

**Class : II PG**

**Time : 3 hours**

**Major : Bioinformatics**

**Max. marks : 60**

**17MBIC16 - Molecular Modeling, Simulation and Drug Designing**

**Part – A**

**10 x ½ = 5**

**Choose the correct answer**

1. Which of the following crosses the blood-brain barrier?  
a. GABA    b. Propranolol    c. Edrophonium    d. Dopamine
2. LD50 is:  
a. Median lethal dose  
b. Determined in phase I clinical trial  
c. Determined from log-dose response curve.  
d. Dose causing death in 50% of animals within 1 to 24 hours
3. An alignment that essentially spans the full extents of the input sequences is called a  
a. Global alignment    b. local alignment    c. multiple alignment    d. pairwise alignment
4. A bifurcating branch point in the phylogenetic tree is known as  
a. A Node    b. A clade    c. A branch    d. A taxon
5. Which of these methods is/are character-based methods in Tree construction  
a. Maximum parsimony    b. Maximum Likelihood  
c. Minimum evolution (ME) method    d. Neighbor joining (NJ)
6. PAM stands for  
a. Point Accepted Mutation    b. Percent Associated Mutation  
c. Protein Accepted Mutation    d. Point Associated Mutation
7. What step of DNA sequencing is skipped during shotgun sequencing?  
a. computer analysis    b. cloning of DNA fragment  
c. primer reactions    d. mapping step
8. The most effective way to analyze variation at the whole genome level is to use  
a. frequency of tandem clusters  
b. number of transposons.  
c. single nucleotide polymorphisms (SNPs).  
d. segmental duplications.
9. Labeling a stretch of DNA according to its function is called  
a. recombinant DNA technology    b. functional analysis.  
c. annotation    d. screening.
10. Sequences that share an arbitrary, threshold level of similarity determined by alignment of matching bases are termed as  
a. Homologous sequences    b. Heterologous sequences  
c. Mismatch sequences    d. Bogus sequences

**Part B**

**Answer all questions**

**Each answer should not exceed 200 words or one page**

**5x4 = 20**

11. (a) Derive the Schrödinger's equation and explain one electron atom.  
Or  
(b) Explain the Born-Oppenheimer approximation.
12. (a) Explain steepest descent minimization and conjugate gradient method  
Or  
(b) Explain in short different force field
13. (a) Explain newtonian dynamics  
Or  
(b) Elucidate implicit solvation model
14. (a) Explain maximum likelihood  
Or  
(b) What is De Novo drug designing?
15. (a) Elaborate on computer aided molecular design  
Or  
(b) What is regression analysis?

**Part B**

**Answer all questions**

**Each answer should not exceed 600 words or three pages**

**5x7 = 35**

16. (a) Explain in detail the four input formats for computation  
Or  
(b) Explain the semi empirical methods and potential energy calculation using *ab initio* methods
17. (a) Explain in detail about the molecular mechanics force field bonded interaction  
Or  
(b) Explain in detail about energy minimization and write brief notes on genetic algorithm
18. (a) What is simulation ? Explain various types of simulation.  
Or  
(b) What is Molecular dynamics? Elucidate the applications in molecular dynamics.
19. (a) What is drug discovery and its steps?  
Or  
(b) What is docking? Explain in detail about types of docking.
20. (a) Explain the significance and validity of QSAR.  
Or  
(b) Explain partial least squares analysis and multi linear regression analysis.