Name

____Section_

7.012 Problem Set 1 Solutions

TA

Question 1

Α.

a) Explain what an ion is.

An ion is an electrically charged particle that forms when an atom gains or loses one or more electrons.

b) Explain what a van der Waals force is.

A weak attraction between molecules resulting from a momentary shift of the electron density in a molecule thus establishing a dipole moment. The dipoles induced by the effect of one molecule on the others lead to a net electrostatic attraction.

c) Match the atoms and molecules listed below with the appropriate descriptive. Choose from:

cation	anion	pol	ar non-charged	nonpolar non-charged
		Cl ⁻	Anion	
		H₂O _	Polar non-charged	,
		Na⁺	Cation	
		CH₄	Nonpolar non-charge	ed

d) Circle the "side chain" in each of the amino acids given below.



e) Check the following terms to describe the side chains of the amino acids you circled above.

Side Chain of	Charged	Non Charged	Polar	Non Polar	Hydrophobic	Hydrophilic
Glutamic Acid	ſ		\checkmark			5
Lysine	ſ		ſ			Г
Tyrosine		ſ		√-/+	√-/+	√-/+
Asparagine		ſ	ſ			Г
Glycine		5		J	ſ	
Glutamine		ſ	ſ			ſ

f) Explain why ethane (C_2H_6) is a non-polar molecule while ethanol (C_2H_5 - OH) is a polar molecule. (A diagram of each molecule may help.)

Ethanol is a polar molecule because there is a partial charge separation associated with the bond between the hydrogen and the oxygen. The higher electronegativity of oxygen pulls the electrons in the bond closer to the oxygen resulting in a greater electron density around the oxygen. This provides a slight negative charge for the oxygen and a slight positive charge for the carbon. The molecule is polar because these opposite charges are separated at the two ends of the bond. Ethane, on the other hand, is nonpolar because there are no such charge separations present in this molecule. The electronegativity difference between carbon and hydrogen are not significant enough to cause this type of charge separation.

Β.

Match the letter of the organelle on the left with the function or characteristic on the right.

Complex or Organelle		Function or Characteristic
a. Mitochondria	<u> </u>	Site of lipid synthesis; where hydrophobic toxins→ water soluable and thus excretable
b. Nucleus	<u> </u>	Digests macromolecules and degrades worn-out organelles
c. Smooth Endoplasmic Reticulum	<u>F</u>	Catalyzes protein synthesis
d. Golgi Apparatus	<u> </u>	Contains all genomic DNA in eukaryotic cell
e. Lysosome	<u> </u>	Uses sunlight to build sugar
f. Ribosome	<u> </u>	Where ribosomal RNA is made
g. Chloroplast	D	Prepares proteins for export from the cell
h. Nucleolus	<u> </u>	Powerhouse of the cell

Question 2

One day in lab while studying your favorite enzyme, Votase, you discover the following potential interactions that could occur between this amazing enzyme and its substrate of choice.



a) At each site between the chemical group on the substrate and the closest side chain of an amino acid on Votase determine if a favorable interaction is likely to take place. If a favorable interaction is likely to take place, give the name for the strongest direct intermolecular interaction. Choose from ionic interaction, covalent bond, hydrogen bond, and van der Waals force.

Amino Acid	Will a favorable interaction take place? Circle one.	Strongest Interaction ionic interaction, covalent bond, hydrogen bond, or van der Waals force.
Gln	Yes/No	hydrogen bond
Asp	Yes/No	ionic interaction
Ile	Yes/No	van der Waals
Lys	Yes/No	
Arg	Yes/No	ionic interaction

b) For all cases where a potential interaction seemed **unfavorable**, explain why that interaction may be unfavorable. The interaction between the side chain of lysine and the NH_3^+ group is unfavorable because both are positively charged. The positive charge of the NH_3^+ group will repel and not attract the positive charge of the lysine side chain.

Question 3

a) In your continuing work with the enzyme votase you discover that the enzyme is also a transmembrane protein (part of the protein crosses the lipid bilayer of the cell). Circle the portion of the sequence below that you would expect to be the transmembrane region of the protein.

+NH₃ - glu - trp - asp - arg - his - asp - phe - glu - ser - gly - pro - thr - phe -

ile - trp - leu - ile - trp - leu - val - ile - ala - val - leu - phe - leu - leu - ile -

trp - ala - val - leu - arg - pro - gly - cys - ser - lys - ala - tyr - ala - lys -- val

- cys - ala - gly - cys - ser - asp - lys - gly - glu - COOH

b) Why will the section you circled be embedded in the membrane?

The lipid bilayer of the cell is largely nonpolar and thus it is expected that the region of the protein that passes through this nonpolar layer will also be nonpolar due to the hydrophobic effect. The section underlined above represents the largest stretch of nonpolar amino acids of the residues given and would most likely be the transmembrane section.

c) Draw a schematic of the phospholipid bilayer. Label the lipid hydrocarbon chains, phosphate hydrophilic heads, and water.



Cell Interior

Question 4

a) What is the chemical difference between the pentose sugar in DNA and the pentose sugar in RNA? (See page 48 in text book.)

The sugar in RNA has a hydroxyl (-OH) group in the 2' position while the sugar in DNA has a hydrogen in the 2' position.

b) In the peptide below, **circle** the peptide bonds. What are the amino acids in this peptide? cysteine-alanine-tyrosine-phenylalanine





STRUCTURES OF AMINO ACIDS at pH 7.0

