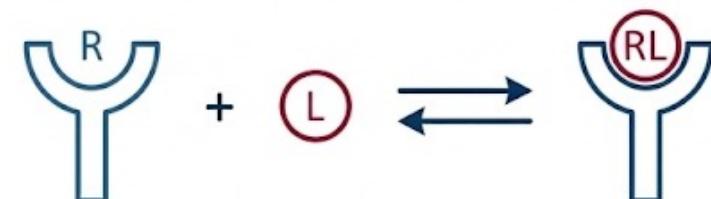
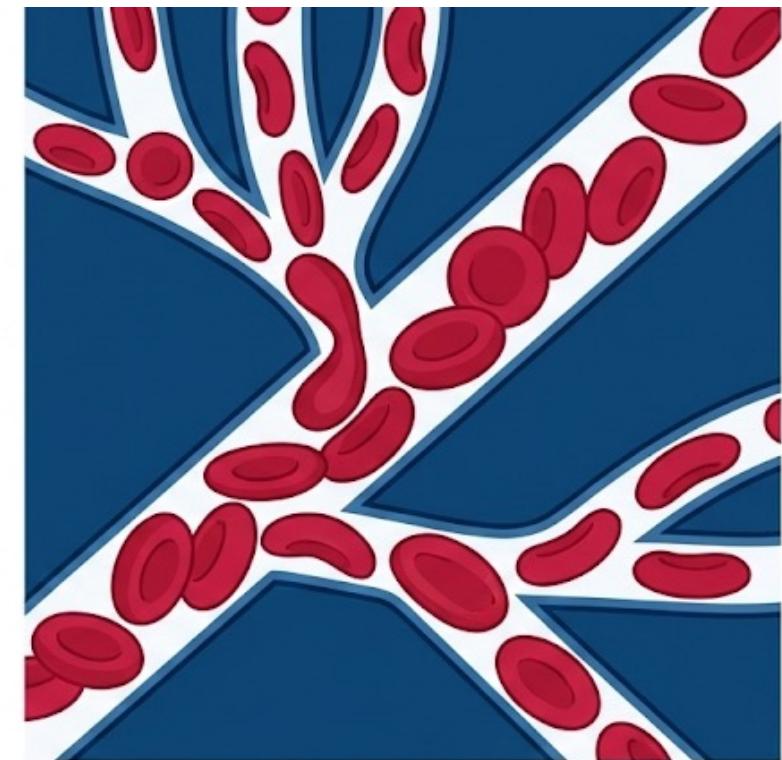
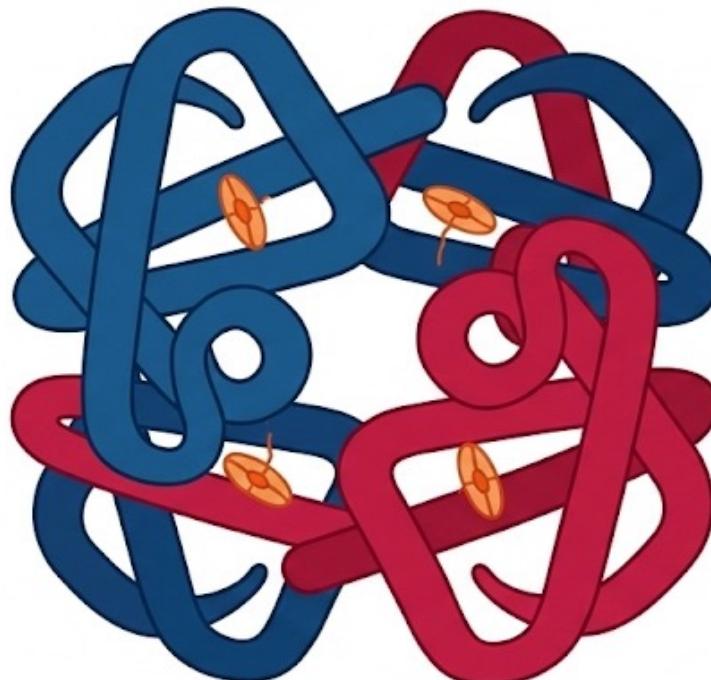
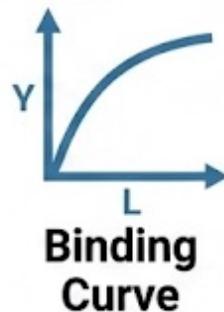


Chapter 5

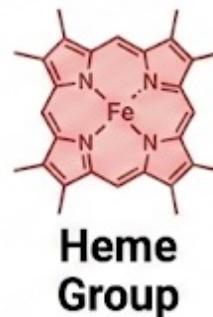
Protein Binding, Molecular Recognition, and Allostery



By end of lecture you can...



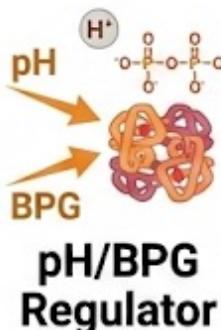
1. Understand the qualitative relationship where ligand concentration affects receptor saturation (hyperbolic curve).



2. Recognize the core principle of oxygen binding to the iron within the heme prosthetic group of myoglobin and hemoglobin.



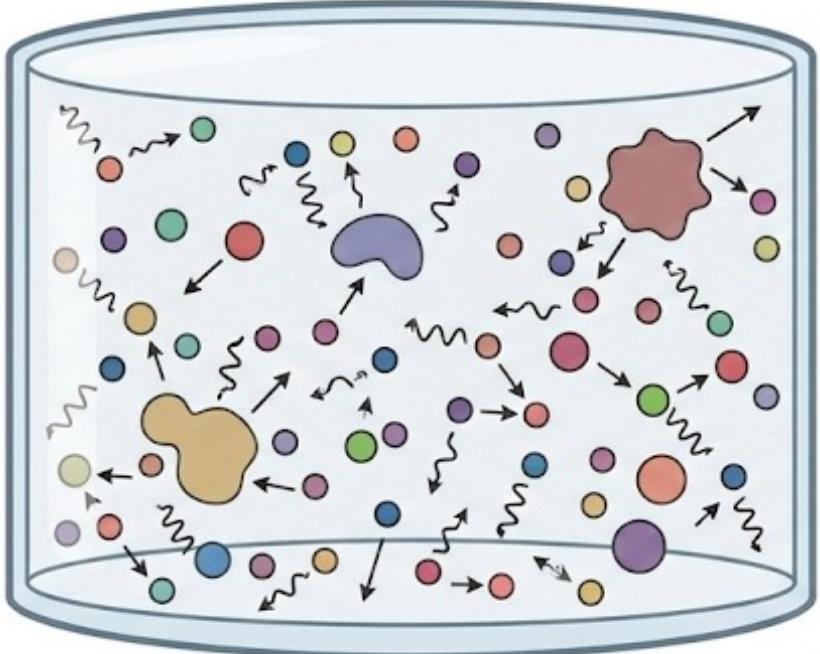
3. Explain how cooperative binding (sigmoid curve) allows hemoglobin to efficiently load and unload oxygen, enhancing its carrier function.



4. Identify key allosteric regulators like pH (Bohr effect) and BPG that modulate hemoglobin's oxygen affinity.

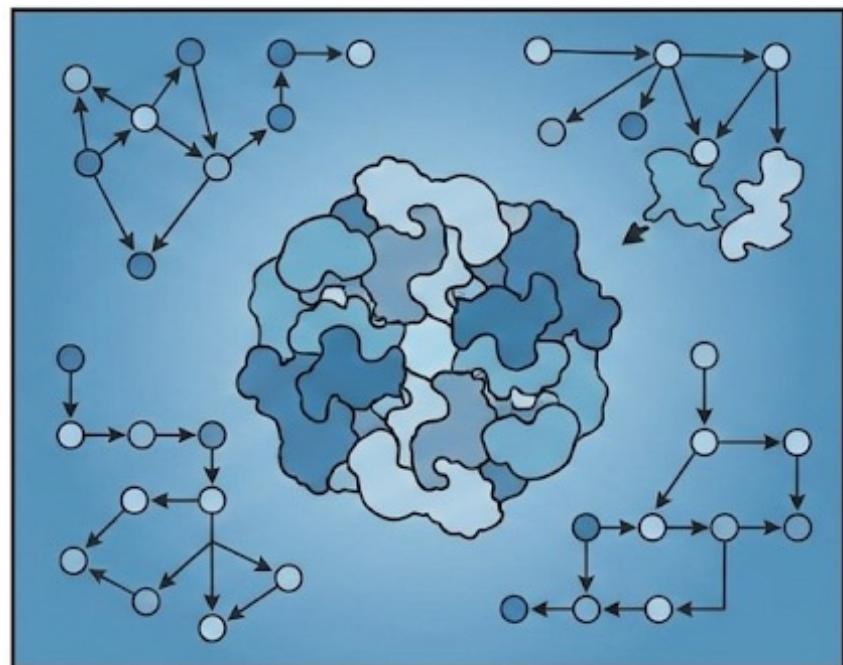
Order is what you get when chemistry gets picky.

RANDOM THERMAL MOTION



Cells are packed with bustling molecules, bustling around due to thermal motion.

ORGANIZED COMPLEXES/PATHWAYS



Order emerges with intricately organized assemblies and specific biochemical pathways.

