Principles of Medical Biochemistry 3rd Edition Meisenberg Test Bank

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Meisenberg: Principles of Medical Biochemistry, 3rd Edition

Chapter 2: Introduction to Protein Structure

Test Bank

MULTIPLE CHOICE

- 1. Which of the following is true about the tertiary structure of proteins?
- a. Disulfide bonds are part of the tertiary structure.
- b. Only proteins with more than one polypeptide subunit have a tertiary structure.
- c. Proteins with tertiary structure do not contain α helix or β -pleated sheet.
- d. Van der Waals interactions play no role in the tertiary structure.
- e. Interactions between hydrophobic amino acid side chains are important for hold the tertiary structure together.

ANS: E

Hydrophobic groups associate with one another to minimize the thermodynamically unfavorable interface between lipid and water. Hydrophobic interactions and van der Waals interactions prevail in the core of globular proteins.

- 2. Amino acids at the isoelectric point in the titration curve have a net charge of:
- a. 1.
- b. +1.
- c. +2.
- d. 0.
- e. –2.

ANS: D

This is the definition of the isoelectric point.

- 3. Which amino acid residues are used as attachment sites for covalently bound oligosaccharides in glycoproteins?
- a. Asparagine and serine.
- b. Tryptophan and glutamic acid.
- c. Alanine and lysine.
- d. Arginine and proline.
- e. Leucine and histidine.

ANS: A

N-linked carbohydrate is bound to asparagine, and *O*-linked carbohydrate is bound to serine or threonine (or, in collagen, to hydroxylysine).

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4. Which of the following is true regarding the structure shown?

-CH2-CH2-COO

- a. Side chain of glutamine; normally found in the interior of globular proteins.
- b. Side chain of glutamine; normally found on the surface of globular proteins.
- c. Side chain of glutamate; normally found in the interior of globular proteins.
- d. Side chain of glutamate; normally found on the surface of globular proteins.
- e. Side chain of the "nonstandard" amino acid γ -carboxyglutamate, found only in some clotting factors.

ANS: D

Unlike glutamine, glutamate is negatively charged. All negatively charged amino acid side chains prefer the surface of globular proteins, where they can interact with water and dissolved ions.

- 5. In the tripeptide glutathione (γ-glutamyl-cysteinyl-glycine), the side chain carboxyl group of glutamate forms a peptide bond with the α-amino group of cysteine. The α-amino and α-carboxyl groups of glutamate do not participate in peptide bonds. Approximately what is the isoelectric point (pK) of glutathione?
- a. 1.5.
- b. 10.0.
- c. 3.0.
- d. 6.5.
- e. 9.0.

ANS: C

There is an α-amino group (of glutamate) with a pK near 9 or 10, an α-carboxyl group (of glutamate) with a pK near 2, a carboxyl terminus (formed by glycine) with a pK near 4, and a cysteine sulfhydryl group (SH) with a pK near 8. The isoelectric point is halfway between the pK values of the two carboxyl groups.

- 6. Noncovalent bonds essential for the formation of the α helix and β -pleated sheet are:
- a. Disulfide bonds.
- b. Van der Waals interactions.
- c. Salt bridges.
- d. Hydrogen bonds.
- e. Hydrophobic forces.

ANS: D

The hydrogen bonds are formed between the components of the peptide bonds.

- 7. Histones are proteins that bind to negatively charged phosphate groups of DNA. An amino acid in the histones that can mediate this binding is:
- a. Valine.

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- b. Lysine.
- c. Aspartate.
- d. Cysteine.
- e. Glutamic acid.

ANS: B

The lysine side chain carries a positive charge at pH values near 7.

- 8. The secondary structure of proteins:
- a. Is maintained by hydrogen bonds.
- b. Is present only in proteins consisting of two or more subunits held together by noncovalent forces.
- c. Refers to any hydrogen-bonded interaction found in proteins.
- d. Implies the presence of a nonprotein moiety bound to the polypeptide.
- e. Is found only in fibrous proteins.

ANS: A

The hydrogen bonds are formed between the components of the peptide bonds.

- 9. Which of the following statements about protein structure is correct?
- a. The \Box helix is stabilized primarily by ionic interactions between the side chains of amino acids.
- b. Cytoplasmic proteins generally contain disulfide bonds.
- c. In comparison with the β -pleated sheet, the α helix is more extended.
- d. The denaturation of proteins is in most cases reversible by slow cooling.
- e. The tertiary structure of the protein forms before the formation of disulfide bonds.

ANS: E

Protein conformation is established by noncovalent interactions between functional groups of the polypeptide, sometimes aided by helper proteins called *chaperones*. Disulfide bonds are formed between cysteine side chains that have been brought into close proximity during the initial folding process.

- 10. An amino acid whose side chain is most likely to be found in the center of a tightly packed, water-soluble globular protein such as myoglobin is:
- a. Serine.
- b. Glutamine.
- c. Aspartate.
- d. Leucine.
- e. Arginine.

ANS: D

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Hydrophobic amino acid side chains aggregate in the core of globular proteins, to avoid contact with the surrounding water.

Better start: The leucine side chain is hydrophobic. Hydrophobic ...

- 11. Which of the following statements is correct for protein structure?
- a. The α helix is stabilized primarily by ionic interactions between the side chains of amino acids.
- b. Disulfide bonds are common in cytoplasmic proteins but are not present in most proteins of the extracellular matrix.
- c. The stability of the quaternary structure in proteins is a result of covalent bonds between the subunits.
- d. The heat denaturation of proteins is in most cases reversible by slow cooling.
- e. Glycine and proline do not usually participate in α -helical structures.

ANS: E

Glycine is too flexible, and proline is too rigid to fit comfortably in an α helix.

- 12. All naturally occurring amino acids:
- a. Have more than one α -carbon atom.
- b. Are uncharged at a pH of 7.
- c. Can only have two pKs.
- d. Are chiral, except glycine.
- e. Occur in nature mainly or exclusively in the optically active D-form.

ANS: D

Only glycine has no asymmetrical carbon.