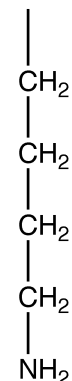
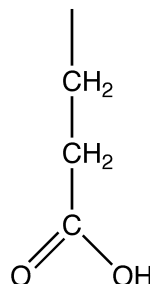
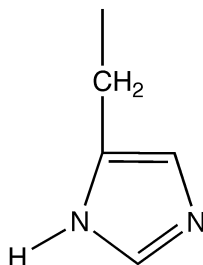
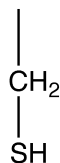
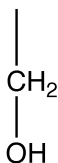


Chem 352, Fundamentals of Biochemistry

Lecture 3, Part I – Supplemental Questions

1. Using the three-letter codes, identify the amino acids that contain the side chains shown below:

A. _____ B. _____ C. _____ D. _____ E. _____



- | | |
|---------------------------------------------------------------------------------------------|-------------------|
| a. Which of these is aliphatic? (<i>Circle all that apply.</i>) | A B C D E |
| b. Which of these can hydrogen bond to water? (<i>Circle all that apply.</i>) | A B C D E |
| c. Which of these is aromatic? (<i>Circle all that apply.</i>) | A B C D E |
| d. Which of these is charged at neutral <i>pH</i> values? (<i>Circle all that apply.</i>) | A B C D E |
| e. Which of these is hydrophobic? (<i>Circle all that apply.</i>) | A B C D E |
| f. Which of these is basic? (<i>Circle all that apply.</i>) | A B C D E |
| g. Which of these is acidic? (<i>Circle all that apply.</i>) | A B C D E |
2. What contributions did G. N. Ramachandran and Linus Pauling make to our understanding of protein structure?
3. Draw the chemical structure for the tripeptide His-Asp-Lys in its proper charged state at *pH* 7.

- On your structure, label one example of a ϕ , a ψ and an ω bond.
- On your structure, label one example of a *peptide bond*.
- Draw a titration curve for this tripeptide?
- What is the isoelectric *pH* (*pI*) for this peptide?

4. Determine the amino acid sequence of a polypeptide given the following data:
- Treatment with the proteolytic enzyme *carboxypeptidase*, which removes amino acid residues sequentially from the C-terminal end, releases **Val**.
 - The first amino acid released from an Edman degradation of the full length polypeptide is **Ser**.
 - Treatment with the proteolytic enzyme *trypsin*, followed by Edman degradation of the resulting fragments, produced the following five fragments:
Phe-Glu-His-Lys, Phe-Gly-Arg, Pro-Val, Ser-Tyr-Ser-Lys, Trp-Gly-Lys.
 - Treatment with the proteolytic enzyme *chymotrypsin*, followed by Edman degradation of the resulting fragments, produced the following five fragments:
Gly-Arg-Trp, Gly-Lys-Pro-Val, Ser-Lys-Phe, Glu-His-Lys-Phe, Ser-Tyr.
 - Total hydrolysis of the polypeptide reveals the following composition:
Arg, Glu, Gly(2), His, Lys(3), Phe(2), Pro, Ser(2), Trp, Tyr, Val.
5. Draw the chemical structure of the peptide that you sequenced in problem 4 in the charged state that it will have at *pH* 8.