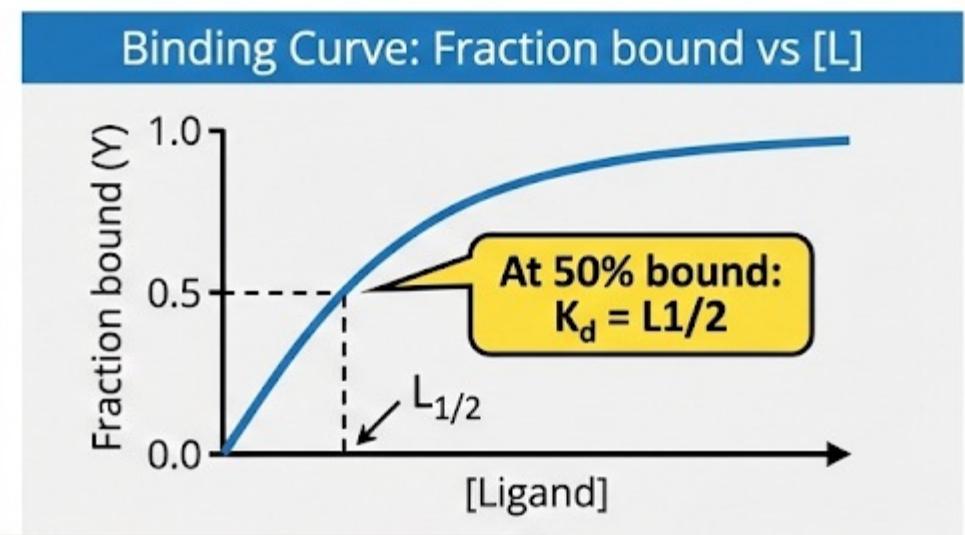
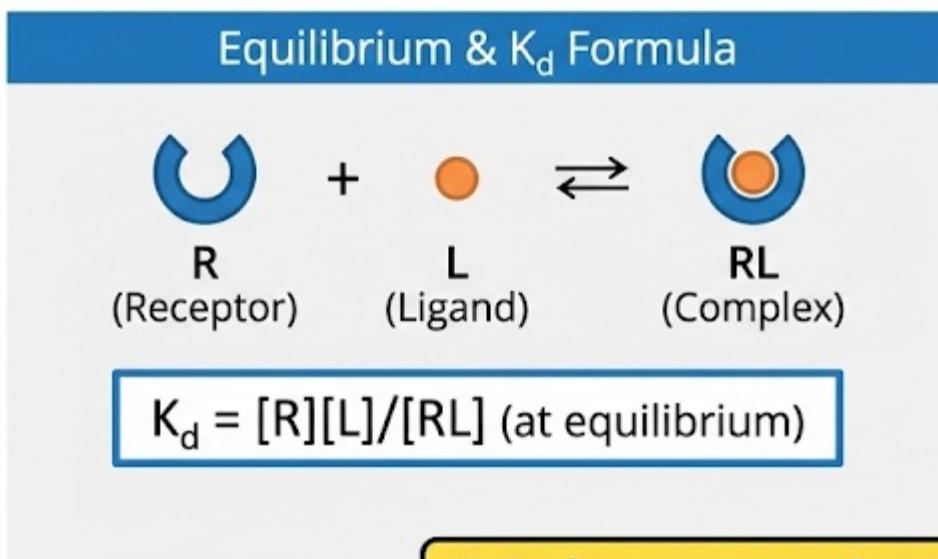
specific binding + selection

Specific molecular binding enables this organization, a consequence of chemical principles and the selective power of molecular evolution.

How do we quantify "pickiness" in a way that's useful?

Binding basics: K_d is the “half-saturation” point

- ✓ Receptor + Ligand \rightleftharpoons Complex (R + L \rightleftharpoons RL)
- ✓ Fraction bound increases as ligand concentration increases
- ✓ $K_d = [R][L]/[RL]$ (at equilibrium)
- ✓ When half the receptors are bound, $[R] = [RL]$ so $K_d = L_{1/2}$
- ✓ Smaller K_d (or $L_{1/2}$) = tighter binding (higher affinity)

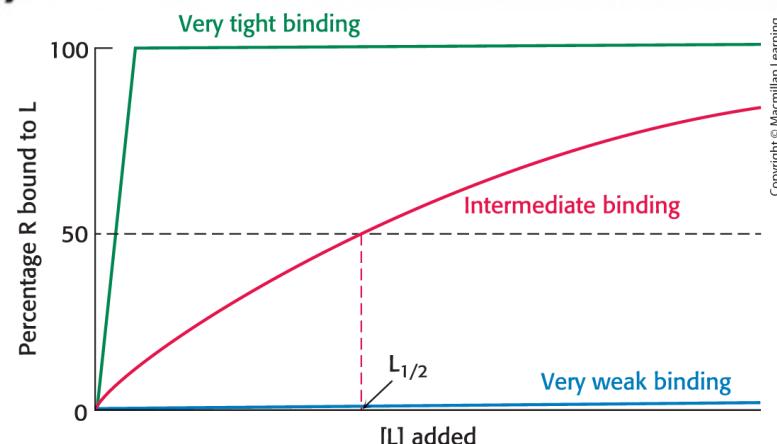


K_d is the concentration that gets you to 50% bound.

Lower K_d = higher affinity

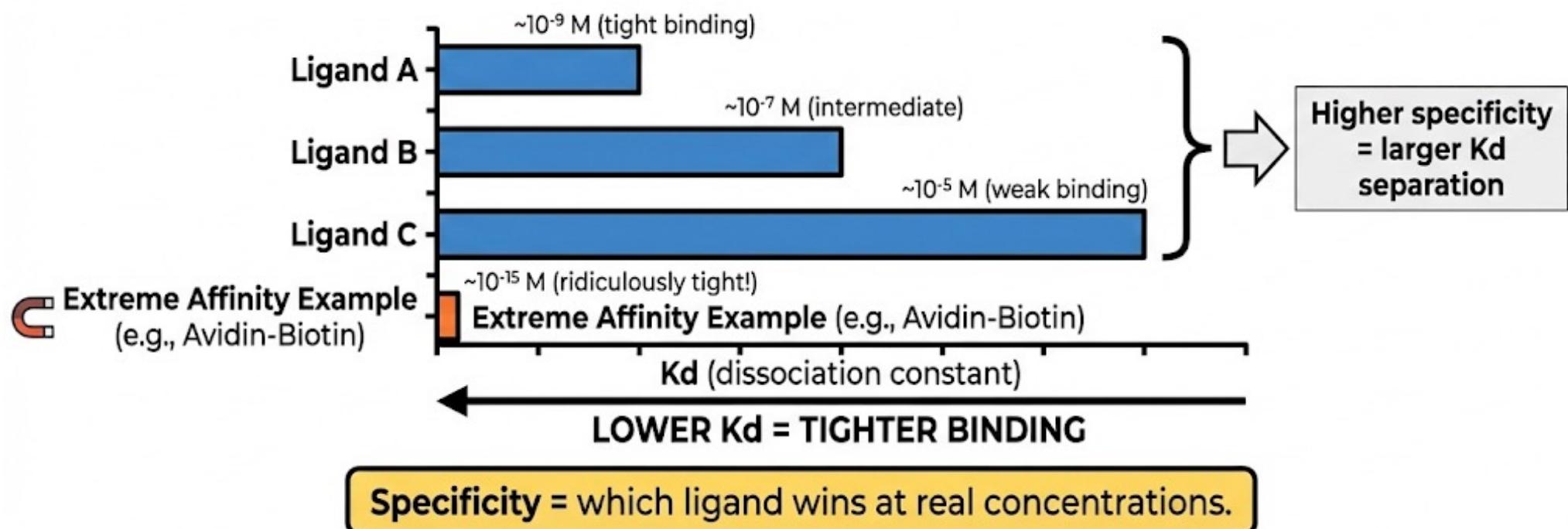
5.1 Binding is a fundamental process in biochemistry.

The ability of specific molecules to bind to one another in the face of many alternatives is referred to as **molecular recognition**.



Specificity: same protein, different K_d values

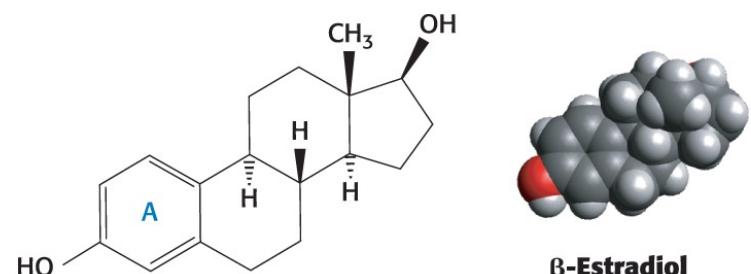
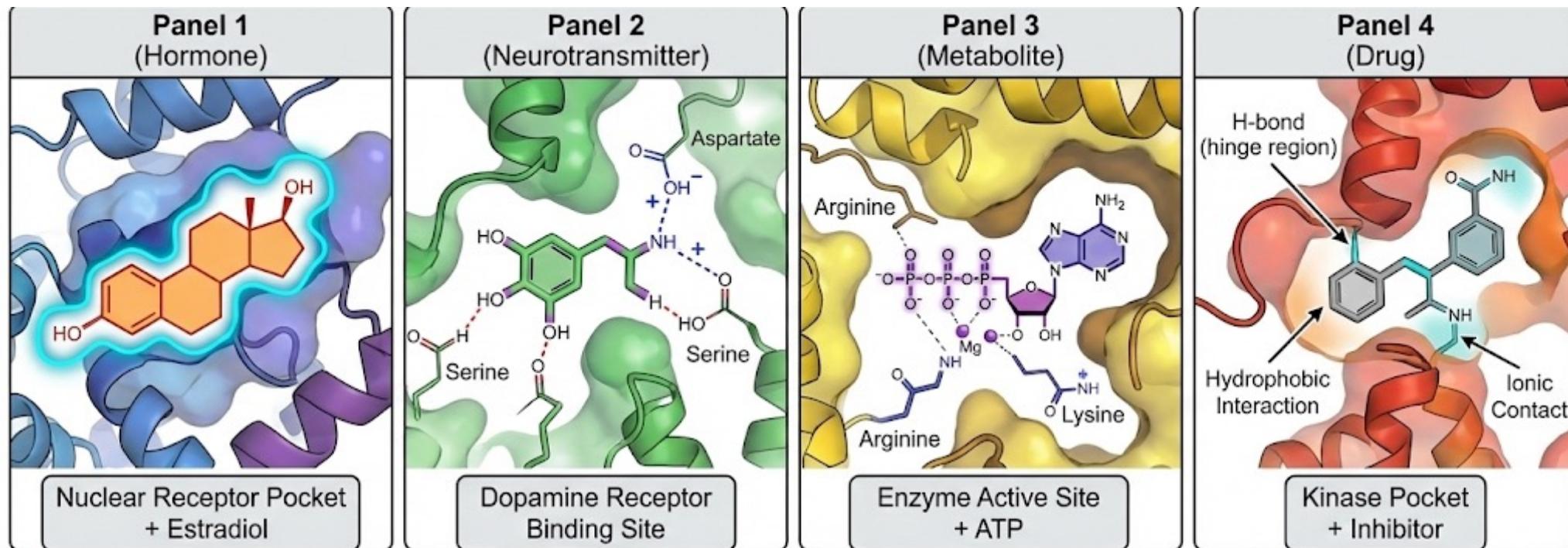
- ✓ Proteins discriminate among similar molecules via shape + charge complementarity
- ✓ Specificity is measured by comparing K_d values for competitors
- ✓ Big message: biology runs on relative affinities, not “perfect fits”
- ✓ Extreme affinity examples exist (used in real lab methods)



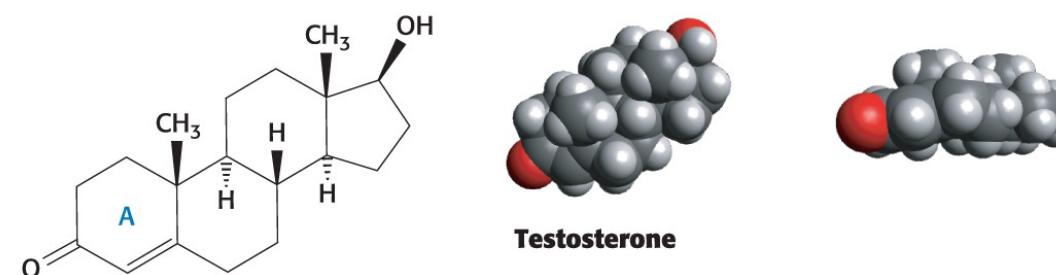
Affinity differences can be orders of magnitude.

