YOUR NAME:		

# NOTES:

- 1. where appropriate please show work if in doubt show it anyway.
- 2. pace yourself you may want to do the easier questions first.
- please note the point value of questions adjust your answers and effort accordingly.
- 4. some questions may have more data than you need.
- 5. please be brief unfocused, rambling answers won't receive as much credit as a few short appropriate phrases.
- 6. Please write CLEARLY if I cannot read it it is wrong.
- 7. Please refer, as needed, to the chart below.

Properties of the amino acids found in proteins

Name	$pK_a$ of $\alpha$ -Carboxyl Group	$pK_a$ of $\alpha$ -Amino Group	pK <sub>a</sub> of Ionizing Side Chain <sup>a</sup>
Alanine	2.3	9.7	
Arginine	2.2	<b>9.0</b> .	12.5
Asparagine	2.0	8.8	
Aspartic acid	2.1	9.8	3.9
Cysteine	1.8	10.8	8.3
Glutamine	2.2	9.1	-
Glutamic acid	2.2	9.7	4.2
Glycine	2.3	9.6	
Histidine	1.8	9.2	6.0
Isoleucine	2.4	9.7	<del></del>
Leucine	2.4	9.6	<u> </u>
Lysine	2.2	9.0	10.0
Methionine	2.3	9.2	
Phenylalanine	1.8	9.1	
Proline	2.0	10.6	
Serine	2.2	9.2	<del></del>
Threonine	2.6	10.4	<del></del>
Tryptophan	2.4	9.4	
Tyrosine	2.2	9.1	10.1
Valine	2.3	9.6	

# Question 1. (21 pts.) Short problems. Show work, but most credit goes to the correct numerical answer.

a. the pH of a solution is 9.8. What is the hydroxide concentration?

 $[OH^{-}] =$ \_\_\_\_\_M

**b.** You mix 500 mL of 50 mM NaOH with 800 mL of 60 mM HCl. What is the pH of the mixture?

pH = \_\_\_\_\_

**c.** You add 0.08 mol of acetic acid (pK = 4.70) in 10 mL of water to 0.03 mol of lithium acetate in 100 mL of water. What is the new pH?

pH = \_\_\_\_\_

d. A solution of 0.3 moles of a weak acid in 1 L of water showed a pH of 4.5. What is the pK of the acid.

pK =\_\_\_\_

**e.** If you add 0.15 moles of sodium hydroxide to a solution of 0.2 moles of formic acid (pK=3.7) in 0.5 L what is the resulting pH?

pH =\_\_\_\_

<ul> <li>A human phospholipas many possible combina</li> </ul>	e contains 8 disulfide bonds a tions of disulfide-bridged pho	and no free cysteine ospholipase molecul	e side chains. How les exist?
		Number =	
g. An oxygen binding prot of 10 μΜ. What is the	ein is half-saturated ( $\theta$ = 0.5)	at a dissolved oxyg	en concentration
or to pivi. What is the	Na for oxygen binding:	K <sub>d</sub> =	μМ
	CLEAR structures of the fo ndicated (refer to the list o		ls in the form that
CYSTEINE at pH 5	ALANINE at pH 12	ASPA	RTIC ACID, pH 1
	orm of tyrosine is shown a	t the right.	ОН
a. You have 0.6 moles o	2, 9.1, 10.1)  f tyrosine in the form show oles do you need to take th	e original 0.6	
9.1	10.1	+	ĊH₂ H₃N—-ĊH—Ç—-Œ
b. What is the pH of the tyrosine in 0.5 L of wa	solution of 0.6 moles of th	is form of	0
		pH =	
c. Is there a pH value at	which tyrosine would pene	trate biological me	embranes?
Circle either "YES"	': at pH =	or "NC	)", at no pH
Explain your answer:			

Question 4 (6 pts.) A peptide composed of 30 lysine residues (LYS-LYS-LYS- ... etc) is an alpha helix at pH 13 but a random, disordered, structure at pH 1. Explain why this could be.

What peptide would show exactly the opposite behavior. It is ...

