"); it _ 1:8; cou _ scan(",")
```

```
UK Swe Fra Germ US
```

```
>x _ expand.grid(cou,it) ; country _ x[,1] ; item _ x[,2]
```

```
>item _ factor(item)
```

```
> first.lm _ lm(p~ country + item,na.action=na.omit)
```

```
> anova(first.lm)
```

Analysis of Variance Table

Response: p

Terms added sequentially (first to last)

	Df	Sum of Sq	Mean Sq	F Value	Pr(F)
country	4	1115.56	278.890	10.57291	3.732294e-05
item	7	16910.20	2415.743	91.58259	0.000000e+00
Residuals	25	659.44	26.378		

```
> next.lm _ lm(p~ item + country, na.action=na.omit)
```

```
> anova(next.lm)
```

Analysis of Variance Table

Response: p

Terms added sequentially (first to last)

	Df	Sum of Sq	Mean Sq	F Value	Pr(F)
item	7	16409.02	2344.146	88.86829	0.000000e+00
country	4	1616.74	404.184	15.32293	1.859221e-06
Residuals	25	659.44	26.378		

2 (i) Let Y_1, \dots, Y_n be independent binary random variables with

$$P(Y_i = 1) = p_i = 1 - P(Y_i = 0), \quad 1 \leq i \leq n,$$

where p_1, \dots, p_n are unknown probabilities. Describe briefly how to fit the model

$$\omega : \log \frac{p_i}{1 - p_i} = \beta^T x_i, \quad 1 \leq i \leq n,$$

where x_1, \dots, x_n are given vectors, each of dimension p , and β is an unknown vector.

What is the maximised log-likelihood under the hypothesis $\Omega : 0 \leq p_i \leq 1, 1 \leq i \leq n$? Why is the usual *deviance* not appropriate as a measure of the fit of ω ?

(ii) Rousseau *et al*, 1983, collected data on males in a heart-disease high-risk region of the Western Cape, South Africa. Our object is to predict $\text{chd} = 1$ or 0 , i.e., coronary heart disease present or absent, from a set of covariates listed below

sbp	systolic blood pressure
tobacco	cumulative tobacco (kg)
ldl	low density lipoprotein cholesterol
adiposity	
famhist	family history of heart disease (Present, Absent)
typea	type-A behaviour
obesity	
alcohol	current alcohol consumption
age	age at onset

Interpret the corresponding S-Plus5 output, which makes use of the function

`stepAIC`

from library (MASS).

```
> SAheart.data[1:3,]
  sbp tobacco  ldl adiposity famhist typea obesity alcohol age chd
1 160 12.00 5.73    23.11 Present   49  25.30  97.20 52  1
2 144  0.01 4.41    28.61 Absent   55  28.87   2.06 63  1
3 118  0.08 3.48    32.28 Present   52  29.14   3.81 46  0
>table(famhist, chd)
      0  1
Absent 206 64
Present 96 96

> first.glm _ glm(chd ~ sbp+tobacco+ldl+adiposity+famhist+typea+obesity+
+ alcohol + age, family = binomial)
> summary(first.glm, cor=F)
```

Coefficients:

	Value	Std. Error	t value
(Intercept)	-6.1506610935	1.306629106	-4.70727390
sbp	0.0065040116	0.005727607	1.13555485
tobacco	0.0793762052	0.026590779	2.98510268
ldl	0.1739231824	0.059627387	2.91683387
adiposity	0.0185864751	0.029270110	0.63499847
famhist	0.9253661529	0.227736242	4.06332406
typea	0.0395947051	0.012308368	3.21689313
obesity	-0.0629099612	0.044222058	-1.42259236
alcohol	0.0001216154	0.004481130	0.02713944
age	0.0452248070	0.012115699	3.73274426

(Dispersion Parameter for Binomial family taken to be 1)

Null Deviance: 596.1084 on 461 degrees of freedom

Residual Deviance: 472.14 on 452 degrees of freedom

Number of Fisher Scoring Iterations: 4

```
> stepAIC(first.glm)
```

```
Start: AIC= 492.14
```

```
chd ~ sbp +tobacco +ldl +adiposity +famhist +typea +obesity +alcohol+
age
```

	Df	Deviance	AIC
- alcohol	1	472.1408	490.1408
- adiposity	1	472.5450	490.5450
- sbp	1	473.4371	491.4371
<none>	NA	472.1400	492.1400
- obesity	1	474.2332	492.2332
- ldl	1	481.0701	499.0701
- tobacco	1	481.6744	499.6744
- typea	1	483.0466	501.0466
- age	1	486.5284	504.5284
- famhist	1	488.8851	506.8851

```
Step: AIC= 490.14
```

```
chd ~ sbp +tobacco +ldl +adiposity +famhist +typea +obesity +age
```

	Df	Deviance	AIC
- adiposity	1	472.5490	488.5490
- sbp	1	473.4651	489.4651
<none>	NA	472.1408	490.1408
- obesity	1	474.2404	490.2404
- ldl	1	481.1541	497.1541
- tobacco	1	482.0563	498.0563
- typea	1	483.0604	499.0604
- age	1	486.6412	502.6412
- famhist	1	488.9925	504.9925

