

UNIVERSITY OF SWAZILAND
FINAL EXAMINATION PAPER: MAY 2018

TITLE OF PAPER: GENOMICS

COURSE CODE: BIO 342

TIME ALLOWED: THREE HOURS

INSTRUCTIONS: 1. ANSWER SECTION A (COMPULSORY) AND ANY TWO OTHER QUESTIONS IN SECTION B.

2. QUESTION 1 CARRIES FIFTY (50) AND MARKS AND EACH QUESTION IN SECTION B CARRIES TWENTY FIVE (25) MARKS

3. ANSWER A TOTAL OF 3 (THREE) QUESTIONS

4. ILLUSTRATE YOUR ANSWERS WITH LARGE AND CLEARLY LABELLED DIAGRAMS WHERE APPROPRIATE

SPECIAL REQUIREMENTS: NONE

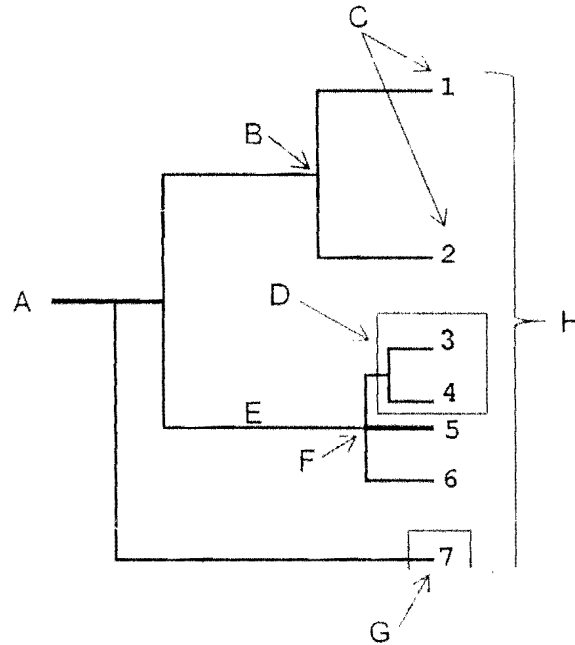
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Section A: Answer ALL questions in this section

Question 1 (Compulsory)

- (a) Define/explain the following terms:
- (i) CpG islands, (1 mark)
 - (ii) Synteny, (1 mark)
 - (iii) Reporter gene, (1 mark)
 - (iv) Expressed sequence tag, (1 mark)
 - (v) Paralogous genes, (1 mark)
 - (vi) Orthologous genes, (1 mark)
- (b) Shown below is a phylogenetic tree. Identify the labels A to H. (8 marks)



- (c) Describe the Sanger method for sequencing DNA. (6 marks)
- (d) Given that, starting from the sequencing primer, the following is the ssDNA sequence: G T A C C C G A A A T C A G G A. Sketch the expected electrophoretogram from this Sanger sequencing experiment. (5 marks)
- (e) Write brief notes on the following as they relate to functional genomics:
- (i) Site-directed mutagenesis, (5 marks)
 - (ii) RNAi, (5 marks)
 - (iii) Yeast two-hybrid system, (5 marks)
 - (iv) DNA microarrays, (5 marks)
 - (v) RNA sequencing. (5 marks)

[Total marks = 50]

Section B: Answer any TWO questions in this section

Question 2

Discuss the concept of genome annotation, highlighting the tools used and purpose thereof. (25 marks)

Question 3

Provide an overview of whole genome sequencing methods. (25 marks)

Question 4

Appraise the relevance of interactomics in biochemical and clinical research. (25 marks)

END OF EXAMINATION PAPER