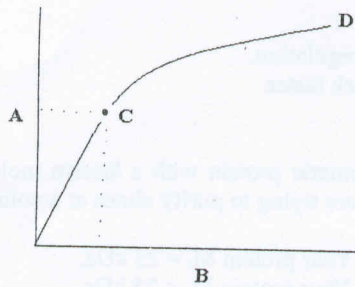


I. Multiple Choice (2% each, 20 questions, total 40%)

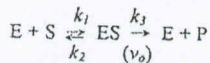
1. Buffers are substances that help resist shifts in pH by
  - A) releasing  $H^+$  in acidic solutions.
  - B) releasing  $H^+$  in basic solutions.
  - C) combining with  $H^+$  in basic solutions.
  - D) combining with  $OH^-$  in acidic solutions.
  - E) releasing  $OH^-$  in basic solutions.
2. What maintains the secondary structure of a protein?
  - A) peptide bonds
  - B) hydrogen bonds
  - C) disulfide bridges
  - D) ionic bonds
  - E) electrostatic charges
3. Alanine is similar to serine in the same way that
  - A) Val is similar to Thr.
  - B) Phe is similar to Trp.
  - C) Phe is similar to Tyr.
  - D) Ser is similar to Thr.
4. An  $\alpha$ -helix with 11 amino acid residues completes three turns. The length of this helix is about
  - A) 10 Å
  - B) 15 Å
  - C) 20 Å
  - D) 25 Å
5. Subunits of a single enzyme undergo conformational changes individually. Which model of allosteric regulation supports this statement?
  - A) The MWC Concerted Model
  - B) The Cooperative Binding Model
  - C) The sequential Model
  - D) The Transition State Model

The Michaelis-Menten equation is  $v_o = V_{max} [S] / (K_m + [S])$ .

Fill in the blanks (questions 6, 7) with the letters shown to correctly label each part of the graph.



6. \_\_\_\_\_  $v_o$
7. \_\_\_\_\_ Point used to determine the  $K_m$
8. The turnover number of an enzyme is the rate at which product is formed.



$$\text{turnover number} = k_3 = V_{max} / [E_T]$$

This is the maximum rate of product formation if all of the enzyme substrate binding sites are filled. What can the turnover number tell us about an enzyme?

- A) Number of binding sites
- B) Reaction mechanism
- C) Substrate concentration
- D) Enzyme efficiency

9. Bends or loops are considered to be non-regular secondary structures of proteins because they do not repeat. Which of the following is **true** of bends?
- A) Bends do not reverse the direction of a polypeptide chain.  
B) Bends connect regions of  $\alpha$ -helices and  $\beta$ -sheets.  
C) Glycine and proline are absent from bends.      D) Bends are usually found at the C terminus.
10. Collagen is best described as
- A) an  $\alpha$ -helical structural protein.      B) a coiled-coil found in hair.  
C) a cross-linked globular protein.      D) a triple-helical fibrous protein.
11. The Hill coefficient ( $n_H$ ) for myoglobin is \_\_\_\_; whereas  $n_H$  is about \_\_\_\_ for hemoglobin.
- A) 2.8; 1.0      B) 1.0; 2.8      C) 1.0; 4.0  
D) 4.0; 1.0      E) None of the above choices are correct.
12. The relationship between an enzyme and a reactant molecule can best be described as:
- A) a temporary association.      B) an association stabilized by a covalent bond.  
C) one in which the enzyme is changed permanently.      D) a permanent mutual alteration of structure.  
E) noncomplementary binding.
13. The  $K_m$  is:
- A) The time for half of the substrate to be converted to product.  
B) The time for all of the substrate to be converted to product.  
C) The [S] that gives half of the maximum reaction rate.  
D) The [S] that gives the maximum reaction rate.  
E) The [P] that is produced when the enzyme is saturated with the substrate.
14. An allosteric inhibitor of an enzyme usually
- A) binds to the active site.      B) participates in feedback regulation.  
C) denatures the enzyme.      D) causes the enzyme to work faster.  
E) is a hydrophobic compound.
15. The void volume of a gel filtration column,  $V_0 = 30$  mL. A monomeric protein with a known molecular weight of 25 kDa elutes at a volume of 45 mL. The protein that you are trying to purify elutes at a volume of 35 mL. Which of the following is a valid conclusion?
- A) Your protein is repelled by the gel filtration material.      B) Your protein  $M_r = 25$  kDa.  
C) Your protein  $M_r > 25$  kDa.      D) Your protein  $M_r < 25$  kDa.
16. A noncompetitive inhibitor of an enzyme-catalyzed reaction
- A) increases  $K_M$  and increases  $V_{max}$ .      B) increases  $K_M$  and reduces  $V_{max}$ .  
C) reduces  $K_M$  and increases  $V_{max}$ .      D) reduces  $K_M$  and reduces  $V_{max}$ .
17. The strong conclusion from Anfinsen's work on RNaseA was that:
- A) 100% enzyme activity corresponds to the native conformation.  
B) urea can reversibly denature proteins *in vivo*.      C) mercaptoethanol can reduce disulfide bonds *in vitro*.  
D) the native conformation of a protein is adopted spontaneously.
18. Catalytic antibodies act like enzymes in that they:
- A) Obey Michaelis-Menten kinetics      B) Bind to substrates in the transition state  
C) Bind to antigen      D) Need cofactors to function

19. The peptide, Val-Lys-Glu-Met-Ser-Trp-Arg-Ala, was digested with chymotrypsin to produce:
- A) Val-Lys + Glu-Met-Ser + Trp-Arg-Ala.
  - B) Val-Lys-Glu-Met-Ser-Trp + Arg-Ala.
  - C) Val-Lys-Glu-Met-Ser + Trp-Arg-Ala.
  - D) Val-Lys-Glu + Met-Ser-Trp-Arg-Ala.
  - E) Val-Lys-Glu-Met + Ser-Trp-Arg-Ala.
20. The molecular formula for glycine is  $C_2H_5O_2N$ . What would be the molecular formula for a linear oligomer made by linking ten glycine molecules together by condensation synthesis?
- A)  $C_{20}H_{50}O_{20}N_{10}$ .
  - B)  $C_{20}H_{32}O_{11}N_{10}$ .
  - C)  $C_{20}H_{40}O_{10}N_{10}$ .
  - D)  $C_{20}H_{68}O_{29}N_{10}$ .
  - E) None of the above.