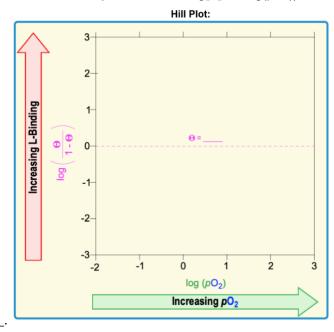
- The Hill Plot is a \_\_\_\_\_ graph plotting the y-value & x-value of the Hill Equation respectively on the y-axis & x-axis.
- •Slope of the line on a Hill Plot (slope = \_\_\_\_) denotes the Hill constant (nH) & the degree of L-binding-site interactions.
  - $\square$  Recall: for both Mb & Hb, L = O<sub>2</sub>, and [O<sub>2</sub>] can be expressed with \_\_\_\_\_ (which means: log[L]  $\rightarrow$  log(pO<sub>2</sub>)).

Equation of a Line: Hill Equation:  $y = mX + b \qquad log(\frac{\theta}{1 - \theta}) = n_H log([L]) - n_H log(K_d)$ 



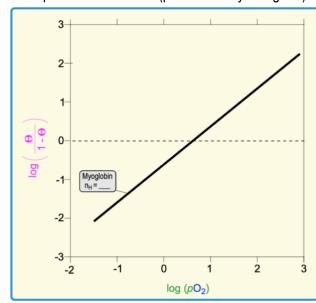
- •On a Hill Plot, the \_\_\_\_\_-intercept reveals when  $\theta$  = \_\_\_\_\_.
  - □ Recall: x-intercept is ALWAYS the x-value when y = \_\_\_\_.
  - $\square$  Note: y-value of Hill Equation ( $\log \left(\frac{\theta}{1-\theta}\right)$ ) will equal zero when the value of θ = \_\_\_\_\_.

### Mb's Hill Plot

- •Since myoglobin (Mb) only has 1-subunit, it is NOT an allosteric protein & has \_\_\_\_\_ cooperativity.
  - $\Box$  Recall:  $n_H = slope = \_$  when there is *no* cooperativity.
  - $\Box$  Since x-intercept indicates the [L] where  $\theta$  = \_\_\_\_\_, x-intercept also indicates K<sub>d</sub> (protein affinity for ligand).

Mb = \_\_\_ subunit = \_\_\_\_ cooperativity (slope = **n**н = 1)

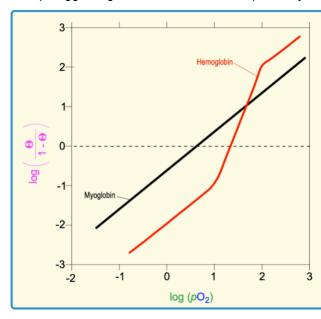




#### **Hb's Hill Plot**

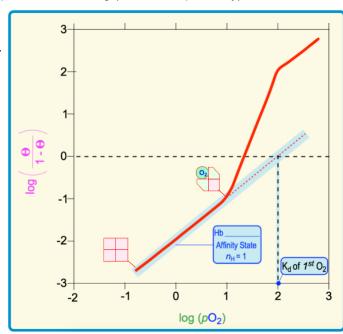
- •Unlike Mb (which forms a single line), Hb's PL-data seems to form \_\_\_\_\_ identifiable lines when plotted on a Hill Plot.
  - □ *Two* of Hb's lines are \_\_\_\_\_ to Mb's line, meaning they have the *same* slope (slope = \_\_\_\_ = 1).
  - □ Recall: a slope or n<sub>H</sub> of 1 means \_\_\_\_\_ cooperativity.
  - ☐ Hb binds its \_\_\_\_\_ and \_\_\_\_\_ O₂ non-cooperatively.
- ●HOWEVER, Hb's 3rd line has a *different/greater* slope (slope = n<sub>H</sub> = \_\_\_\_\_), suggesting \_\_\_\_\_ cooperativity.





## Hb's Lowest O2-Affinity State

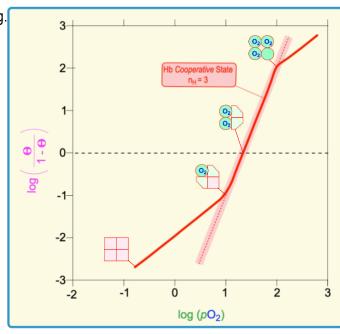
- •Recall: Hb is NOT always displaying cooperativity; Hb binds its \_\_\_\_\_ & \_\_\_\_ O₂ non-cooperatively (slope = n<sub>H</sub> = 1).
  - ☐ Hb subunits equally & \_\_\_\_\_ compete for O₂-binding (without cooperativity) until 1st O₂ binds.
  - □ This line represents Hb's \_\_\_\_\_ O₂-affinity.
  - □ Lowest O₂-affinity means \_\_\_\_\_ K<sub>d</sub> for 1<sub>st</sub> O₂.