```
Step: AIC= 488.55
```

```
chd ~ sbp + tobacco + ldl + famhist + typea + obesity + age
```

	Df	Deviance	AIC
- sbp	1	473.9799	487.9799
<none>	NA	472.5490	488.5490
- obesity	1	474.6548	488.6548
- tobacco	1	482.5353	496.5353
- ldl	1	482.9470	496.9470
- typea	1	483.1925	497.1925
- famhist	1	489.3779	503.3779
- age	1	495.4754	509.4754

Step: AIC= 487.98

chd ~ tobacco + ldl + famhist + typea + obesity + age

	Df	Deviance	AIC
- obesity	1	475.6856	487.6856
<none>	NA	473.9799	487.9799
- tobacco	1	484.1760	496.1760
- typea	1	484.2967	496.2967
- ldl	1	484.5327	496.5327
- famhist	1	490.5818	502.5818
- age	1	502.1120	514.1120

Step: AIC= 487.69

chd ~ tobacco + ldl + famhist + typea + age

	Df	Deviance	AIC
<none>	NA	475.6856	487.6856
- ldl	1	484.7143	494.7143
- typea	1	485.4439	495.4439
- tobacco	1	486.0322	496.0322
- famhist	1	492.0948	502.0948
- age	1	502.3788	512.3788

Call:

```
glm(formula = chd ~tobacco +ldl +famhist +typea +age,binomial)
```

Coefficients:

(Intercept)	tobacco	ldl	famhist	typea	age
-6.446392	0.08037506	0.1619908	0.9081708	0.0371149	0.05045984

Degrees of Freedom: 462 Total; 456 Residual

Residual Deviance: 475.6856

```
>summary(glm(chd ~tobacco+ldl+famhist+typea+age,binomial),cor=F)
```

Coefficients:

	Value	Std. Error	t value
(Intercept)	-6.44639157	0.91929370	-7.012331
tobacco	0.08037506	0.02586750	3.107183
ldl	0.16199083	0.05493652	2.948691
famhist	0.90817082	0.22560312	4.025524
typea	0.03711490	0.01215529	3.053395
age	0.05045984	0.01019143	4.951201

(Dispersion Parameter for Binomial family taken to be 1)

Null Deviance: 596.1084 on 461 degrees of freedom

Residual Deviance: 475.6856 on 456 degrees of freedom

Number of Fisher Scoring Iterations: 4

3 The table below shows the number of road accidents at eight different locations, over a number of years, before and after installation of some traffic control measures. The question of interest is whether there has been a significant change in the rate of accidents. Let

y_{ij} = number of accidents in location i under ‘treatment’ j

with $j = 1$ corresponding to ‘before’, and $j = 2$ to ‘after’

installation of traffic control.

Let p_{ij} be the corresponding period of observation, so that for example $p_{11} = 9$ years, during which a total of $y_{11} = 13$ accidents were observed. (The total of ‘Before’ accidents was 114 over 68 years (rate 1.676/year), and the total of ‘after’ accidents was 15 over 18 years (rate 0.833/year).)

(i) Write down the equations to find the maximum likelihood estimates of the unknown parameters in the model in which y_{ij} are assumed independent Poisson variables with

$$\begin{aligned} \mathbb{E}(y_{ij}) &= p_{ij}\mu_{ij}, \text{ and} \\ \log \mu_{ij} &= \mu + \alpha_i + \beta_j, \quad 1 \leq i \leq 8, \quad 1 \leq j \leq 2, \end{aligned}$$

and $\alpha_1 = \beta_1 = 0$.

Indicate briefly how `glm()` solves the corresponding equations, and interpret the attached S-Plus output.

(ii) Let e_{ij} be the corresponding ‘fitted values’ in this model. Show that

$$\begin{aligned} \sum_j e_{ij} &= \sum_j y_{ij} \text{ for each } i, \text{ and} \\ \sum_i e_{ij} &= \sum_i y_{ij} \text{ for each } j. \end{aligned}$$

Location	Before		After	
	Years	Accidents	Years	Accidents
1	9	13	2	0
2	9	6	2	2
3	8	30	3	4
4	8	20	2	0
5	9	10	2	0
6	8	15	2	6
7	9	7	2	1
8	8	13	3	2

```
>summary(glm(acc ~ treat + site,poisson,offset=log(year)),cor=F)
```

```
Call:glm(formula =acc~treat+site,family=poisson,offset=log(year))
```

```
Deviance Residuals:
```

Min	1Q	Median	3Q	Max
-2.027386	-0.591431	-0.02094977	0.3122669	2.141791

```
Coefficients:
```

	Value	Std. Error	t value
(Intercept)	0.2707792	0.2784869	0.9723229
treat	-0.7806616	0.2751810	-2.8369024
site2	-0.4855078	0.4493122	-1.0805578
site3	1.0176088	0.3263931	3.1177397
site4	0.5370828	0.3562308	1.5076822
site5	-0.2623643	0.4205764	-0.6238207
site6	0.5858730	0.3528776	1.6602725
site7	-0.4855078	0.4493133	-1.0805552
site8	0.1992985	0.3791789	0.5256054

