



Hemoglobin

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Another look at binding

- $\text{Mb-O}_2 \xrightleftharpoons[k_{\text{on}}]{k_{\text{off}}} \text{Mb} + \text{O}_2$
- K_d is a result of two rate constants
 - Rate of dissociation, k_{off}
 - Rate of association, k_{on}
- k_{on} is the result of a collision
 - k_{on} for myoglobin relies on the presence of O_2
- $k_{\text{on}} = \#$ per mole per second
- $k_{\text{off}} = \#$ per second





What is K_d ?

- On and off rates are defined by their rate constants

- $k_{\text{on}} = \# \text{ mole}^{-1} \text{ sec}^{-1}$

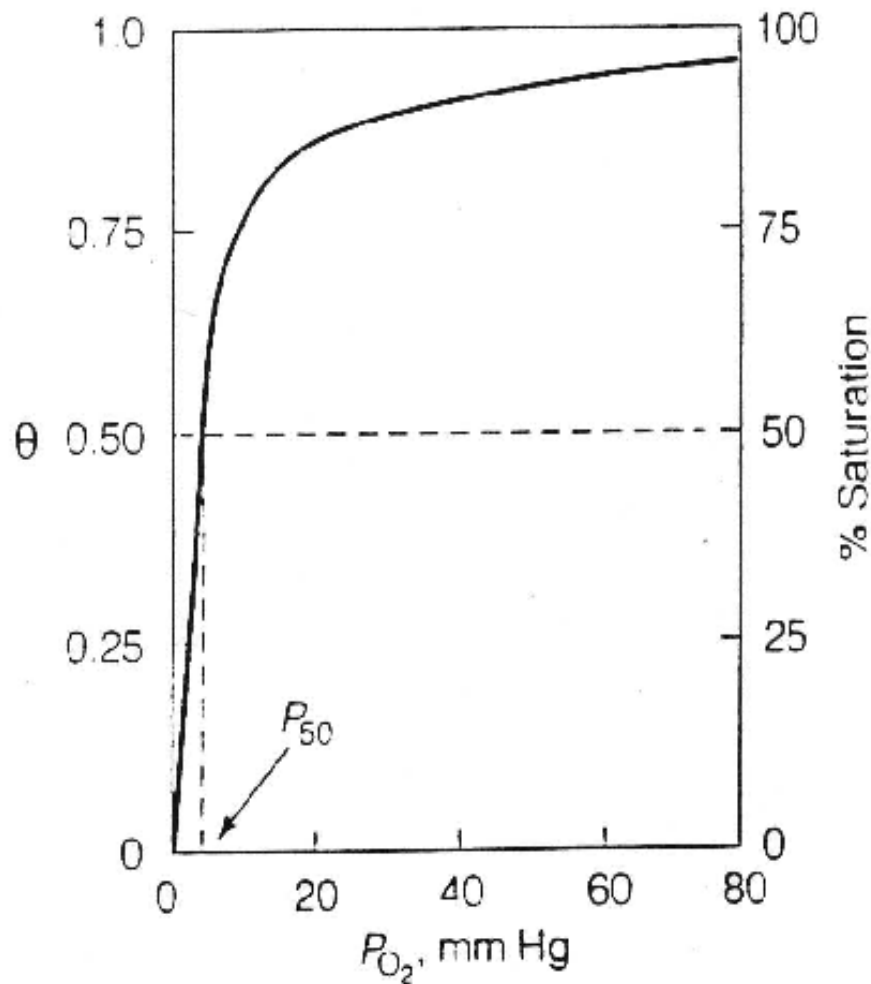
- $k_{\text{off}} = \# \text{ sec}^{-1}$

- $K_a = \frac{k_{\text{on}}}{k_{\text{off}}} = \frac{\# \text{ M}^{-1} \text{ sec}^{-1}}{\# \text{ sec}^{-1}} = \text{M}^{-1} = \text{“per molar?”}$

- $K_d = \frac{k_{\text{off}}}{k_{\text{on}}} = \frac{\# \text{ sec}^{-1}}{\# \text{ M}^{-1} \text{ sec}^{-1}} = \text{M} = \text{Useful !}$



Why K_d is very useful



$$K_d = \frac{[\text{Mb}_{\text{free}}][\text{O}_2]}{[\text{Mb-O}_2]}$$

$$\text{Mb}_{\text{free}} = [\text{Mb}_T] - [\text{Mb-O}_2]$$

We can make a substitution and rearrangement to get something very useful!

