NAME: $\qquad$ ANSWERS

Quiz 2 October 7, 2009

1. Shown below is the backbone of the protein myoglobin. The positions of the alpha carbons of the two histidines that are important in iron binding are shown with an asterisk, though the side groups are not drawn in. (6 points)

Identify the following landmarks in this image by writing very small letters a-f on the image:
a) the iron atom
b) the heme group
c) the white gly carboxy terminus
d) the distal His (keeps $\mathrm{O}_{2}$ bent)
e) the proximal His (ligates metal)
f) the oxygen ligand (it's TEENY)
2. First give a definition, and then identify the forces (i.e. covalent, $\mathrm{H}-$ bonds, disulfide, van der Waals, salt bridges) and magnitude (weakest, weak, strong) that stabilize primary, secondary, tertiary, and quaternary structures? ( 20 points)


| Structure Level for <br> MYOGLOBIN | Definition (2 pts each) | Forces (2 pts ea) | Magnitude <br> (1 pt ea) |
| :--- | :--- | :--- | :--- |
| Primary: | The sequence of amino acids N <br> to C terminal bound together by <br> peptide bonds. | covalent | strong |
| Secondary | The individual shapes such as $\alpha$ <br> helix and $\beta$ strands that a stretch <br> of amino acids can adopt. | H bonds | weak |
| Tertiary | The manner in which individual <br> shapes (helices, sheets) pack up <br> against one another. | Van der waals, H <br> bonds, some <br> covalent disulfide | Weakest, <br> weak, and <br> strong |
| Quaternary | Interactions made between <br> subunits in a multi-subunit <br> protein. | Why none? There <br> are no subunits in <br> Mb. | Weak to <br> strong |

## 3. Binding Curves for Myoglobin and Hemoglobin (16 points)


a. Explain the directionality of the transport of $\mathrm{O}_{2}$ in the tissues at normal blood/lymph pH of 7.4 using the graph at left. (Hint: your answer should begin with the phrase....As you can see from the graph..) 4 pts

As you can see from the graph, at $\mathrm{P}_{\mathrm{O} 2}$ of $25-40 \mathrm{~mm} \mathrm{Hg}$, the Hb is only about $25 \%$ saturated while Mb is about $95 \%$ saturated. This suggests that the Hb has released the $\mathrm{O}_{2}$ and the Mb has taken it up, which is as it should be as the Hb 's job is to transport AND RELEASE $\mathrm{O}_{2}$ to the tissues for Mb to take it up and bring it to the mitochondria for aerobic respiration.
b. What do the three curves below tell you about how Hb 's oxygen binding affinity changes as the pH drops from 7.6 to 7.2 to 6.8 ? Your answer should include a comparison of how the Kds change for Hb as the pH drops. 4 pts


As the pH drops, the Hb binding curve is shifting to the right. This means that if we compare the saturation of Hb with $\mathrm{O}_{2}$ (Y value) at any one X value, we see that the saturation decreases as the pH decreases. This also means that Hb is becoming a poorer and poorer binder of $\mathrm{O}_{2}$, or that the Kd (dissociation) is increasing as the pH is decreasing.
c. What "state" (R or T) of the Hb is most prevalent in the tissues at pH 6.8 ? Explain. 4 pts

Since the deoxygenated state is more prevalent at lower pH , this suggests that the T state, or tense deoxy state predominates. This makes sense because these tissues are stressed and need oxygen! NOW
d. What are some of the small molecules or ions that bind to this state of Hb ? 4 pts
$\mathrm{H}^{+}$, bisphosphoglycerate (DPG or BPG), $\mathrm{CO}_{2}$ all bind to the T state of Hb
4. The last of the eight helices in Mb is very long one, extending from Ala125 to Leu149. The end to end distance of this helix is shown as a dotted line on this image below (rotated from the view on the first page) and measured to be $36.04 \AA$. ( 8 points)

a. How many amino acids are in this helix? (note the answer to this is a whole number, which never limits the significant figures) 1 pt Ala 149 to Leu 125 is $149-125+1=25$ amino acids
b. The length of the helix was given above as $36.04 \AA$. Using the conversion factors on the back calculate the length in picometers, nanometers, micrometers, and meters. Pay attention to significant figures. 4 pts
$36.04 \AA\left(10^{-10} \mathrm{~m} / 1 \AA\right)\left(10^{12} \mathrm{pm} / 1 \mathrm{~m}\right)=3.604 * 10^{3} \mathrm{pm}$ $36.04 \AA\left(10^{-10} \mathrm{~m} / 1 \AA\right)\left(10^{9} \mathrm{~nm} / 1 \mathrm{~m}\right)=3.604 \mathrm{~nm}$ $36.04 \AA\left(10^{-10} \mathrm{~m} / 1 \AA\right)\left(10^{6} \mathrm{um} / 1 \mathrm{~m}\right)=3.604 * 10^{-3} \mathrm{um}$ $36.04 \AA *\left(10^{-10} \mathrm{~m} / 1 \AA\right)=3.604 * 10^{-9} \mathrm{~m}$ All answers above have four sig figs
c. Calculate the distance per amino acid for this helix (use distance units of $\AA$ ). Pay attention to significant figures. 1.5 pts
$36.04 \AA / 25=1.4416 \AA$ or $1.442 \AA$ to the correct (4) significant figures
d. These amino acids find themselves in an alpha helix, which is a fairly compressed structure for the amino acid backbone. IF these amino acids found themselves in a beta strand ( $3.4 \AA$ per amino acid), how long would the strand be? Watch your significant figures. 1.5 pts
$3.4 \AA \times 25=85 \AA$ (note two significant figures in answer)

These conversion factors are exact, and will not limit the number of significant figures in your answer:
$1 \mathrm{~m}=10^{2} \mathrm{~cm}=10^{3} \mathrm{~mm}=10^{6} \mathrm{um}=10^{9} \mathrm{~nm}=10^{12} \mathrm{pm}$
$1 \AA=10^{-10} \mathrm{~m}=10^{-8} \mathrm{~cm}$