```
(Dispersion Parameter for Poisson family taken to be 1 )
```

```
Null Deviance: 132.9485 on 15 degrees of freedom
```

```
Residual Deviance: 16.27524 on 7 degrees of freedom
```

```
Number of Fisher Scoring Iterations: 4
```

4 A client has come to two statisticians (Dr. Mean and Dr. Variance) with data collected from a one-academic year randomised-controlled study on m students, known for their tendency to get into fights in school. The study randomised students to receive, at the beginning of the academic year, either the new Counselling and Managing Behaviour (CAMB) therapy treatment or the standard Warning treatment (which is administered at the time of a fight) in order to determine whether the new treatment procedure was effective in reducing the number of fight episodes seen during the academic year.

The client has brought the fight-episode data in the form of counts $\mathbf{Y}_i = (Y_{i1}, Y_{i2}, Y_{i3})$, $1 \leq i \leq m$, recorded for each term in the academic year. Additional information on a student is recorded in covariate vectors \mathbf{x}_{ij} , $1 \leq i \leq m$, $1 \leq j \leq 3$, which includes information on what treatment was received.

Both Drs. Mean and Variance realise that there will be a correlation between the components of \mathbf{Y}_i . Dr. Mean decides to model the data as follows. He assumes that

$$\begin{aligned}\log E(Y_{ij} | \mathbf{x}_{ij}) &= \beta_0 + \beta^T \mathbf{x}_{ij} = \log \mu_{ij} \\ \text{Var}(Y_{ij} | \mathbf{x}_{ij}) &= \mu_{ij} \\ \text{Corr}(Y_{ij}, Y_{ik} | \mathbf{x}_{ij}, \mathbf{x}_{ik}) &= \rho (j \neq k).\end{aligned}$$

However, Dr. Variance decides to adopt the following alternative approach. She assumes that conditional on b_i , the responses Y_{ij} 's on the i th student are independent Poisson random variables with

$$\begin{aligned}E(Y_{ij} | \mathbf{x}_{ij}; b_i) &= \eta_{ij} \\ \text{Var}(Y_{ij} | \mathbf{x}_{ij}; b_i) &= \eta_{ij} \\ \text{Cov}(Y_{ij}, Y_{ik} | \mathbf{x}_{ij}, \mathbf{x}_{ik}; b_i) &= 0, (j \neq k) \\ \log \eta_{ij} &= b_i + \beta_0 + \beta^T \mathbf{x}_{ij}\end{aligned}$$

She also assumes that the $\exp(b_i)$'s are independent and identically distributed Gamma(τ^2/θ , τ/θ) (i.e. with mean τ and variance θ).

(i) What are the differences between the two approaches?

(ii) How would you interpret, for the client, the intercept parameter, β_0 , and the treatment parameters, say β_1 , from the two models? How would you interpret the parameter θ ?

(iii) Find $\log \mathbb{E}(Y_{ij} | x_{ij})$ for Dr. Variance's model and compare it with the expression given in Dr. Mean's model. If Dr. Variance's model was correct in this situation, would Dr. Mean be *consistently* estimating what *he thinks* he is estimating? Explain your answer.

(iv) If the variance and correlation structures in Dr. Mean's model were incorrectly specified, but the mean structure was correctly specified, how would Dr. Mean be able to make valid inferences about the parameters of interest?

5 (i) Suppose that y_1, \dots, y_n are independent Poisson random variables, and $\mathbb{E}(y_i) = \mu_i$, $1 \leq i \leq n$. We wish to fit the model ω , defined as

$$\omega : \log \mu_i = \mu + \beta^T x_i, \quad 1 \leq i \leq n,$$

where μ, β are unknown parameters and x_1, \dots, x_n are given covariates. Show that the deviance D for testing the fit of ω may be written as

$$D = 2 \sum y_i \log(y_i/e_i)$$

where (e_i) are the “expected values” under ω , and show that $D \simeq \sum (y_i - e_i)^2/e_i$.

(ii) Now suppose that y_1, \dots, y_n are independent negative binomial variables, and that y_i has frequency function

$$f(y_i | \theta, \mu_i) = \frac{\Gamma(\theta + y_i)}{\Gamma(\theta) y_i!} \frac{\mu_i^{y_i} \theta^\theta}{(\mu_i + \theta)^{\theta + y_i}}$$

for $y_i = 0, 1, 2, \dots$, thus $\mathbb{E}(y_i) = \mu_i$, $\text{var}(y_i) = \mu_i + \mu_i^2/\theta$.

Assume that θ is known. Show that the deviance for testing

$$\omega_n : \log \mu_i = \beta^T x_i \quad , \quad 1 \leq i \leq n$$

is say D_n , where

$$D_n = 2 \sum y_i \log \frac{y_i}{e_i} - 2 \sum (y_i + \theta) \log \frac{(y_i + \theta)}{(e_i + \theta)}$$

where (e_i) are the “expected values” under ω_n .