K_d has concentration units

Protein Selective Binding—Shape + Chemistry Match



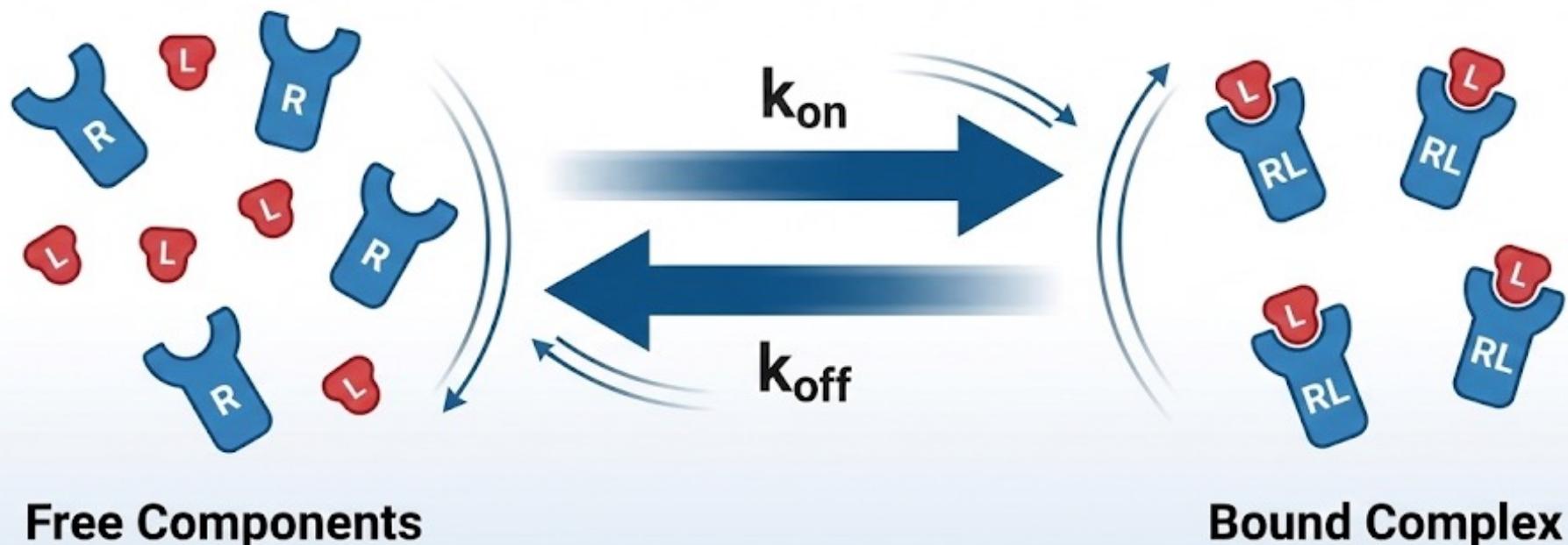
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BINDING IS A DYNAMIC PROCESS:

DYNAMIC AT EQUILIBRIUM

At **equilibrium**, the **concentrations** of R, L, and RL are **CONSTANT**.
Yet, each receptor is constantly releasing one **ligand** and **rebinding** another.



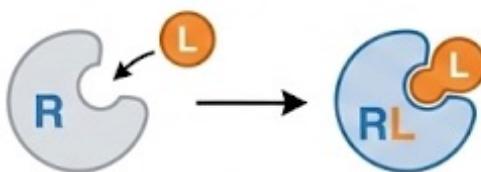
Free Components

Bound Complex

K_d : The Affinity Number (and when it equals the 50% point)

K_d is the ligand concentration (free ligand) that gives 50% occupancy under common assumptions.

ZONE 1 The binding reaction

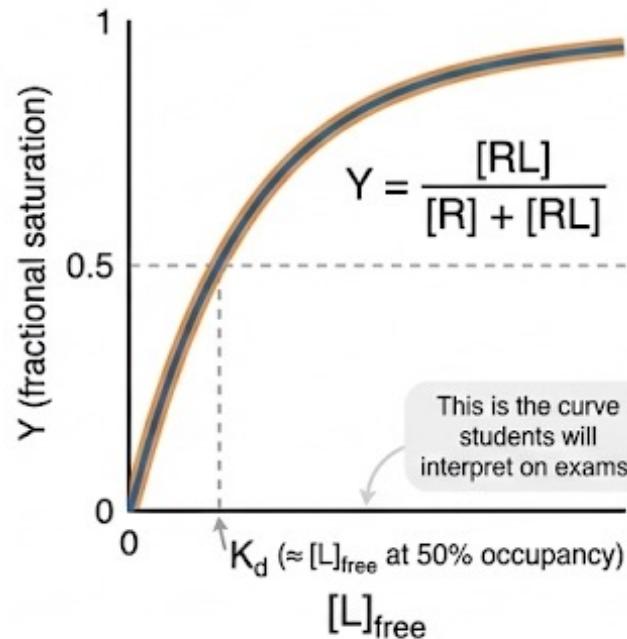


$$K_d = \frac{[R][L]}{[RL]}$$

**Smaller K_d = tighter binding
(higher affinity)**

[R] = free receptor, [L] = free ligand, [RL] = bound complex

ZONE 2 Occupancy curve + the 50% point



**Exam-use
translation**

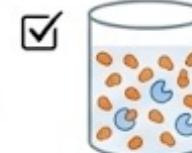
Lower K_d → higher affinity

Right-shifted curve →
weaker binding (higher K_d)

At $Y = 0.5$, read x-axis: $\sim K_d$
(if assumptions hold)

ZONE 3 When does K_d equal the 50% point?

The shortcut is true when...



- 1. Ligand is not depleted
 $[L]_{total} \approx [L]_{free}$
(R is small relative to L)



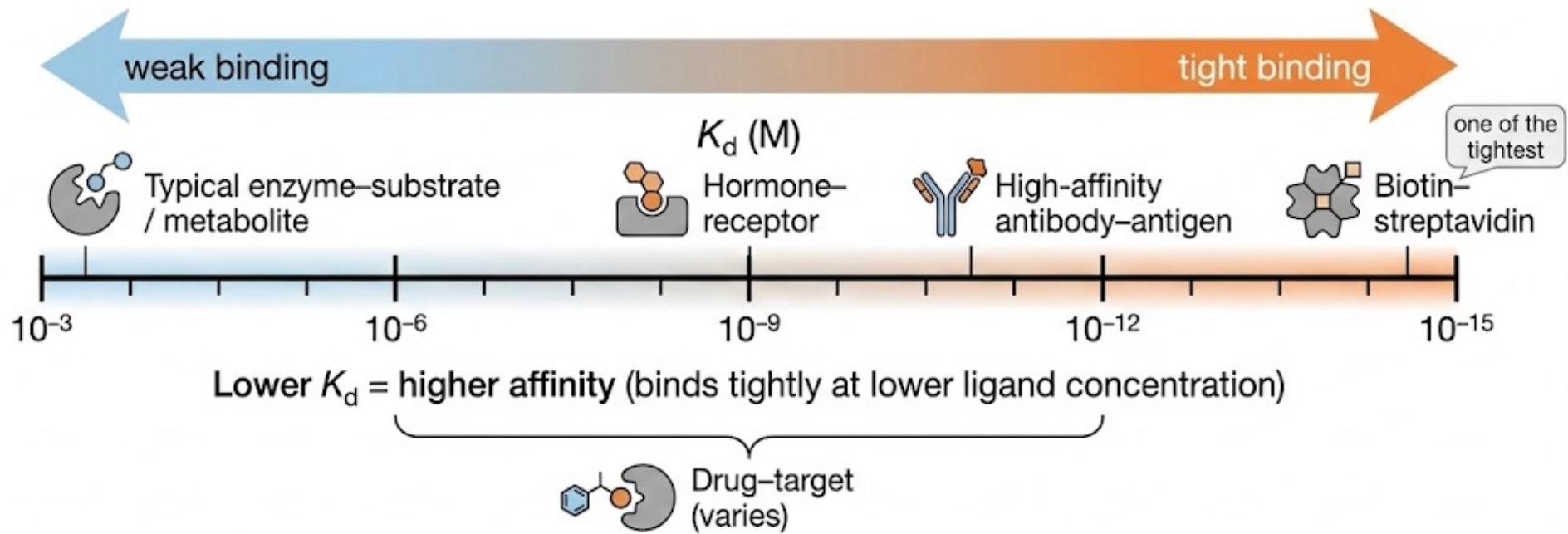
- 2. Single binding site,
simple equilibrium
No cooperativity / no
multiple sites



When it can fail

- High receptor concentration depletes ligand
- Multiple sites / cooperativity shifts the relationship

Affinity in One Glance: K_d Ranges (What ‘tight binding’ means)



