Chemistry 333
Principles of Biochemistry
Fall 2004
First Exam
September 21, 2009

NAME:

To be eligible for a regrade, the exam must be done in ink.

| l | /5 points |
|--------|--------------------------|
| 2 | /12 points |
| 3 | /20 points |
| 4 | /4 points |
| 5 | /5 points |
| 6 | /4 points |
| 7 | /24 points |
| 8 | /26 |
| 9 | /2 points (EXTRA CREDIT) |
| | |
| | |
| TOTAL: | /100 points |

| | HA AND THE PARTY OF THE PARTY O | Ka | pK _a |
|-----------------|--|----------------------------|-----------------|
| Acid | | 1.78×10^{-4} | 3.75 |
| Formic acid | HCOOH | 1.76×10^{-5} | 4.75 |
| Acetic acid | CH3COOH | 3.16×10^{-3} | 2.50 |
| Pyruvic acid | CH₃COCOOH CH₃CHOHCOOH | 1.38×10^{-4} | 3.85 |
| Lactic acid | HOOC—CH ₂ —CHOH—COOH | (1) 3.98×10^{-4} | 3.40 |
| Malic acid | OH OH | (2) 5.5×10^{-6} | 5.26 |
| Citric acid | H00C-CH2-C-CH2-C00H | (1) 8.14×10^{-4} | 3.09 |
| | Соон | | |
| | | (2) 1.78×10^{-5} | 4.75 |
| | | $(3) 3.9 \times 10^{-6}$ | 5.41 |
| | н со | (1) 4.3×10^{-7} | 6.4 |
| Carbonic acid | H ₂ CO ₃ | $(2) 5.6 \times 10^{-11}$ | 10.2 |
| n | U PO | (1) 7.25×10^{-3} | 2.14 |
| Phosphoric acid | H ₃ PO ₄ | (2) 6.31×10^{-8} | 7.20 |
| | | (3) 3.98×10^{-13} | 12.4 |
| Ammonium ion | NH ⁺ | 5.6×10^{-10} | 9.25 |

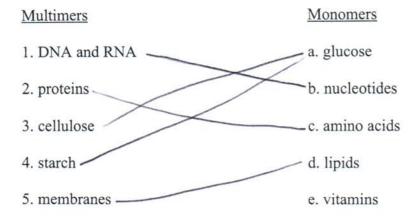
TABLE 4.2

Table of pK values for the 20 amino acids found in proteins a

| Name | p.K ₁ | pK ₂ | pK _B |
|---------------|------------------|-----------------|-----------------|
| Glycine | 2.4 | 9.8 | |
| Alanine | 2.3 | 9.9 | |
| Valine | 2.3 | 9.6 | |
| Leucine | 2.4 | 9.6 | |
| Isoleucine | 2.4 | 9.7 | |
| Methionine | 2.3 | 9.2 | |
| Phenylalanine | 1.8 | 9.1 | |
| Proline | 2.0 | 10.6 | |
| Serine | 2.1 | 9.2 | |
| Threonine | 2.6 | 10.4 | |
| Cysteine | 1.8 | 10.8 | 8.3 |
| Asparagine | 2.0 | 8.8 | |
| Glutamine | 2.2 | 9.1 | |
| Tyrosine | 2.2 | 9.1 | 10.5 |
| Tryptophan | 2.4 | 9.4 | |
| Aspartate | 2.0 | 10.0 | 3.9 |
| Glutamate | 2.2 | 9.7 | 4.3 |
| Histidine | 1.8 | 9.2 | 6.0 |
| Lysine | 2.2 | 9.2 | 10.8 |
| Arginine | 1.8 | 9.0 | 12.5 |

 $^{^{0}}$ p K_{1} values are assigned to the α -carboxyl group, p K_{2} values to the α -amino group, and p K_{R} to ionizable groups in the R group (side chain).

Biological macromolecules are multimers made up of monomeric building blocks.
Use arrows to match the monomers used to synthesize these naturally occurring
polymers. Each monomer may be used more than once or not at all. (5 points)



2. How many **grams** each of solid KH₂PO₄ (MW = 136 g/mol) and solid K₂HPO₄ (MW = 174 g/mol) would you use to prepare 2 L of a 0.3 M phosphate buffer (phosphoric acid) at pH 6.8? (12 points)

Write out and identify the ACID and the CONJUGATE BASE. (Ignore the K⁺ except when calculating the grams needed.)