Y = fractional saturation

$$Y = [\text{Mb-O}_2] / [\text{Mb}]_T$$

$$Y = [\text{O}_2] / (K_d + [\text{O}_2])$$

* Now we can use K_d to know the fraction of protein bound at ANY O_2 concentration!

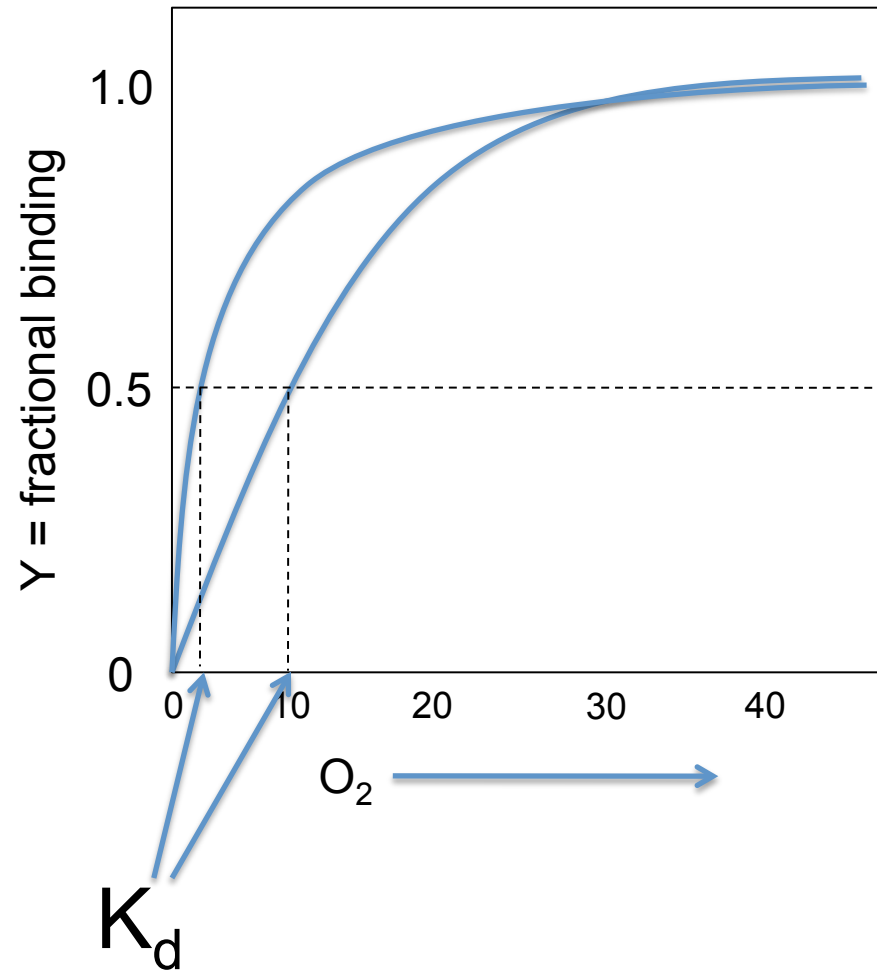
What does a change in K_d look like?

- Fractional Binding curves

- $$K_d = \frac{[Mb_{free}][O_2]}{[Mb-O_2]}$$

– $Mb_{free} = [Mb_T] - [Mb-O_2]$

- $$Y = \frac{[O_2]}{(K_d + [O_2])}$$





Thinking some more.

- Constant K_d .
 - Increase the amount of Oxygen.
 - What happens?
- At the same oxygen concentration.
 - You measured more Mb-O₂ for Myoglobin A than for mutant Myoglobin B?
 - MyoA Mb-O₂ > MyoB Mb-O₂
 - Does Mb_{free} or Mb_T change? How?
 - MyoA K_d ? MyoB K_d






Now On to Hemoglobin!


- Background
- Structure
- Cooperative binding.