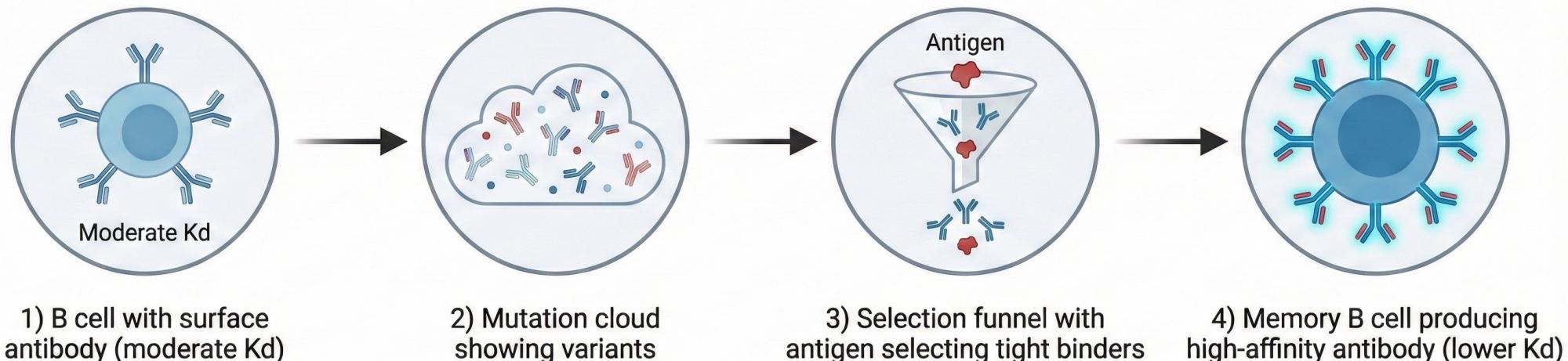
Exam-use translation

✓ Compare K_d values:
smaller wins

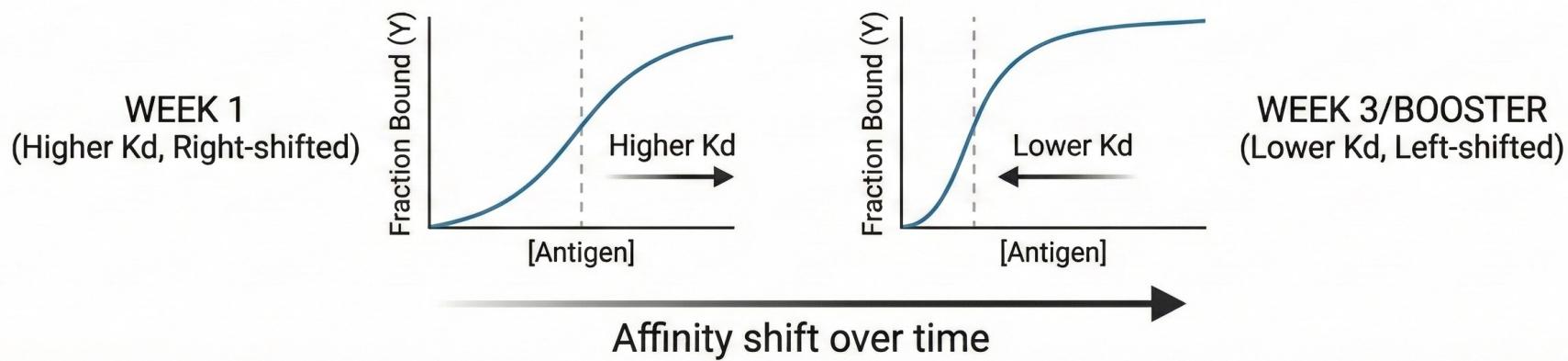
✓ A 10 \times lower K_d means
much **tighter binding**

✓ You rarely calculate K_d ;
you **interpret it**

How Vaccines Generate High-Affinity Antibodies (Affinity Maturation)



KEY MESSAGE: “Selection favors lower Kd (tighter binding)”

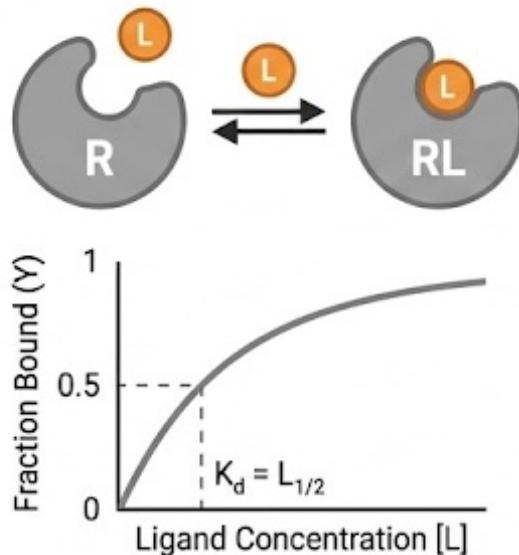


True or False

The dissociation constant of the ligand-binding reaction is equal to the concentration of L at which half of the receptor is bound to L (and half is free).

FROM SIMPLE BINDING TO COOPERATIVITY: THE HEMOGLOBIN STORY

WHAT WE JUST LEARNED



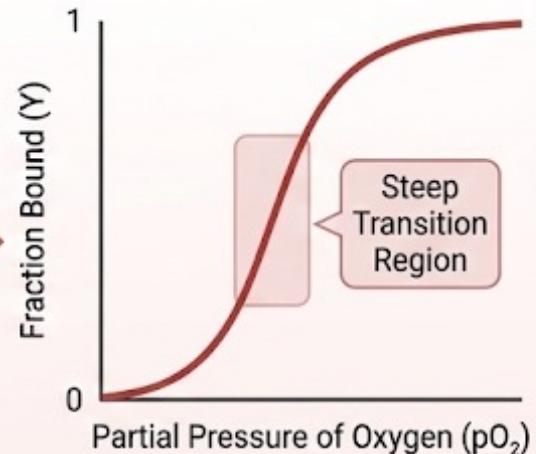
Single binding site
 $K_d = L_{1/2}$
One protein, one ligand

THE HEMOGLOBIN TWIST



Multiple binding sites
Sites communicate
Binding changes conformation

THE RESULT



Sigmoidal curve
Better physiological control
Sensitive to small pO_2 changes

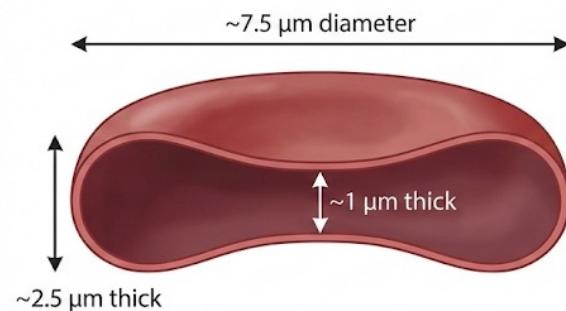
Cooperativity = binding at one site changes affinity at other sites

How does hemoglobin load O₂ in lungs but unload in tissues?

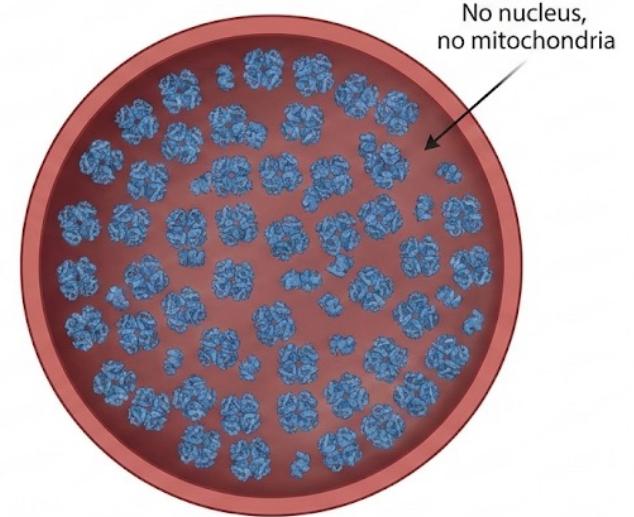


- Red blood cell:
- We have 5 million RBCs per microliter of blood
- -Biconcave---lacks some cytoskeletal elements
- Lacks several key organelles like nuclei

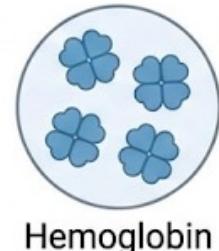
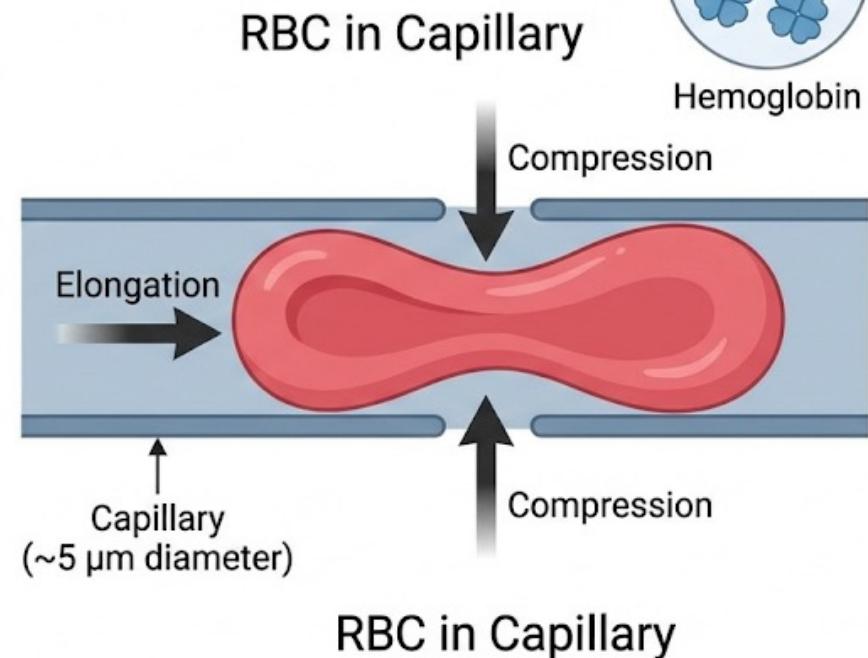
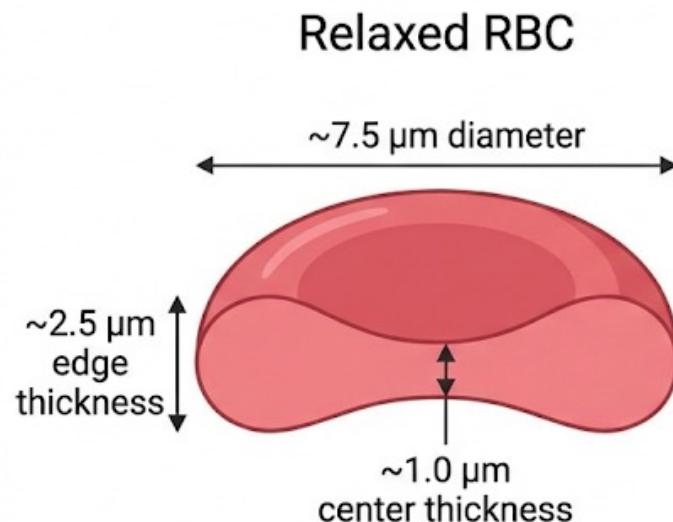
SIDE VIEW & MEASUREMENTS