Partial credit will be awarded, so please show all of your work including any equations!

KH2PO4 - ACID

KZHPO4 - Conjugate Base

$$CA^{-1}$$
 CA^{-1}
 CA

0.714 x 0.6 moles = 0.4284 moles HA

- Answer the following questions regarding the peptide LCYRAIDCG (20 points) 3.
 - What is the sequence of amino acids written as the THREE letter code? A.

Draw the structure of the peptide LCYRAIDCG as it would exist at pH 6.5 under B.

oxidizing conditions. Be sure to draw any disulfide bonds that could form.

- C. Label the N-terminus of the peptide you drew in 3B above with a ✓.
- Label the C-terminus of the peptide you drew in 3B above with a ★. D.
- Calculate the net charge of LCYRAIDG at pH 1.5. E.

Net Charge = +2

Calculate the approximate pI of the peptide DSYRLKCF (4 points) 4.

Draw the form(s) of the amino acid lysine that would exist at pH 10.8.
 In what relative percentages (% of each) will these forms exits? Why? (5 points)

6 A &B (4 points)

There are six forms of a protein; one is normal and five contain different mutations that change the amino acid at position 150 (called mutants). The normal enzyme has a glutamine residue at amino acid position 150 that is located on the protein surface. Each mutant form of the protein has an amino acid substitution at position 150 as indicated:

- 6A. Which mutant form of the protein is most like the normal form because it contains the most **conservative** substitution? (i.e. which side chain most resembles that of glutamine?)
 - a. Form A: glutamine has been replaced by alanine
 - b. Form B: glutamine has been replaced by arginine Form C: glutamine has been replaced by asparagine
 - d. Form D: glutamine has been replaced by glutamate
 - a. Form E: glutamine has been replaced by phenylalanine
- 6B. Which mutant form has the highest propensity to place amino acid 150 in the **interior** of the protein?
 - a. Form A: glutamine has been replaced by alanine
 - b. Form B: glutamine has been replaced by arginine
 - c. Form C: glutamine has been replaced by asparagine
 - d. Form D: glutamine has been replaced by glutamate
 - (e.) Form E: glutamine has been replaced by phenylalanine

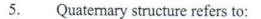
| 7. | Match the characteristics at left with the best matching amino acids at right by entering each characteristic's number in the appropriate blank. Unless otherwise stated, all of the |
|----|--|
| | amino acids are at physiological pH 7.4. There are more than one characteristic for many |
| | of the amino acids and 24 total answers each worth one point. There will be a deduction |
| | for incorrect answers, so don't fill in the lines with unnecessary characteristics. (24 points) |

| 1. | acidic side chain | | leucine |
|-----|---|---------|---------------|
| 2. | non-chiral amino acid | 1,10 | aspartate |
| 3. | basic side chain | 4,12,13 | serine |
| 4. | side chain can be modified by adding phosphates | 3,8,9 | lysine |
| 5. | involved in disulfide cross links | 7,14 | phenylalanine |
| 6. | often found in the turns of proteins | 9,12,13 | asparagine |
| 7. | is converted to tyrosine by hydroxylation | 2,6 | glycine |
| 8. | important for the structure of collagen | 4,12,14 | tyrosine |
| 9. | has more than one amino group | 6,8 | proline |
| 10. | has an overall net charge of -1 at pH 8.5 | 5,10,12 | cysteine |

- 11. only straight or branched hydrocarbons in the side chain
- 12. polar, non-charged side chain
- 13. side chain can be modified by adding sugars
- 14. aromatic side chain

