

6/H-77 (viii) (Syllabus-2015)

2 0 2 3

(May/June)

BIOTECHNOLOGY

(Honours)

**(Genomics, Proteomics and Computer
Application)**

Marks : 56

Time : 3 hours

*The figures in the margin indicate full marks
for the questions*

**Answer Question No. 1 which is compulsory
and any four from the rest**

- 1. Write notes on the following : 2×6=12**
- (a) BLAST**
 - (b) Quaternary structure of protein**
 - (c) C-value paradox**
 - (d) SNP**
 - (e) Protein domain**
 - (f) Transcriptomics**

2. (a) What is 'genome project'? Discuss the salient features of *Arabidopsis* genome project. 2+3=5
- (b) List the salient features of the human genome. What are the ethical issues involved in human genome sequencing project? 5+1=6
3. (a) Explain how STSs can be used to build a clone contig. 7
- (b) What is a flowchart? Explain the basic data structure of C++ with suitable example. 1+3=4
4. (a) Discuss the primary, secondary and tertiary structures of proteins. 7
- (b) Explain the role of bioinformatics in genomics with emphasis on comparative genomics. 4
5. Describe the role of computers in the automation of bioreactors. 11
6. (a) What is an operating system? What are the advantages of Windows operating system over Linux operating system? 2+4=6
- (b) Discuss in brief the concept of data mining and its significance in biological sciences. 5

7. (a) Discuss the role of bioinformatics in the field of proteomics. 5
- (b) What is an algorithm? Explain the batch-online and real time application of computers in industries. 1+5=6
8. What is bioinformatics? Enumerate the role of bioinformatics in various fields with suitable examples. Add a note on the limitations of bioinformatics. 1+8+2=11