TOP-VIEW CROSS-SECTION & CONTENTS



RBC Deformation in Capillaries



Red blood cell:

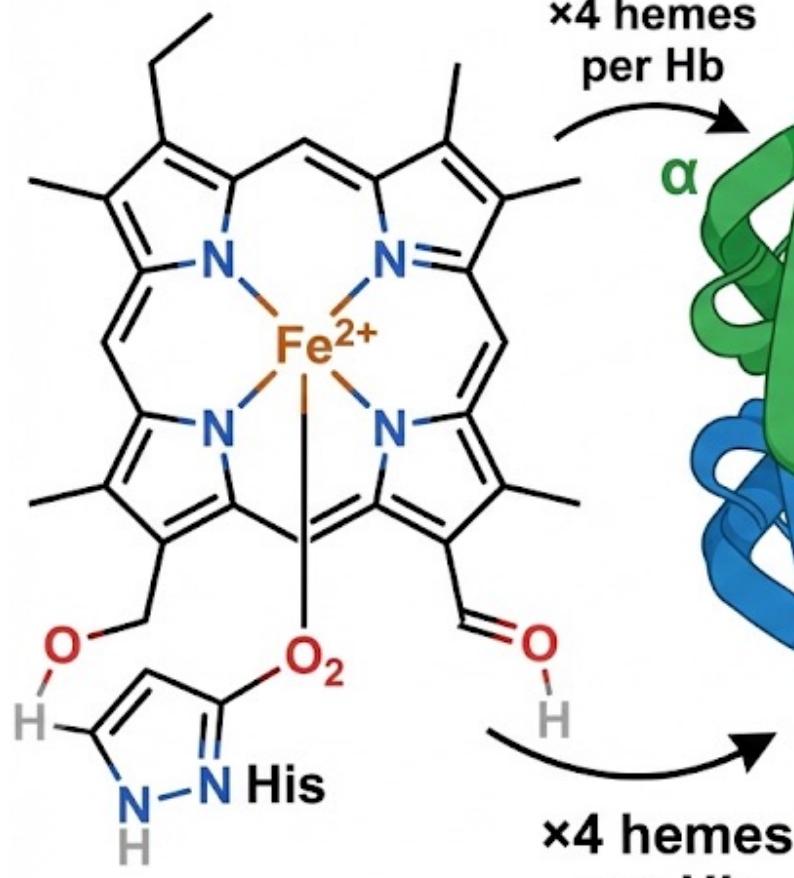
- Super flexible
- no nucleus
- can squeeze through capillaries
 - the squeezing is important to maximize gas exchange

RBC lifespan (~120 days)

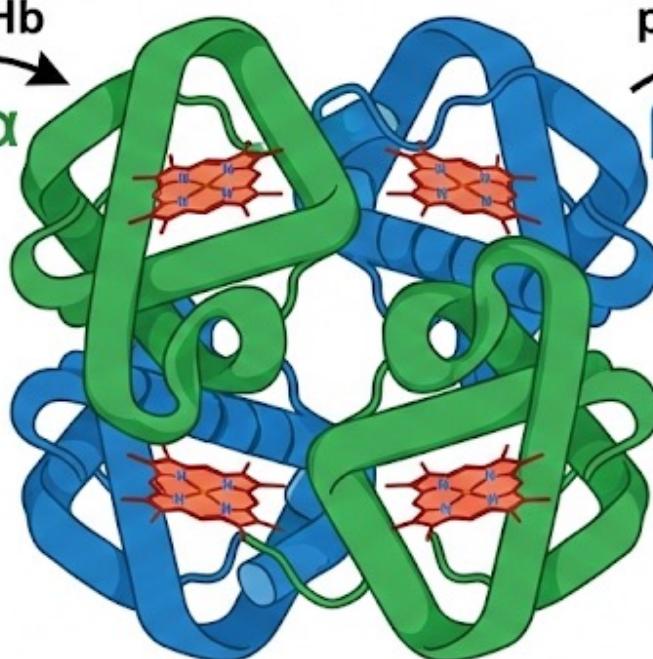
1 red blood cell contains about 300 million molecules of hemoglobin.

Each RBC can transport about 1.2 billion O₂ molecules.

Heme Group



Hemoglobin Tetramer



$\times 4$ hemes
per Hb

α

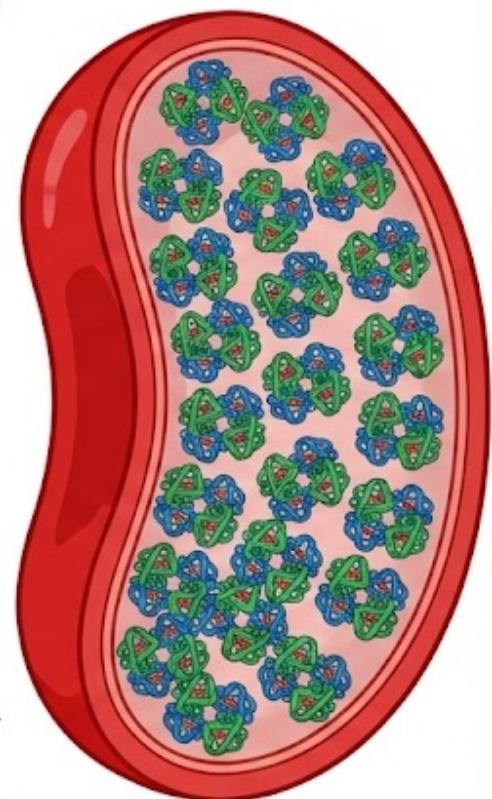
$\times 4$ hemes
per Hb

$\times 300\text{M Hb}$
per RBC

β

$\times 300\text{M Hb}$
per RBC

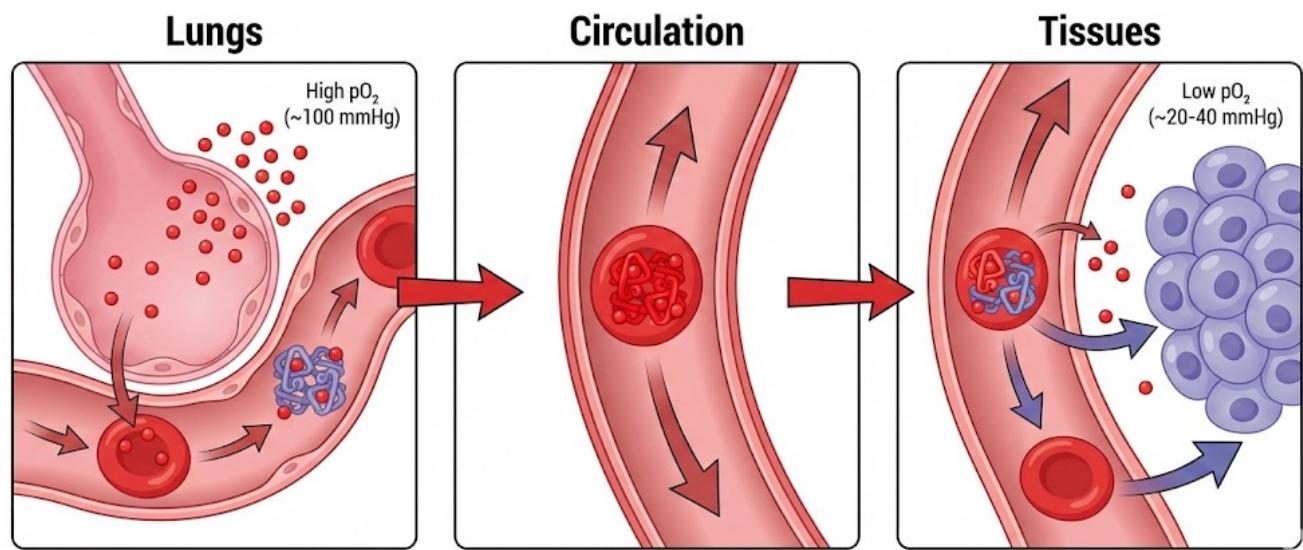
Red Blood Cell (RBC)



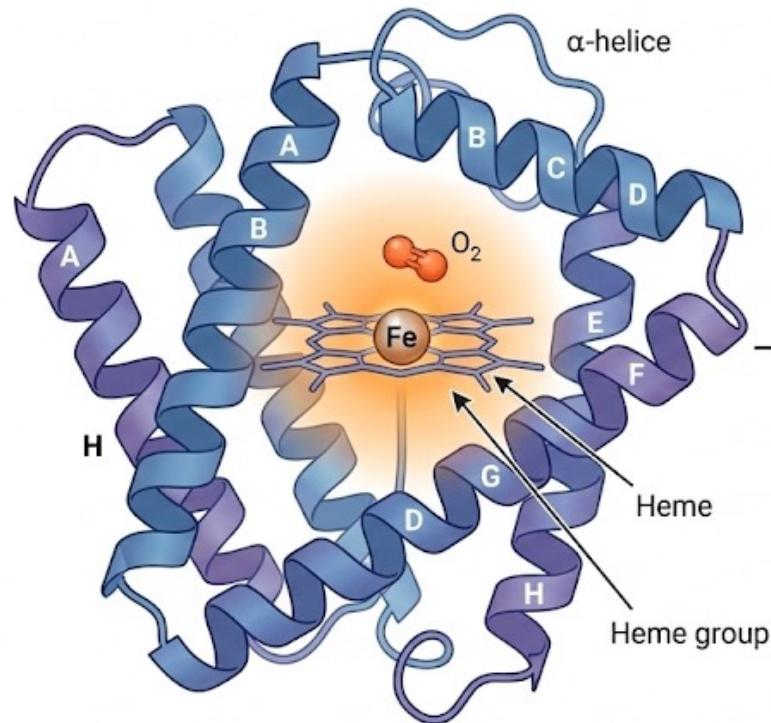
- Learning objective 7:
Explain how allosteric properties contribute to hemoglobin function.

- Hemoglobin is a red blood cell protein that carries oxygen from the lungs to the tissues.
- Hemoglobin is an allosteric protein that displays cooperativity in oxygen binding and release.
- Myoglobin binds oxygen in muscle cells. The binding of oxygen by myoglobin is not cooperative.
- Oxygen binding is measured as a function of the partial pressure of oxygen (pO_2).

