

STA 423

# OFFICE OF THE DEPUTY PRINCIPAL ACADEMICS, STUDENT AFFAIRS AND RESEARCH

# **UNIVERSITY EXAMINATIONS**

**2020 /2021 ACADEMIC YEAR** 

# FOURTH YEAR SECOND SEMESTER REGULAR EXAMINATION

FOR THE DEGREE OF BACHELOR OF SCIENCE (APPLIED STATISTICS WITH COMPUTING)

**COURSE CODE:** 

**STA 423** 

**COURSE TITLE:** 

**BIOMETRY METHODS** 

DATE: 22/7/2021

TIME: 1300-1600HRS

#### **INSTRUCTION TO CANDIDATES**

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# <u>REGULAR – MAIN EXAM</u> STA 423: BIOMETRY METHODS

**STREAM:** 

**DURATION: 3 hours** 

#### INSTRUCTION TO CANDIDATES

Answer ALL questions from section A and any THREE from section B.

#### **SECTION A [31 Marks] Answer All questions]**

#### **QUESTION ONE [16 Marks]**

a) Define clearly the following terms

[5 Marks]

- i) Clinical Trials
- ii) Blinding
- iii) Cross-sectional study
- iv) Longitudinal study
- v) Randomization
- b) Discuss any four types of endpoints in clinical trials

[8 Marks]

c) Identify three objectives of Phase II trials.

[3 Marks]

## **QUESTION TWO [15 Marks]**

a) Discuss any two advantages of a case-control study

[4 Marks]

b) Consider a placebo control trial investigating lung function in Asthmatic patients, the primary endpoint is the change from baseline in lung function at 16 weeks, at baseline we observe slight imbalances in characteristics such as age and baseline lung function, FEV. With the continuous and normally distributed endpoint of change from baseline in FEV, a simple linear regression analysis to assess treatment effect was conducted. Each patient, *i*, has a value for the outcome or dependent variable Y, FEV change while X is the explanatory variable is treatment which is binary being either active drug or placebo. The following table provides summary output of the simple linear regression model.

```
lm(formula = fev.change ~ group, data = asthma.trial)
Residuals:
                   Median
                  0.01849
-1.45445 -0.24530
Coefficients:
                          Std. Error t value Pr(>|t|)
                  0.01500
(Intercept)
                             0.03613
                                      -0.415
groupActive Drug
                  "***" 0.001 "**" 0.01 "*" 0.05 "." 0.1 " 1
Signif. codes:
Residual standard error: 0.3649 on 199 degrees of freedom
Multiple R-squared: 0.01935, Adjusted R-squared: 0.01442
F-statistic: 3.926 on 1 and 199 DF, p-value: 0.04891
```

i) Interpret the coefficients of the model

[3 marks]

ii) Make relevant conclusions

[2 marks]

c) For the same dataset in Question (b) multivariate linear regression was conducted to adjust for age effects. The regression model output is given below

```
lm(formula = fev.change ~ group + age, data = asthma.trial)
Residuals:
              10 Median
    Min
-1.48008 -0.24273 -0.00455 0.23823 0.84782
Coefficients:
                 Estimate Std. Error t value Pr(>|t|)
(Intercept)
                 0.590232
                           0.188586
                                     3.130 0.00201
                                      1.571 0.11773
groupActive Drug 0.079707
                            0.050729
                0.009250
                            0.002831
                                     -3.267
Signif. codes: 0 '*** 0.001 '** 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.3563 on 198 degrees of freedom
Multiple R-squared: 0.06951, Adjusted R-squared: 0.06011
F-statistic: 7.395 on 2 and 198 DF, p-value: 0.0007991
```

i) Interpret the coefficients of the model

[3 marks]

ii) Could there be any change from the conclusions made in Question (b)? Explain [3 marks]

## SECTION B [39 Marks] Answer any THREE questions]

# **QUESTION THREE [13 Marks]**

a) State some reasons a clinical trial may be stopped early.

[3 Marks]

b) Use the snapshot below of the structure of Acupuncture dataset from a randomized controlled trial investigating the effects of acupuncture on headache scores compared to a control intervention.

For the data, write the R codes to:

i) Display the numbers of patients randomized to the two treatment groups (Hint: use the **table()** function). [2 marks]

- ii) Generate summaries of the variables (score.baseline, age, and sex by treatment.group) by treatment group and save results as baselines (Hint: use the compareGroups() function) [4 marks]
- iii) Display the results saved in baselines (Hint: use the createTable() function) [2 marks]
- iv) Display the created summary table, that is, the numbers of patients randomized to the two treatment groups. [2 marks]

#### **QUESTION FOUR [13 Marks]**

- a) Identify the USA Institutional Review Board's (IRB) specific prerequisites that human research studies must meet. [6 marks]
- b) Suppose you want to conduct a clinical trial whereby simple randomization is used to design the study. You are required to produce a randomization list for the trial with 140 patients and two treatment arms, A and B.
  - i) Generate a vector to store treatment labels "A" and "B". Set the seed to a preferable number. [2 marks]
  - ii) Randomly select (with replacement) treatment arm 140 times with the **sample()** function and store in a vector [2 marks]
  - iii) Display the contents of the vector

iv) Tabulate the numbers assigned to each treatment [2 marks

#### **QUESTION FIVE [13 Marks]**

a) Giving examples, discuss any three categories of sources of bias in clinical studies [6 marks]

b) Let  $\overline{Y}_A$  be the average response for the sample of individuals assigned to treatment A and  $\overline{Y}_B$  the similar quantity for treatment B. Show that under stratified randomization,

$$\bar{Y}_A - \bar{Y}_B = \beta + \alpha \left( \frac{n_{A1}}{n_A} - \frac{n_{B1}}{n_B} \right) + (\bar{\epsilon}_A - \bar{\epsilon}_B).$$

[7 marks]

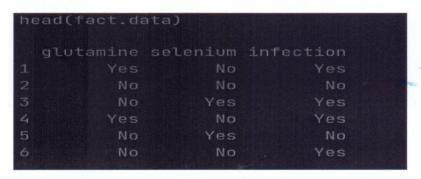
[1 mark]

# **QUESTION SIX [13 Marks]**

a) Identify any 5 purposes of a protocol document

[5 Marks]

b) A trial was designed to assess whether supplementing with glutamine or selenium, or both affected the odds of infections in critically ill patients. A factorial, randomized design was used. The data stored as **fact.data** has the head, as shown in the snapshot below;



#### Write R codes to:

i) Display the numbers with and without infections by supplement combination.

[2 marks]

- ii) Display the numbers and proportions with/without infections for those given glutamin e. [3 marks]
- iii) Display the numbers and proportions with/without infections for those given selenium
  [3 marks]

# **QUESTION SEVEN [13 Marks]**

- a) Identify three reasons for computing a correct sample size in clinical trials [3 marks]
- b) Give any four conditions for deriving the sample size necessary to detect a clinically important difference with some desired power [4 Marks]
- c) Suppose the standard treatment of care (treatment 2) has a response rate of about .35 (best guess). After collaborations with your clinical colleagues, it is determined that a clinically important difference for a new treatment is an increase in .10 in the response rate. That is, a response rate of .45 or larger. If we are to conduct a clinical trial where we will randomize patients with equal allocation to either the new treatment (treatment 1) or the standard treatment;
  - i) How large a sample size is necessary to detect a clinically important difference with 90% power using a one-sided test at the .025 level of significance? [5 marks]

ii) How many patients will be assigned to each treatment arm? [1 mark]

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