Hemoglobin

- Oxygen carrying protein in your blood.
 - Oxygen picked up in lungs and transported to tissue throughout the body.
 - Oxygen gets transferred to Myoglobin in muscle
 - There are 2.3×10^8 hemoglobin molecules in a single red blood cell.
 - Hemoglobin looks different and binds oxygen differently than Myoglobin
- 



Hemoglobin is “related” to Myoglobin

- But:
 - Hemoglobin is a tetrameric protein
 - Two different isoforms α and β
 - Each subunit has a very similar “fold” to myoglobin
 - Yet only 27% of the residues are identical



Primary Sequence Alignment

```

sp|P68871|HBB_HUMAN      MVHLTPEEKSAVTALWGKVVN--DEVGGEALGRLLVVYPWTQRFFESFGDLSTPDAVMGN
sp|P69905|HBA_HUMAN      -MVLSPADKTNVKAAWGKVGAHAGEYGAEALERMFLSFPTTKTYFPHFD-----LSHGS
sp|P02144|MYG_HUMAN      -MGLSDGEWQLVLNVWGKVEADIPGHGQEVLIIRLFKGHPEKLEKFDKFKHLKSEDEMKAS
                          : * : : * **** . * * * * : : . * * * * * ..

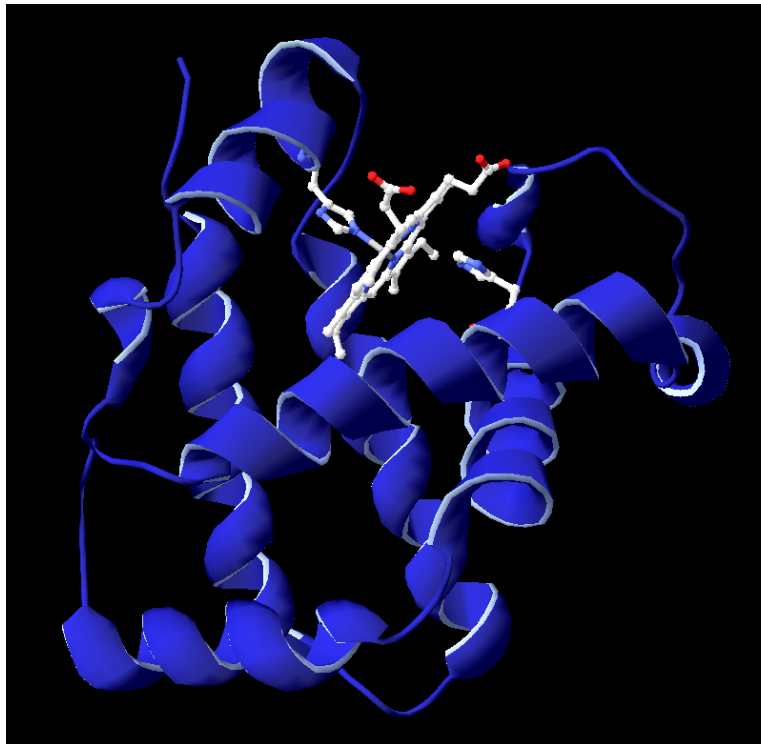