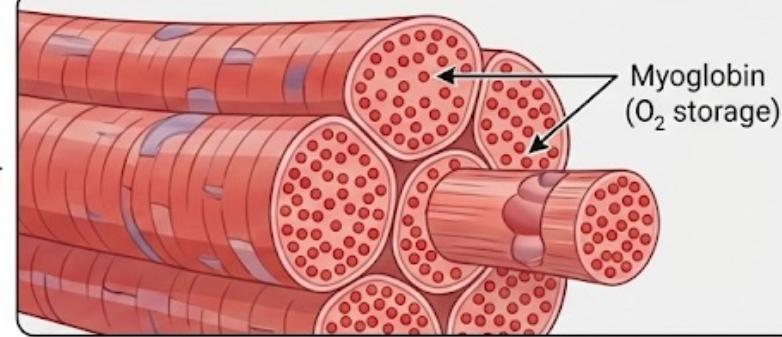
HEMOGLOBIN'S OXYGEN JOURNEY



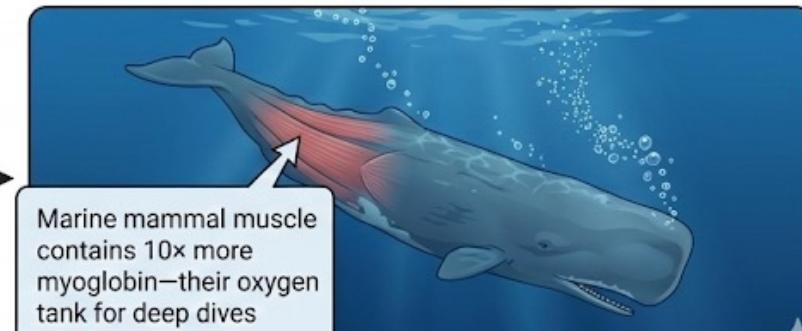
MYOGLOBIN STRUCTURE



OXYGEN RESERVE IN MUSCLE



MARINE MAMMAL ADAPTATION

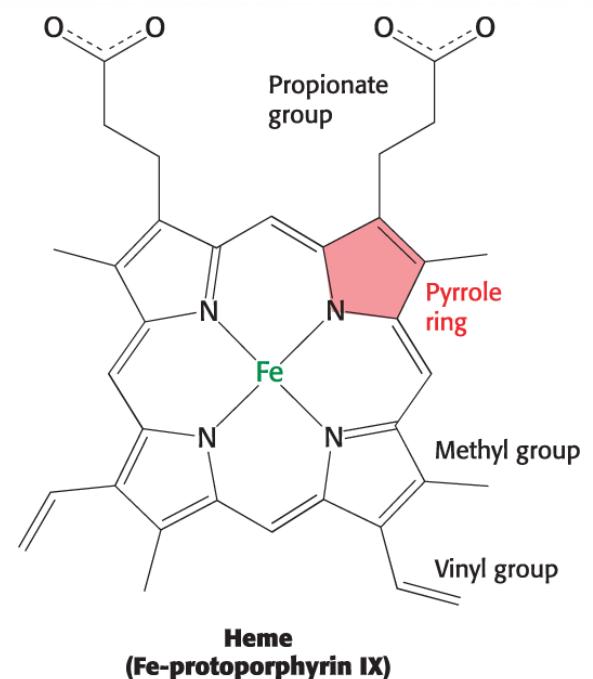
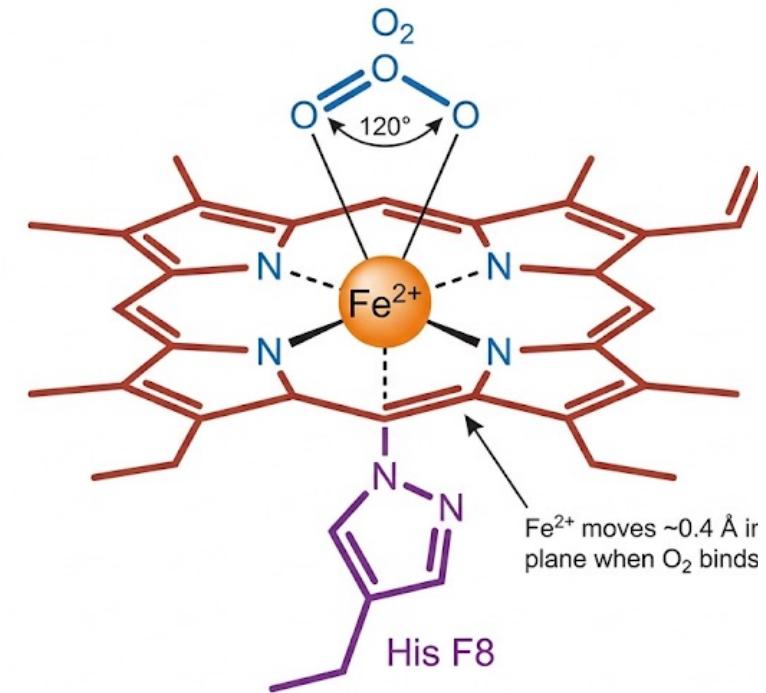


Section 5.2 Myoglobin and Hemoglobin Bind Oxygen in Heme Groups

- Myoglobin is a single polypeptide chain consisting mainly of α helices arranged to form a globular structure. This is a simple reference case.
- Myoglobin, like hemoglobin, binds oxygen at a heme, a bound prosthetic group.

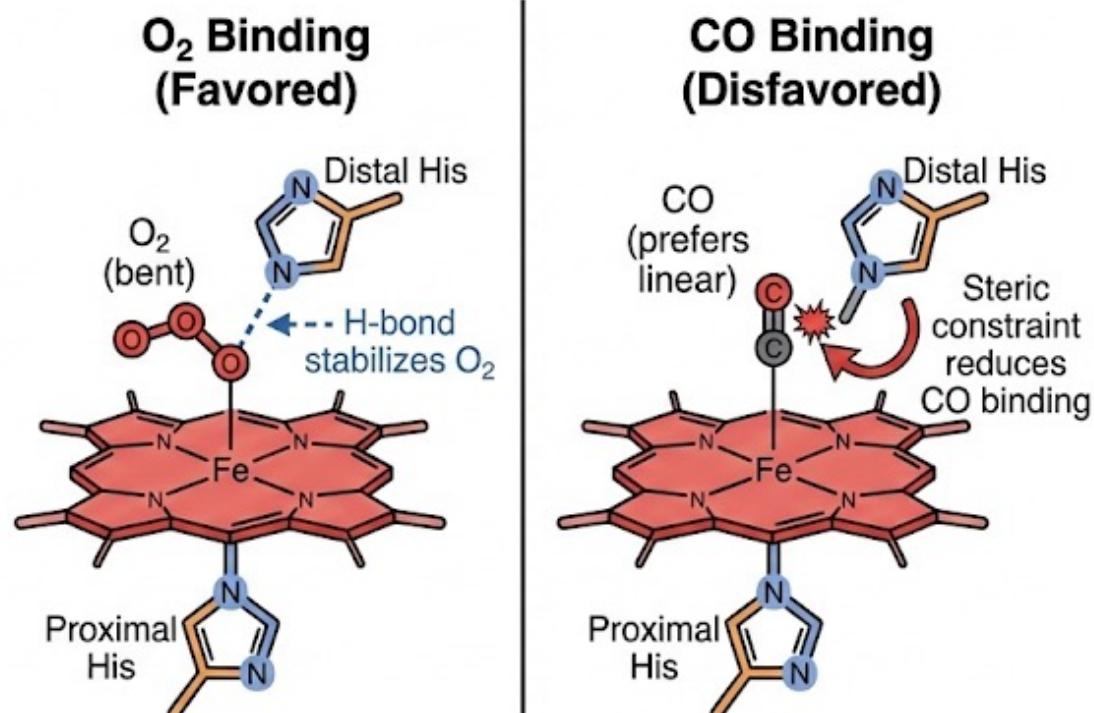
Section 5.2 Myoglobin and Hemoglobin Bind Oxygen in Heme Groups (2/3)

- The heme group consists of an organic component called protoporphyrin and a central iron ion in the ferrous (Fe^{2+}) form.
- The iron lies in the middle of the protoporphyrin bound to four nitrogens.
- Iron can form two additional bonds, called the fifth and sixth coordination sites.
- The fifth coordination site is occupied by an imidazole ring of a histidine called the proximal histidine.
- The sixth coordination site binds oxygen.
- Upon oxygen binding, the iron moves into the plane of the protoporphyrin ring.



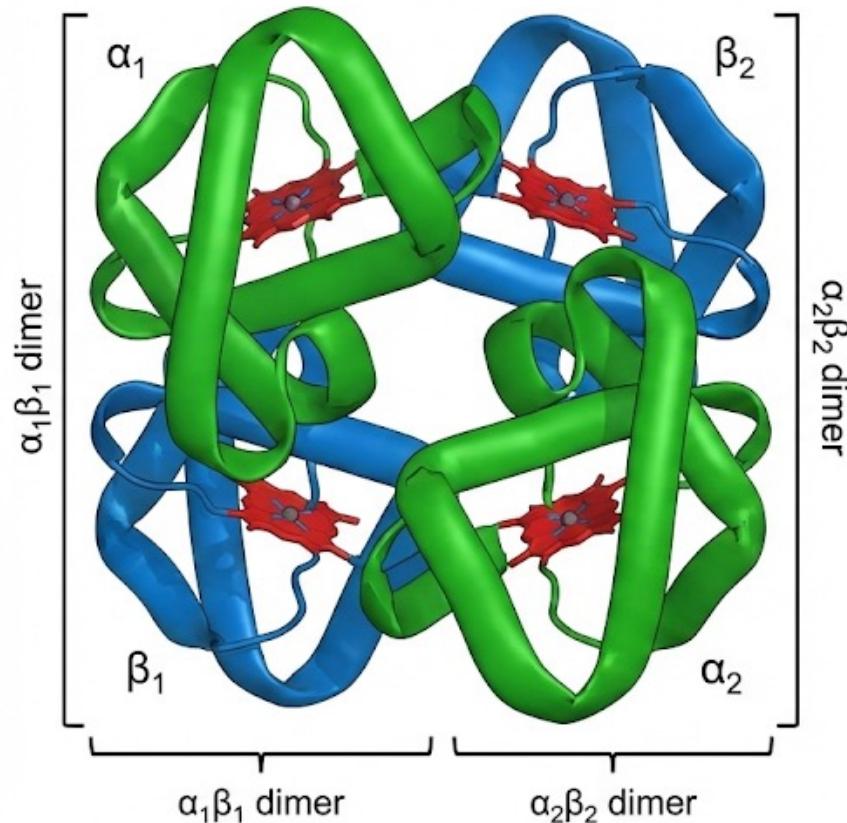
Distal histidine: the heme “bouncer” that favors O₂ over CO