8. Multiple Choice: 2 points each (26 points)

- 1. The **primary** purpose of membranes in eukaryotic cells is:
 - A. to increase surface area inside the cell
 - (B) to create compartments to separate biochemical functions
 - C. to provide anchors for the cytoskeleton
 - D. to provide attachment sites for ribosomes
 - E. none of the above
- 2. What are the energy-producing organelles in eukaryotic cells?
 - A. endoplasmic reticulum
 - B. lysosomes
 - C. nucleus
 - mitochondria
 - E. golgi
- 3. Aspartame (Nutrasweet) is
 - A. deadly in small amounts
 - a dipeptide
 - C. a modified sugar
 - D. a carbohydrate
 - E. an amino acid
- 4. Free rotation about the peptide bond in a protein is restricted primarily because of
 - A partial double-bond character of the peptide bond.
 - B. hydrogen bonding to the amide backbone groups.
 - C. partial double-bond character of the N-C alpha bond.
 - D. restrictions caused by local folding patterns.
 - E. steric interference of neighboring amino acid side chains.



- A. the overall shape of the polypeptide chain
- B. the sum of secondary and tertiary interactions
- C. simple proteins with one subunit
- the relative orientation of one polypeptide to another polypeptide in a multisubunit protein complex
- E. the linear sequence of amino acids in a protein

6. Enantiomers are:

- A. compounds with very different chemical composition
- B. molecules with both hydrophobic and hydrophilic characteristics
- (C) a pair of compounds that are non-superimposable mirror images of one another
- D. not chiral
- E. superimposable mirror images

7. The lysosome is important for:

- A. Biosynthesis of amino acids
- B Disposal of molecular waste/excess
- C. Protein translation
- D. Glucose breakdown
- E. None of the above
- The primary structure of a protein describes the 8. number of each type of amino acid (percent composition) A.

 - B linear sequence of amino acids
 - C. overall three-dimensional shape
 - D. rotation angles for each amino acid
 - E. assembly of protein subunits into complexes

| 9. | S | tructural proteins that typically assemble into large cables or threads to provide aechanical support to cells or organisms are classified as proteins. |
|-----|---------------------|---|
| | | fibrousproteins. |
| | В | . enzyme |
| | C | . globular |
| | D | . receptor |
| | E. | α-helical |
| | | |
| 10. | Th | e [H ₊] concentration at pH 4.4 is: |
| | A. | 10 times higher than the [H+] concentration at pH 6.4 |
| | B. | 10 times lower than the [H+] concentration at pH 5.4 |
| | C. | 100 times higher than the [H+] concentration at pH 3.4 |
| | 0 | 100 times higher than the [H+] concentration at pH 6.4 |
| | E. | 100 times lower than the [H+] concentration at pH 5.4 |
| | | |
| | | |
| 11. | Ifl | numan blood is not maintained at close to pH = 7.4, a person can develop |
| | A. | Acidosis |
| | B. | Alkalosis |
| | C. | Diabetes |
| | 0 | Both a and b |
| | E. | None of the above |
| | | |
| 12. | P ₁ w | roteins are biopolymers of amino acids in which monomers are linked in a linear fashion ith which type of bond? |
| | (| amide |
| | B. | ester |
| | C. | ether |
| | D. | phosphoanhydride |
| | E. | phosphate ester |
| | | |
| | | |

- Which of the following compounds would you select to construct a buffer at pH 8? 13.
 - A. acetic acid, pKa = 4.8
 - N-tris(hydroxymethyl)methylglycine (Tricine), pKa = 8.15 (B)
 - N-2-hydroxyethylpiperazine-N'-ethansulfonic acid, pKa2 = 7.6C.
 - D. glycine, pKa2 = 9.9
 - E. tris(hydroxymethyl)aminomethane (Tris), pKa = 8.3

9. EXTRA CREDIT: (2 points)

If 70 mL of 0.01 M HCl is added to 150 mL of 0.1M acetic acid, pH 5.0, what is the 1. resulting pH? Identify the appropriate acid and conjugate base and determine their concentrations in the final solution. (2 points)

CH₃ COOH
$$\Rightarrow$$
 CH₃COO+HD

 $PH = PKQ + log \frac{[A-]}{[CHA]}$
 $70mL = 0.07L$
 $5.0 = 4.75 + log \frac{[A-]}{[CHA]}$
 $0.07L \times \frac{0.01 \, mol}{L} = 1.7 \, mmol$
 $CA-] = \frac{1.8}{[CHA]}$
 $1.8 + l = 2.8$

added