sp|P68871|HBB_HUMAN      PKVKAHGKKVLGAFSDGLAHLNLIKGTFFATLSELHCDKLHVDPENFRLLGNVLCVLAHH
sp|P69905|HBA_HUMAN      AQVKGHGKQVADALTNVAHVDDMPNALSALSDLHAHKLRVDPVNFKLLSHCLLVTLAAH
sp|P02144|MYG_HUMAN      EDLKKHGATVLTALGGILKKKGHHEAEIKPLAQSHATKHKIPVKYLEFISECIIQVLQSK
                          . : * * * . * * : : : . : * : : * . * : : : : : : : : . * :

sp|P68871|HBB_HUMAN      FGKEFTPPVQAAYQKVVAGVANALAHKYH-----
sp|P69905|HBA_HUMAN      LPAEFTPAVHASLDKFLASVSTVLTISKYR-----
sp|P02144|MYG_HUMAN      HPGDFGADAQGAMNKALELFRKDMASNYKELGFQG
                          : * . . . : * : . . : : * :

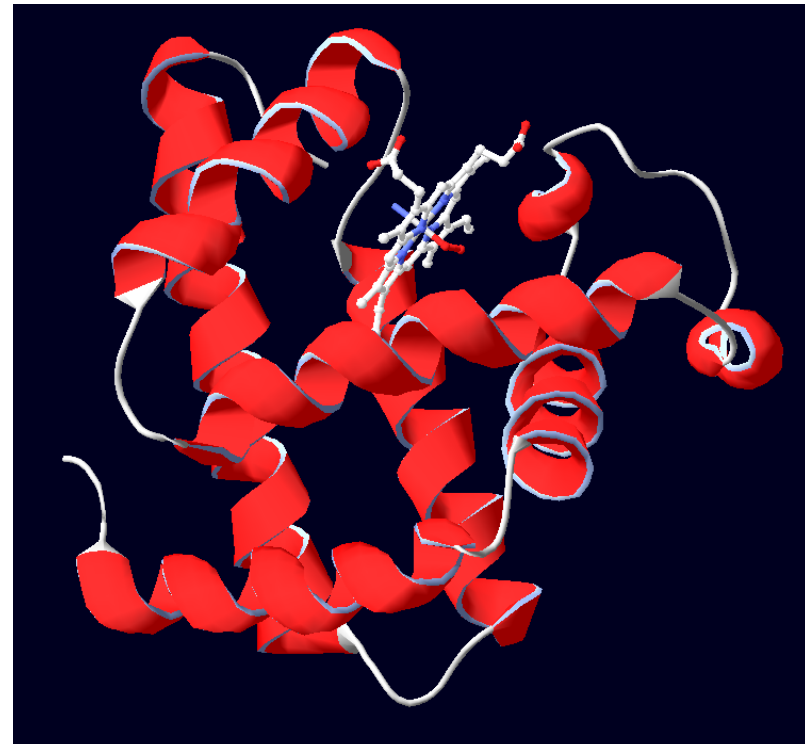
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- One dot?
- Two dots?
- Star?
- Colors?

Hemoglobin

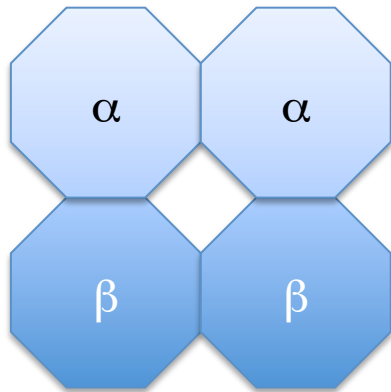


Hemoglobin β



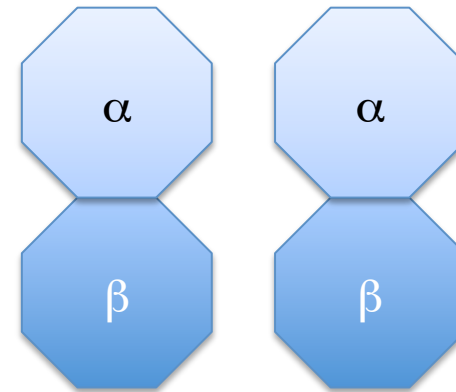
Myoglobin

Hemoglobin is a tetramer



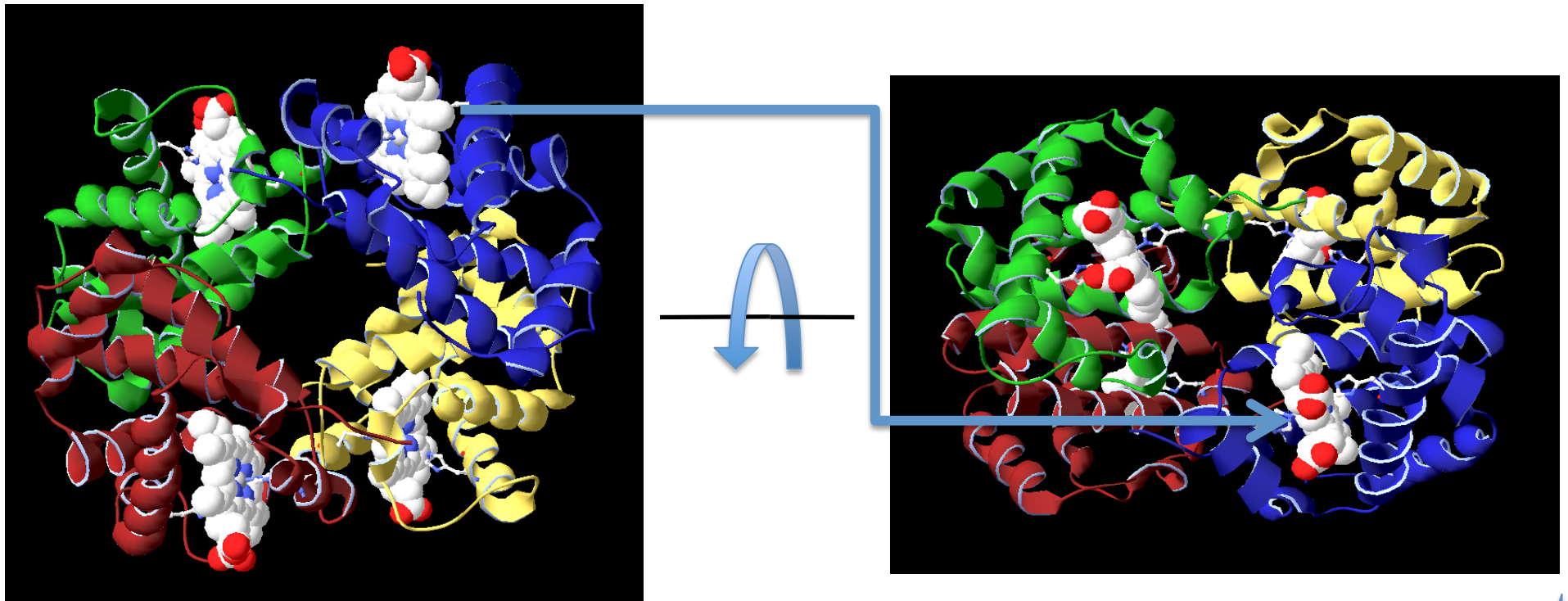
2 α 2 β units
4 peptide chains
Tetramer
Binds four Hemes
Binds four O₂

“Dimer of Dimers”