- Heme iron can bind O₂ at the 6th coordination site, but CO can compete
- Distal His partially blocks straight-line binding geometries
- Distal His provides a stabilizing H-bond that favors the bent O₂ binding mode
- Net effect: boosts O₂ binding and reduces CO advantage
- Helps protect heme from undesirable chemistry (must remain functional Fe²⁺)

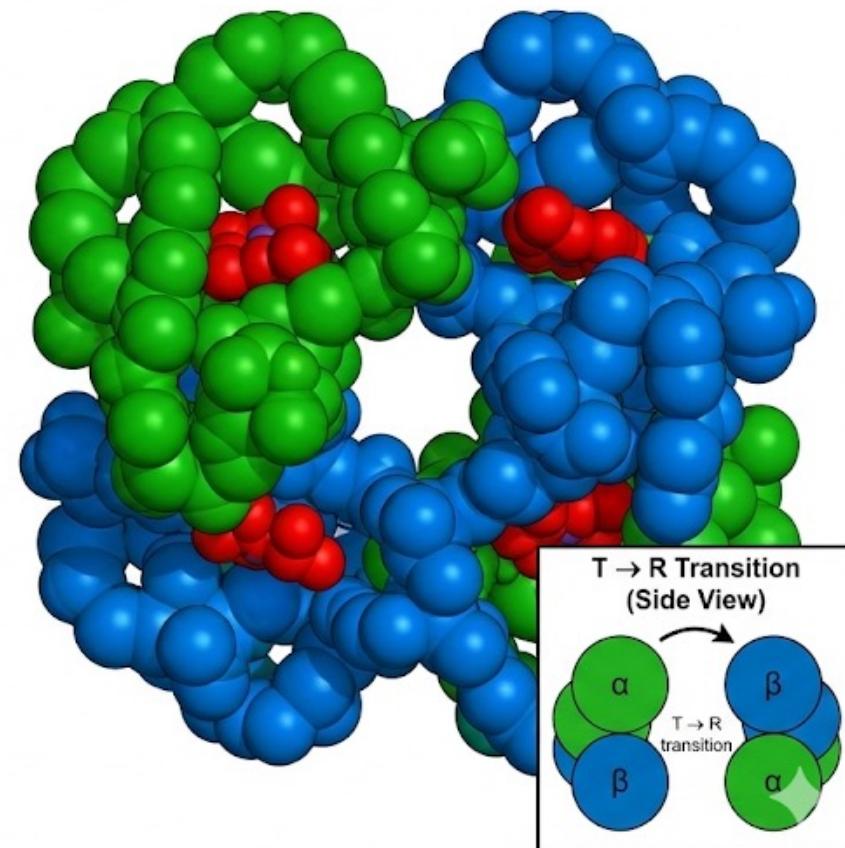


Protein environment, not heme alone, determines “safe” ligand binding.

Panel A: $\alpha_2\beta_2$ Tetramer (Top View, Ribbon)



Panel B: Space-Filling Model with Central Cavity



Section 5.3 Hemoglobin Binds Oxygen Cooperatively

- Hemoglobin is a tetramer consisting of two α subunits and two β subunits. Each subunit has a bound heme.
- The quaternary structure is best described as a pair of identical $\alpha\beta$ dimers ($\alpha_1\beta_1$ and $\alpha_2\beta_2$).
- In deoxyhemoglobin, which corresponds to the T state of allosteric enzymes, the $\alpha\beta$ dimers are linked by an extensive interface.

The Mechanical Linkage: From Iron to Quaternary Structure

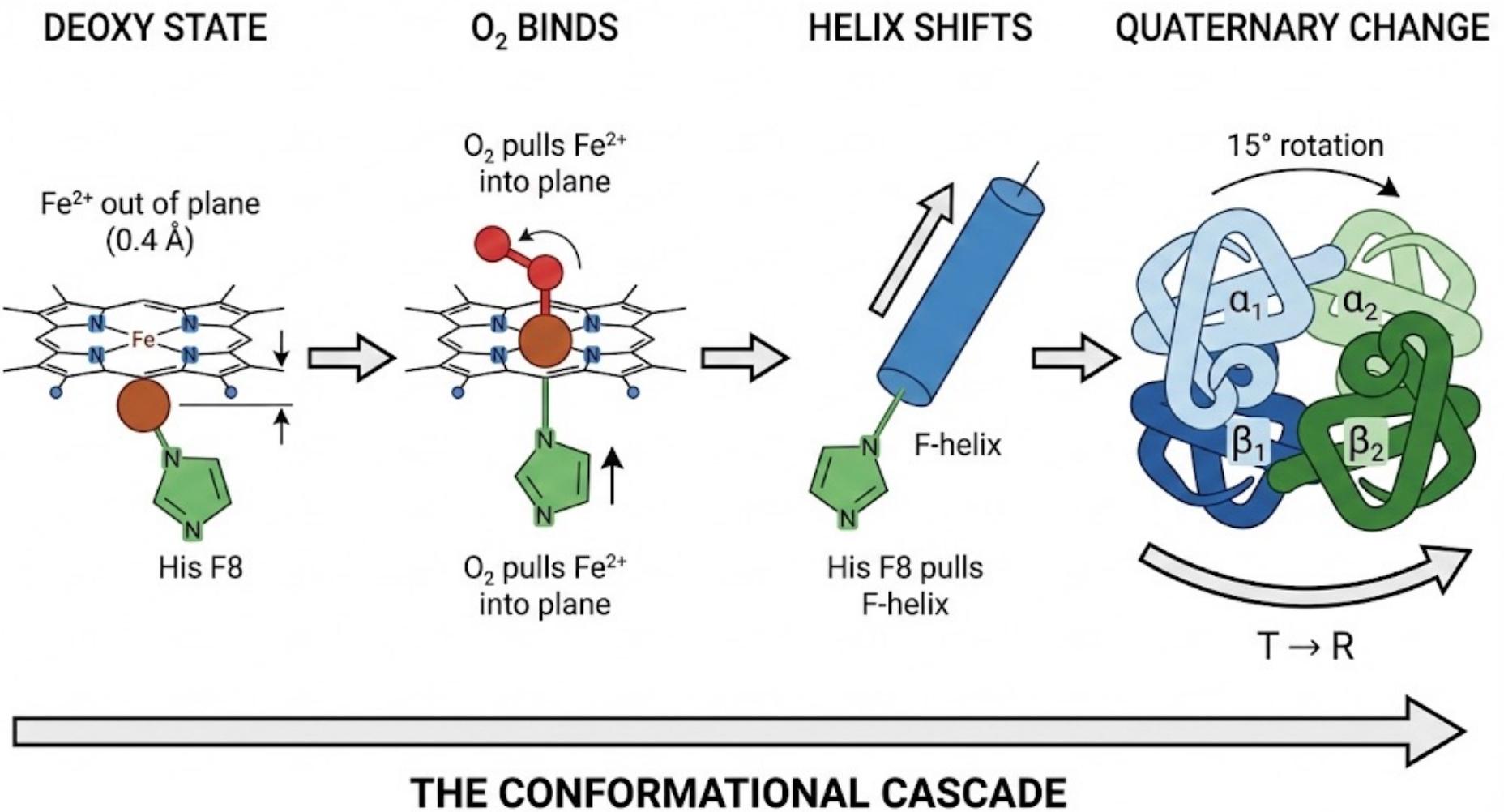
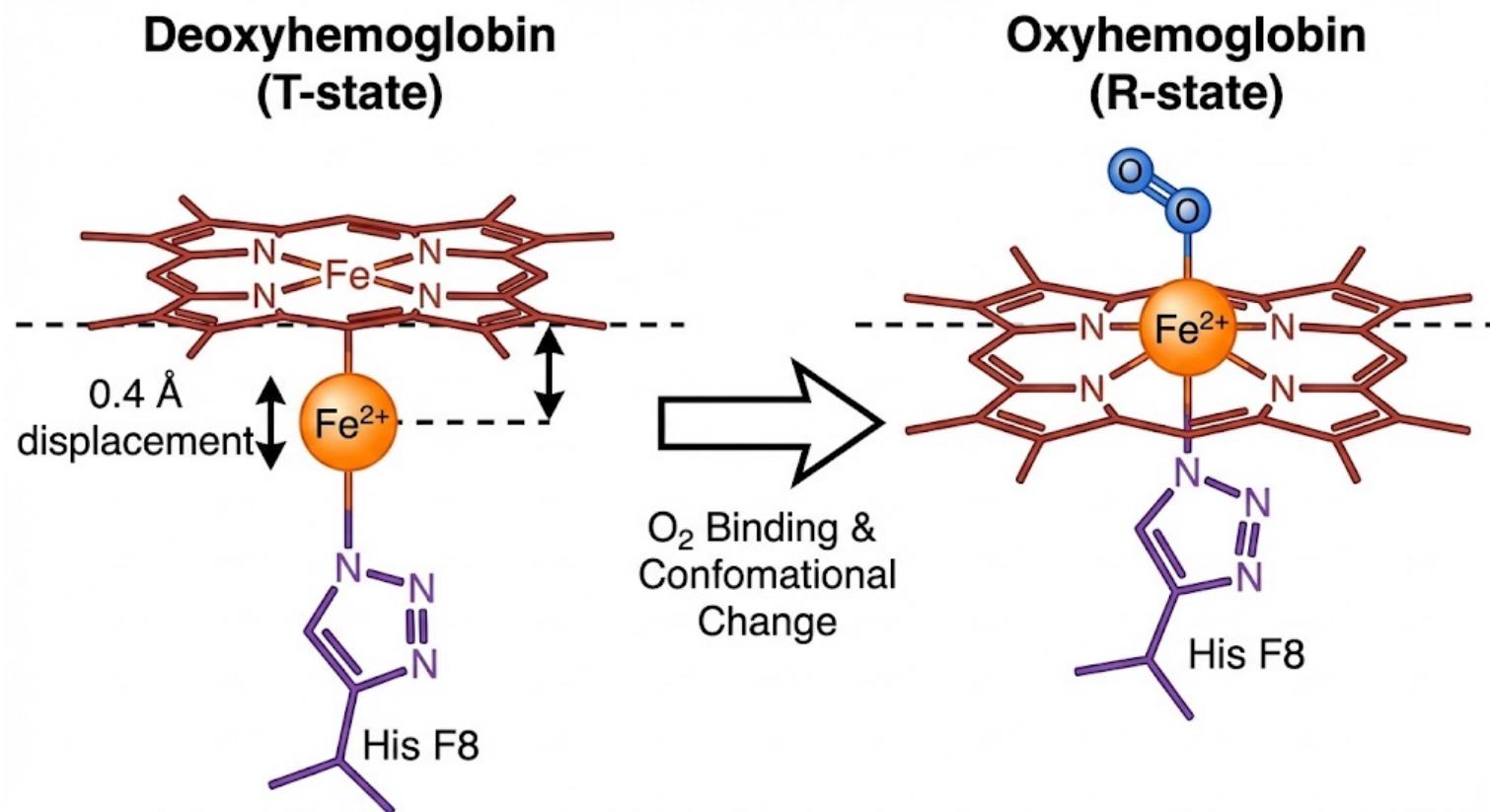
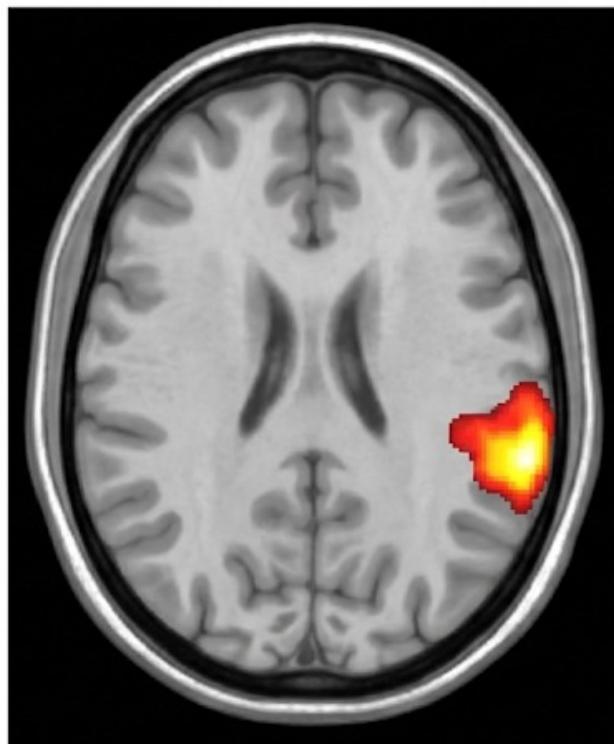