### **Hb's Cooperative State**

- After the 1st O₂ binds to Hb, Hb subunits begin to display \_\_\_\_\_ cooperativity (slope = nH = 3).
  - □ Recall: Hb's O<sub>2</sub>-binding behavior explained via a \_\_\_\_\_\_ of *concerted* & *sequential* models (n<sub>H</sub> ≠ n).
  - ☐ Hb displays *positive* cooperativity from the 1st bound O<sub>2</sub> up *until* the 3rd O<sub>2</sub> binds.
  - ☐ Hb subunits are NOT equally competing for O2-binding.

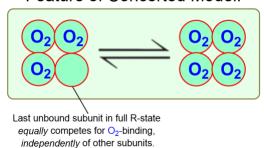


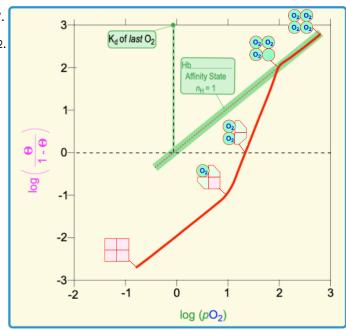


## Hb's Highest O<sub>2</sub>-Affinity State

- After the 3rd O₂ binds to Hb, the 4th and final O₂ binds \_\_\_\_\_-cooperatively, just like the 1st O₂ did (slope = nH = 1).
  - □ Once again, Hb subunits *equally* & *independently* compete for binding to the *last* O<sub>2</sub> (\_\_\_\_\_ cooperativity).
  - □ Last unoccupied Hb subunit is in the full R-state experiencing \_\_\_\_\_ model features (symmetry rule).
  - □ This line represents Hb's \_\_\_\_\_\_ O₂-affinity.
  - □ Highest O2-affinity means \_\_\_\_\_ Kd for 4th O2.

### Feature of Concerted Model:

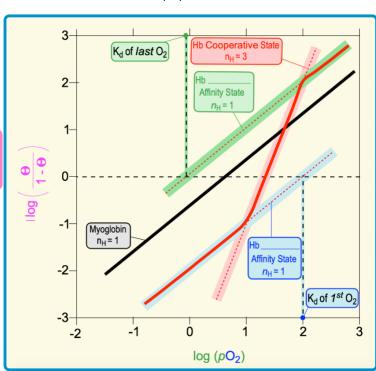




### **Hill Plot Breakdown**

●Hill plots *visually* display **PL**-affinities (K<sub>d</sub>) and the degree of \_\_\_\_\_\_(n<sub>H</sub>) in a PL-interaction.

 $\log\left(\frac{\theta}{1-\theta}\right) = n_{H}\log(L) - n_{H}\log(K_{d})$ 



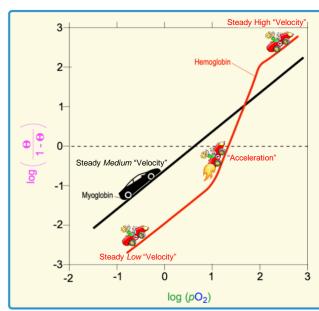
# "Velocity" & "Acceleration"

- ●Mb's & Hb's O₂-binding can be thought of as \_\_\_\_\_ (for O₂-affinity) and \_\_\_\_\_ (for cooperativity).
  - $\ \square$  Mb ALWAYS maintains the same, relatively high/medium velocity with  $\_\_\_$  acceleration.
  - $\hfill\Box$  Hb starts off with low velocity and does NOT begin accelerating \textit{until} it binds it's 1st O2.
  - □ Once Hb binds 3 O<sub>2</sub>, it reaches/maintains its max velocity & stops accelerating.

Steady = No "acceleration" = No cooperativity

"Velocity" = O<sub>2</sub>-affinity

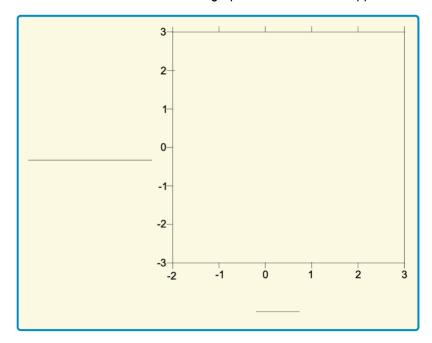
"Acceleration" = Positive Cooperativity



**PRACTICE:** Which of the following situations would produce a Hill plot with n<sub>H</sub> < 1.0?

- a) The protein is a single polypeptide with a single ligand binding site. As purified, the protein preparation is heterogeneous, containing some protein molecules that are partially denatured and having a lower binding affinity.
- b) The protein is a single polypeptide with two ligand binding sites, each having a different affinity for the ligand.
- c) The protein has multiple subunits, each with a single ligand-binding site. Binding of ligand to one site decreases the binding affinity of other sites for the ligand.

**PRACTICE:** Label the axis of the Hill Plot below & fill-in the graph with Mb's & Hb's approximate O<sub>2</sub>-binding data.



PRACTICE: The slope of a Hill plot for hemoglobin \_\_\_\_\_\_\_; whereas that for myoglobin \_\_\_\_\_\_.

- a) is about 3 in its cooperative state; is 1.0.
- b) decreases at low pO<sub>2</sub>; is constant at all pO<sub>2</sub>.
- c) increases at high pO<sub>2</sub>; curves upward for all pO<sub>2</sub>.
- d) Is 1.0; is about 4.
- e) Both A and B.