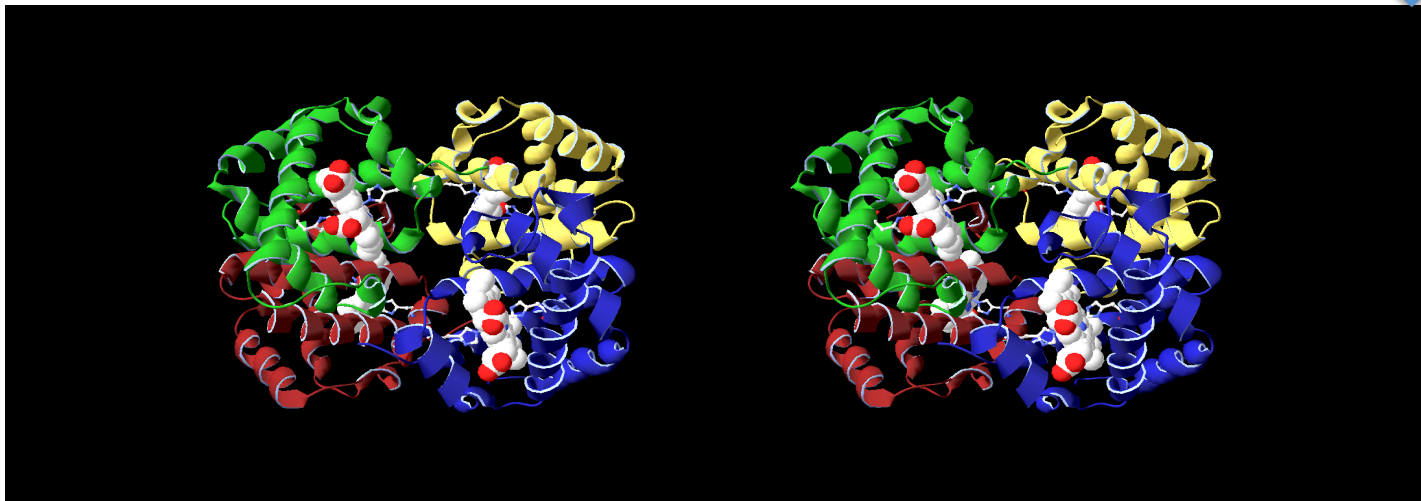
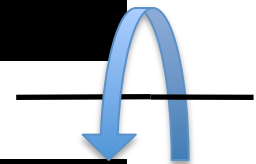
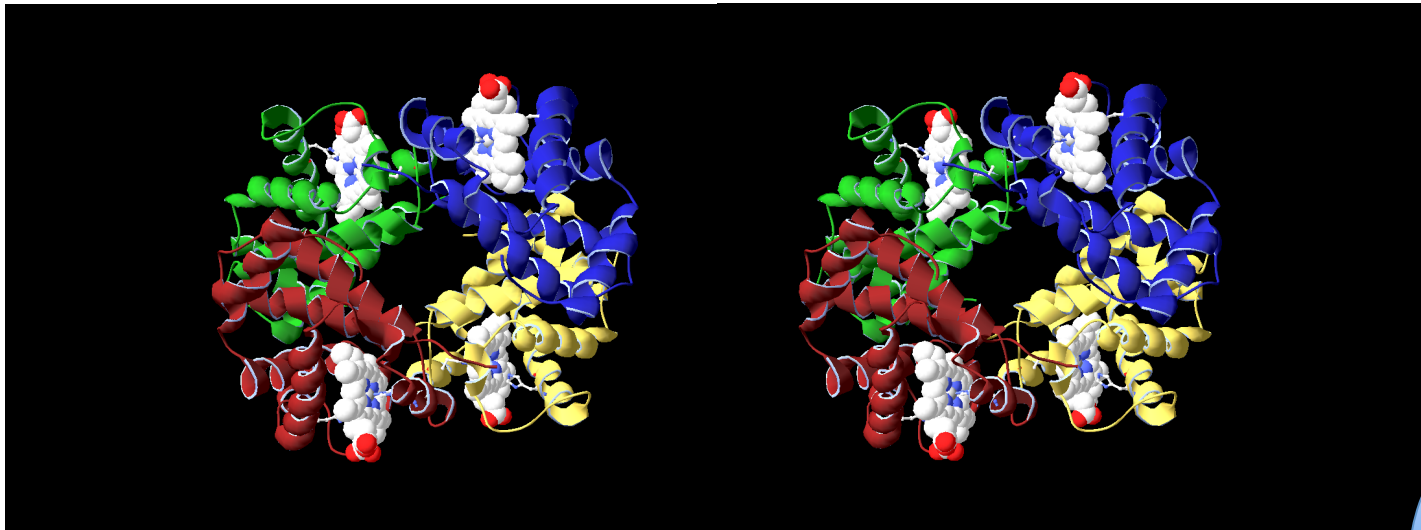


Single unit
1 peptide chains
Monomer
Binds one Heme
Binds one O₂

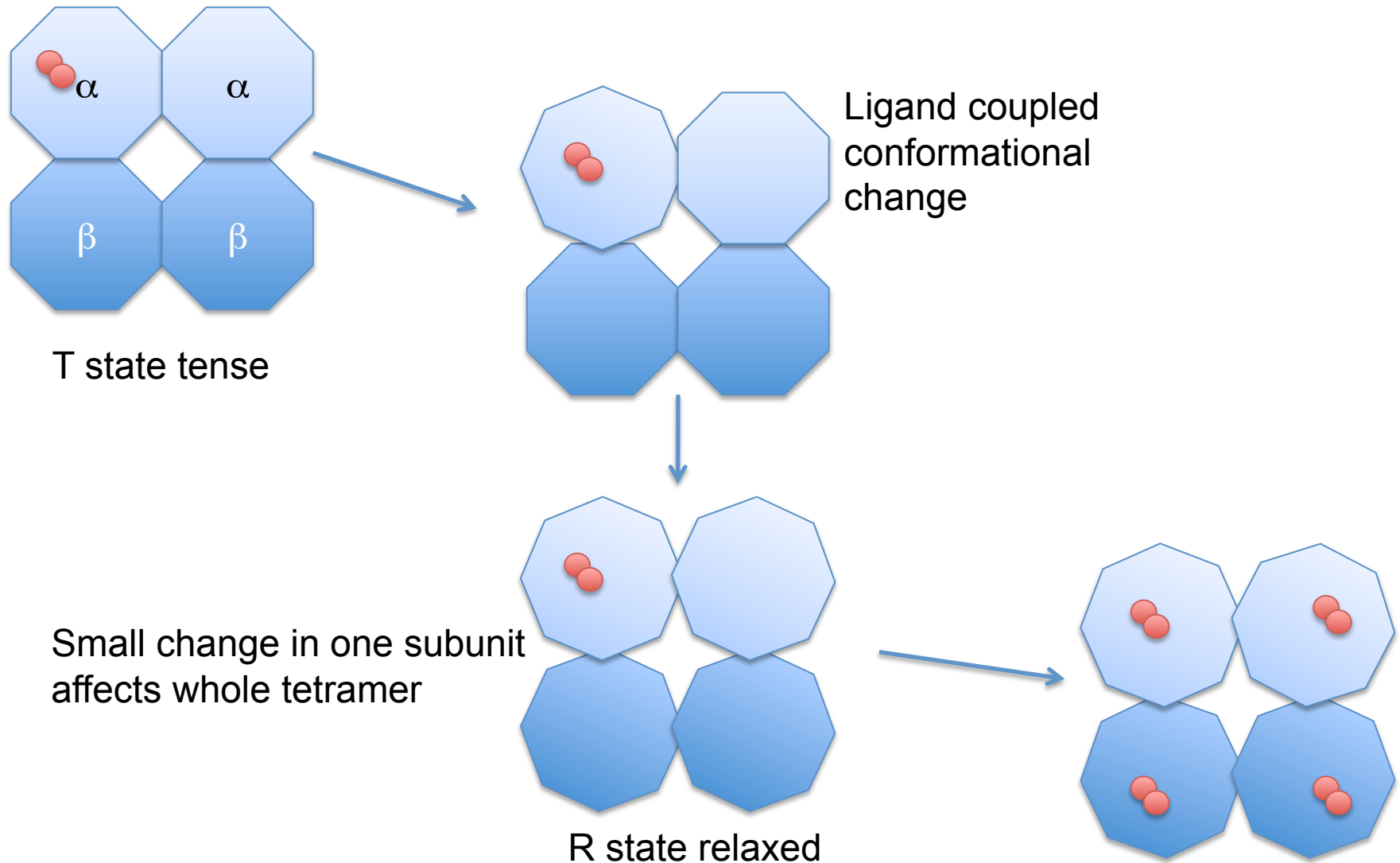
Hemoglobin Quaternary Structure



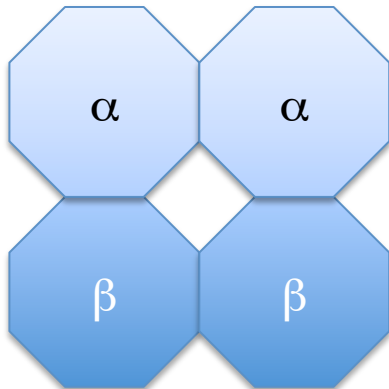
Hemoglobin Quaternary Structure



Hemoglobin has allostery

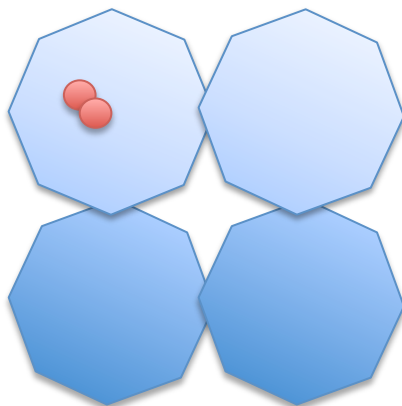


Each state has a different K_d



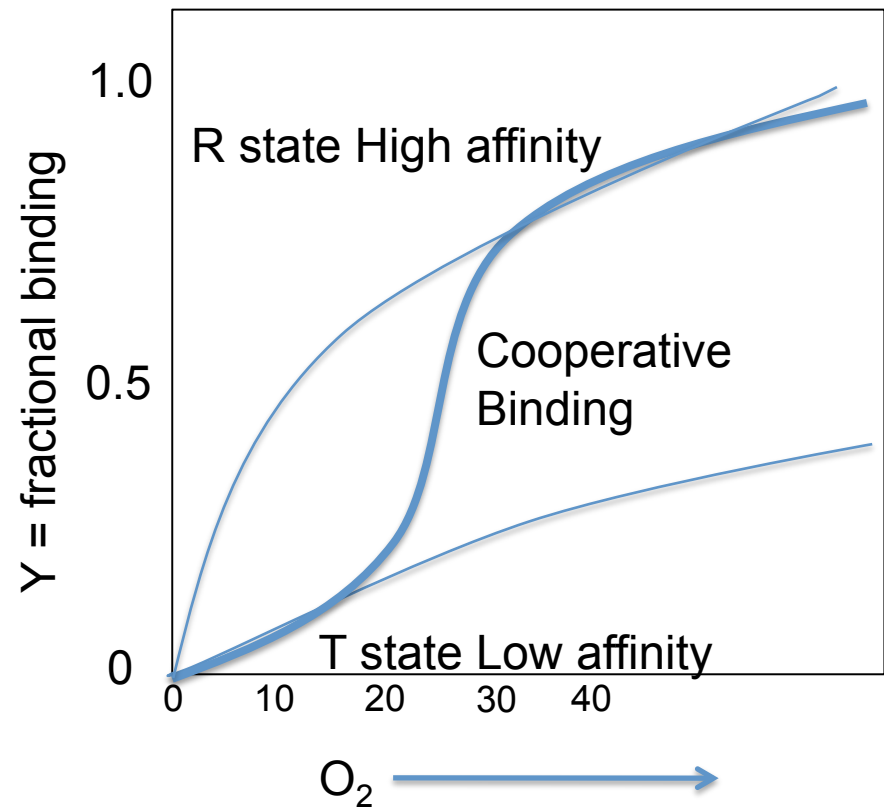
