

2021

M.Sc. (Bio-Informatics) Third Semester  
MBIN-8013: Programming Language in Bio-Informatics – II

Time allowed: 3 Hours

Max. Marks: 60

**NOTE:** Attempt five questions in all, including Question No. 1 which is compulsory and selecting atleast one question from each Unit.

x-x-x

I. Attempt the following:-

- a) Enlist various string operators used in Perl.
- b) Define Subroutines
- c) How Bio Perl is different from Perl?
- d) Empty XML Element
- e) MathML
- f) Well-Formed XML Document
- g) XML Prolog
- h) XML:: Simple Perl module

(8x1½)

**UNIT – I**

- II. a) What are control statements? Explain with examples.
- b) Discuss array functions available in PERL with example. (2x6)
- III. a) Write a PERL script to compute frequency of nucleotides from DNA Sequence.
- b) Explain the following briefly:
  - i) Reading data from FASTA files
  - ii) Regular Expression Operators (2x6)

**UNIT – II**

- IV. a) Create a Well-Formed XML instance document and write a DTD to validate it.
- b) Write the naming rules of Elements and Attributes in XML.
- c) Compare the features of HTML with XML (6+3+3)
- V. a) Explain Namespaces in XML using suitable example.
- b) What is DTD in XML? Give its types and significance. (2x6)

P.T.O.

(2)

**UNIT – III**

- VI. a) Write about major features of XML.  
b) Create an XML document to represent the following information and write a CSS script to display this XML document (use assumptions if any required).  
Accession: AAA1234, Type: mRNA, Organism: Homo sapiens  
Accession: BBB1234, Type: DMA, Organism: Pan troglodytes (2x6)
- VII. a) Write a script to demonstrate the use of Perl to process XML documents.  
b) Write short note on:-  
i) XML Tree  
ii) XML Parsing (2x6)

*x-x-x*