Explain why:

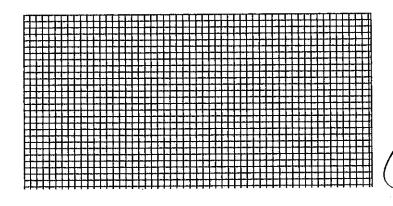
Question 5 (6 pts). Here is the structure of nicotine – an active ingredient in tobacco. What form of nicotine will predominate at the pH of tobacco leaves (about 6). If it is the same as the figure, circle it, if it is different, change structure as necessary.

Tobacco companies treat tobacco leaves with  $NH_3$  - they say "to improve the taste" – prior to packaging into cigarettes. Give a more reasonable answer.

Question 6 (12 pts) Draw the tripeptide LYS-GLY-ALA in the form appropriate for pH 7.

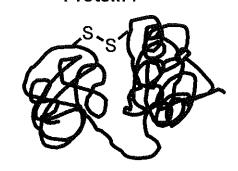
At the right, draw an ACCURATE titration curve of this tripeptide starting with the most protonated form you could obtain at the left. Include number of equivalents and pKs etc. Where necessary, use pK values on page 1

рΗ



### **Protein P**

Question 7 (25 pts) The protein "P" (40,000 MW; or 40 kDa) shown to the right is a single polypeptide chain. The protein is a monomer and contains: 4 cysteine residues (2 of them in a disulfide bond as shown) and 7 ARG and 3 LYS (including a LYS at the C terminus). Edman degradation of "P" gave the following initial sequence:



#### MIKEY-

a. What is the three letter abbreviations for this

b. P is dissolved in buffer alone and treated with trypsin. Draw an accurate representation of the

chemical reaction that takes place with trypsin treatment. Include all relevant species.

c. Only one cleavage site occurs in 'b'. How many might you expect? # =

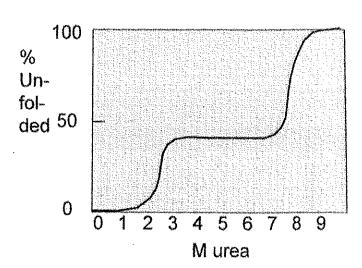
## and why only one?

- **d.** the trypsin treated "P" was purified and subjected to very high resolution mass spectrometry. Compared to P, <u>treated</u> "P" had gained in molecular weight slightly. Explain **how is this possible**.
- f. treated 'P' (as in step b) is then exposed to 2-mercaptoethanol (sometimes called β-ME). Draw the structure of 2-ME.

Then, two fragments of treated 'P' were obtained (of approximately 18 and 22 kDa). Why?

g. Samples of the original 'P' were dissolved in increasing concentrations of urea. Draw the structure of urea:

The degree of unfolding of the protein was then determined and gave the result shown in the diagram. Give a short explanation of the behavior of the curve in light of lecture material.



Question 8 (6 pts.) Proteins A-E show the following pl values:

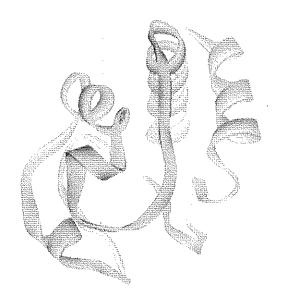
During ion exchange chromatography at pH 7.50

- i. which protein will elute fastest on anion exchange
- ii. which protein will elute slowest on anion exchange
- iii. which protein will elute fastest on cation exchange
- iv. which protein will elute slowest on cation exchange

Protein C elutes much faster than protein D on anion exchange at pH 7.50. Suggest a simple reason for this:

# Question 9 (7 pts.) Fill in the blanks with not more than 3 legible words.

a.	a technique for determining 3D structure of proteins
b.	a method for visualizing complex protein mixtures in proteomics
c.	oxygen is always bound <b>directly</b> to in all oxygen-binding proteins (worms to humans)
d.	these catalysts aid folding of large proteins
e.	every third amino acid in collagen is almost always
f.	In one collagen disease, the amino acid in e is replaced by a
77	the word that heet describes this evam



the end