T state tense

High K_d



R state relaxed

Low K_d



Modeling cooperative binding

For Myoglobin:

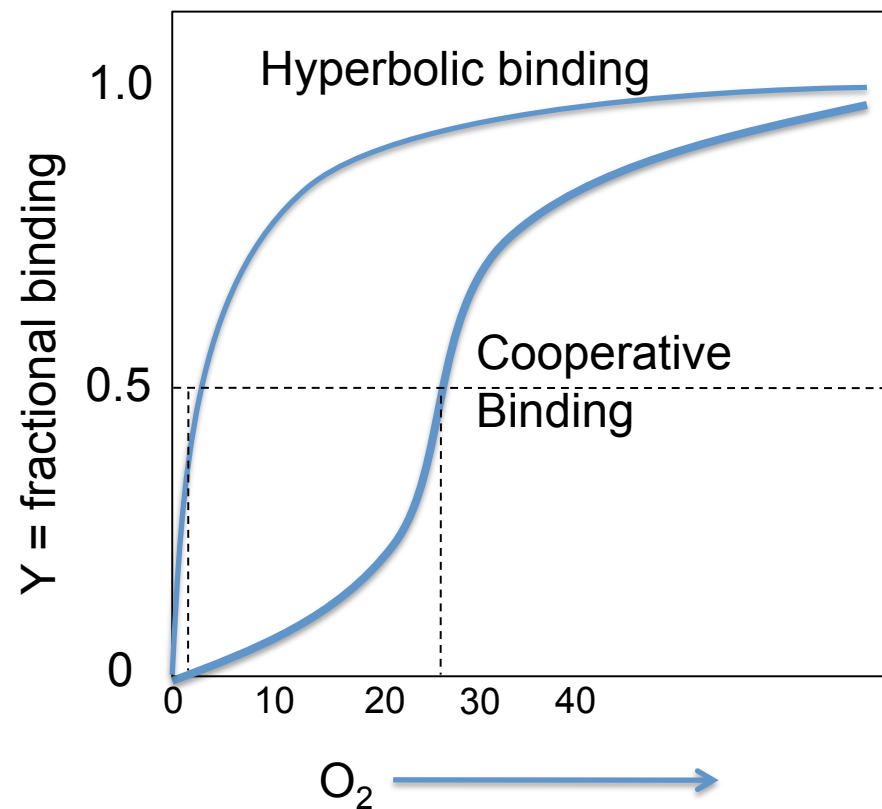
$$Y = \frac{[O_2]}{(K_d + [O_2])}$$

Hyperbolic binding
Simple one phase binding

For Hemoglobin:

$$Y = \frac{[O_2]^n}{(K_d^n + [O_2]^n)}$$

Sigmoidal binding
Where **n** is extent of cooperativity





More binding this afternoon

- In lab we will be looking at two things that bind Hemoglobin and Myoglobin
 - Oxygen
 - Carbon Monoxide

