



DM-M-2011125

Seat No. _____

M. Pharm. (Sem. II) (CBCS) Examination

April / May – 2015

**Bioinformatics Theory
(Core-VI)**

Time : 3 Hours]

[Total Marks : 80

- Instructions :**
- (1) Answer and tie up both the sections separately.
 - (2) Figures to the right indicate marks.
 - (3) Answer any three (03) questions from each section
 - (4) Question one (1) and questions five (5) are compulsory
 - (5) Draw neat and clean diagrams as required

SECTION – I

- 1 Write notes on following : (any seven) 7×2=14**
- (A) Operating system
 - (B) e-Value
 - (C) Query coverage
 - (D) DBMS
 - (E) Bioinformatics
 - (F) NCBI
 - (G) FASTA
 - (H) DNASStar
 - (I) SMART
 - (J) Pfam

- 2 Write a detailed note on different types of BLAST. 13**

- 3 Write a detailed note on GCG Wisconsin package of sequence analysis program. 13
- 4 Answer the following questions :
- (A) Difference between Linux and Unix system. 7
- (B) What do you mean by databases? Write a short note on different databases. 6

SECTION – II

- 5 Answer any two out of three. 7×2=14
- (a) Write a note on Pubmed and e-Journals.
- (b) Write a short note on Protein-Ligand Docking.
- (c) Enlist various funding agencies for scientific fundings, and give a brief guideline about applying for a research grant.
- 6 Enlist different alignment algorithms and discuss any one in detail. 13
- 7 Discuss in detail about submission of DNA sequences to database. 13
- 8 Answer the following questions :
- (a) Describe different types of protein structures and write a note on secondary structure prediction. 7
- (b) Write a note on CLUSTAL W and CLUSTAL X for multiple sequence alignment. 6