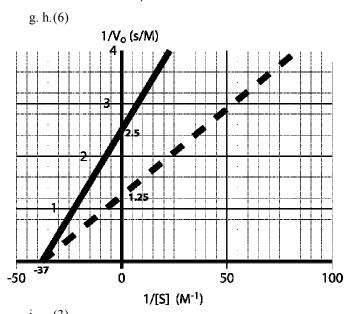
- 1. (5) a, c, d, f, g
- 2. (2) c, a, b
- 3. (4 pts) (1) b; (2) c; (3) a; (4) d
- (3) A β-barrel is like a β-sheet rolled up so the edges H-bond. If there are an odd number of strands, two bonded strands will be parallel (see edge strands in pic), and the barrel is not wholly antiparallel.
- 5. (2) False *cholesterol clusters in rafts*
- 6. (2) False [S] is constant because  $[S] >> [E_T]$
- 7. (2) True
- 8. (2) True

9. a. (3)  $\theta = \frac{p0_2}{P_{50} + p0_2}$   $\theta(P_{50} + p0_2) = p0_2$   $\theta \cdot P_{50} + \theta \cdot p0_2 = p0_2$   $\theta \cdot P_{50} = p0_2 - \theta \cdot p0_2$   $\theta \cdot P_{50} = p0_2(1 - \theta)$  $P_{50} = p0_2 \frac{(1 - \theta)}{\theta} = 44 \text{ torr } \cdot \frac{0.1}{0.9} = 4.9 \text{ torr}$ 

- b. (4) x-axis: pO<sub>2</sub> (torr); y-axis: θ, with values 0 to 1.0;
  'N' curve: hyperbolic, passing through (2.8, 0.5) and approaching 1.0 in y; 'A' curve (right of 'N' curve): hyperbolic, passing through (4.9, 0.5) and (44, 0.9)
- c. (2) lower
- d. (3) Affinity depends on the rate constants for binding and unbinding. To have a lower affinity, O<sub>2</sub> would unbind (dissociate) faster from the altered myoglobin.
- 10. a. (2) hydrolase, phosphatase
  - b. (2) ligase, synthetase
  - c. (1) isomerase
- 11. a. (2)  $\Delta G_6$ 
  - b. (2)  $\Delta G_2 \Delta G_6$  or  $\Delta G_2 \Delta G_8$  accepted
  - c. (1) unimolecular  $(1^{st} order)$
- a. (3) Substrates are positioned for maximal reactivity, because they bind the enzyme at an optimal orientation and distance relative to each other and to the enzyme's reactive groups.
  - b. (2) True
- 13. a. (1) in red blood cells
  - b. (2)  $CO_2 + H_2O \implies HCO_3^- + H^+$
  - c. (2) lyase
  - d. (5) In the capillaries, where CO<sub>2</sub> levels are high, the enzyme catalyzes the formation of bicarbonate and H<sup>+</sup>. The protons protonate Hb and stabilize the T-state, promoting release of O<sub>2</sub>. Bicarbonate exiting the RBCs results in the entry of Cl<sup>-</sup>, which also stabilizes the T-state.
  - e. (2) The maximal number of  $S \rightarrow P$  conversions per single enzyme per unit time.

f. (2) cat. eff. = 
$$\frac{k_{cat}}{k_m}$$
  
 $K_m = \frac{k_{cat}}{\text{cat. eff.}} = \frac{4 \times 10^5 \text{s}^{-1}}{1.5 \times 10^7 \text{M}^{-1} \text{s}^{-1}}$   
= 0.027 M or 27 mM

(f. continued) (2) 
$$V_{max} = k_{cat} [E_T] = 4 \times 10^5 s^{-1} \cdot 10^{-6} M = 0.4 M/s$$

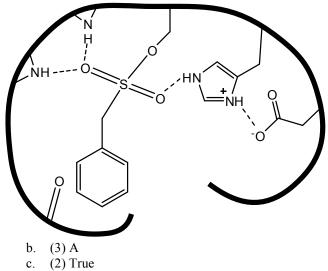


$$K_m = \frac{k_{cat}}{cat \text{ eff.}} = \frac{10^6 \text{s}^{-1}}{8.3 \times 10^7 \text{M}^{-1} \text{s}^{-1}} = 0.012 \text{ M } or 12 \text{ mM}$$
  

$$K_m \text{ for CO}_2 \text{ is lower, so affinity for } \underline{CO}_2 \text{ is higher}$$
  
j. (1) False

- 14. a. (2) Retaining glycosidases hydrolyse glycosidic bonds and maintain the anomeric configuration in the product
  - b. (2) There are two  $SN_2$  steps
- 15. (4) b, d, e





- 17. a. (3) Cysteine has a lower pKa, so it will deprotonate more readily, to form the reactive nucleophile.
  - b. (3) Because of cysteine's lower pKa, its deprotonated form is less nucleophilic than serine's deprotonated hydroxyl.