Diagram of Oxygen Binding Changing the Position of the Iron Ion

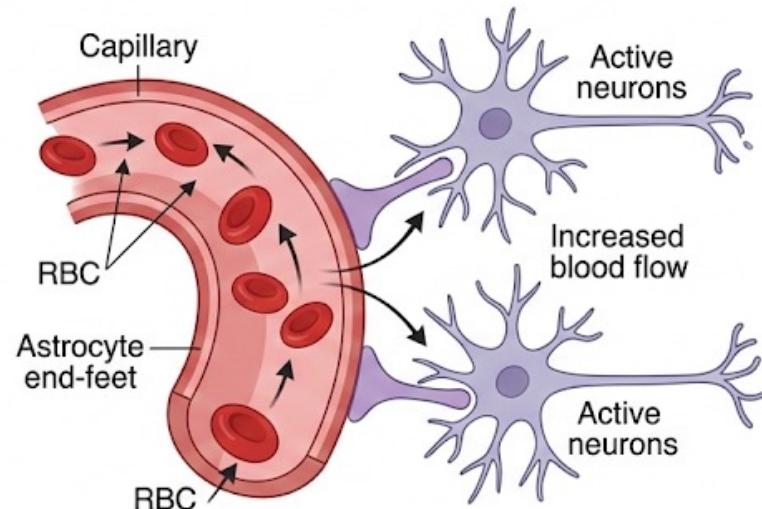
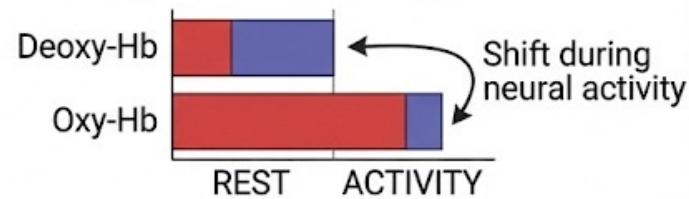


This will be the state of hemoglobin in tissues to offload O₂

This will be the state of hemoglobin in lungs to onload O₂

A**fMRI BOLD RESPONSE**

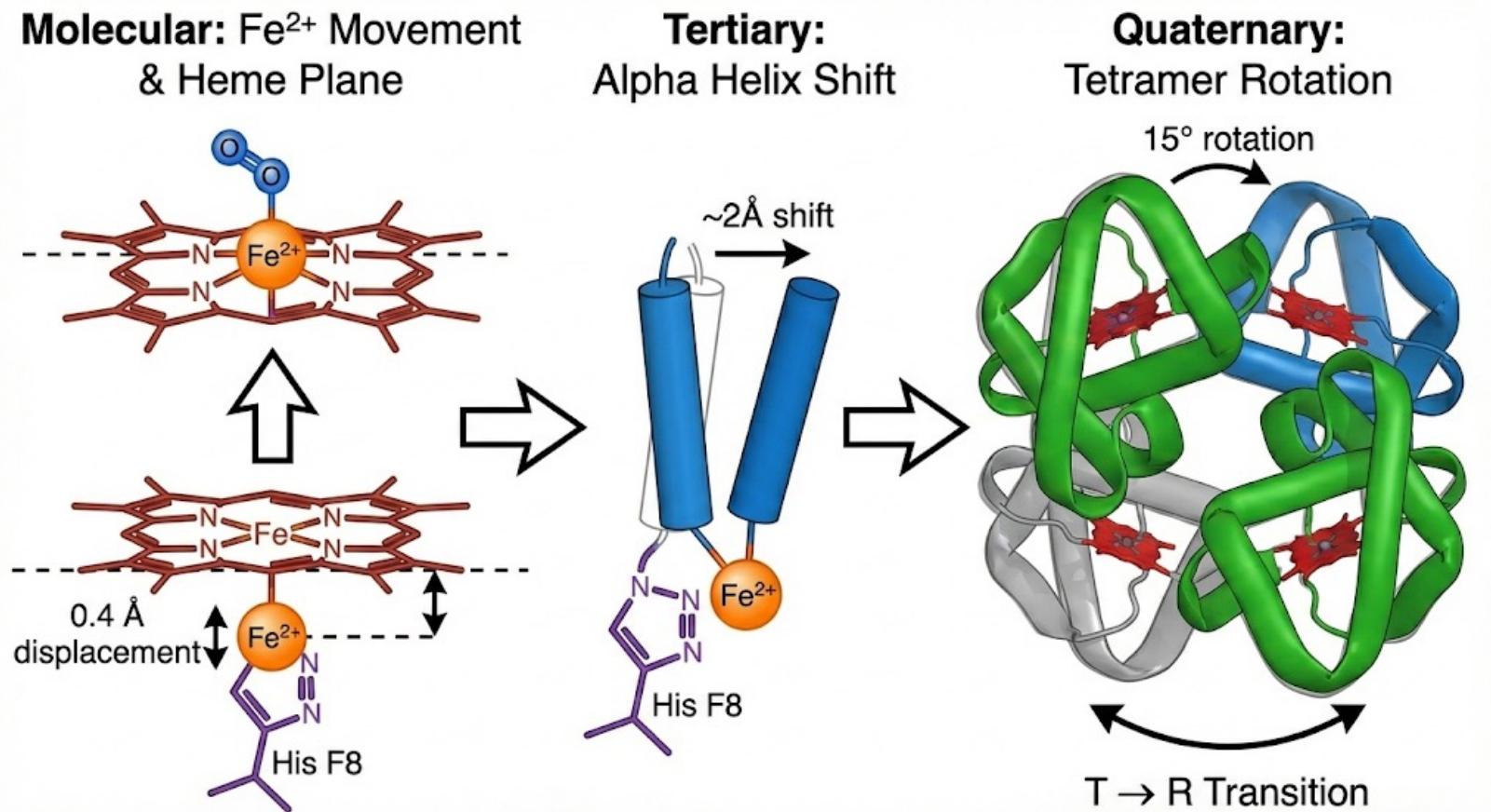
Brain response to sensory stimulus

B**NEUROVASCULAR COUPLING DIAGRAM****Oxy-Hb/Deoxy-Hb Ratio Shifts**

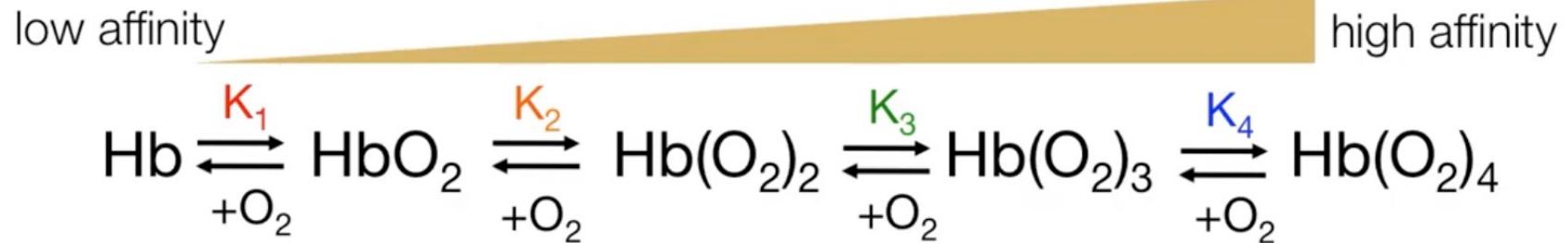
- **Clinical Insight: Functional Magnetic Resonance Imaging Reveals Regions of the Brain Processing Sensory Information**
- The magnetic properties of the heme iron change when it moves into the plane of the protoporphyrin ring.
- Functional magnetic resonance imaging (fMRI) can distinguish the relative amounts of oxy- and deoxyhemoglobin.
- Functional magnetic resonance can be used to monitor activity in specific regions of the brain by measuring the increase in oxyhemoglobin.

Section 5.3 Hemoglobin Binds Oxygen Cooperatively

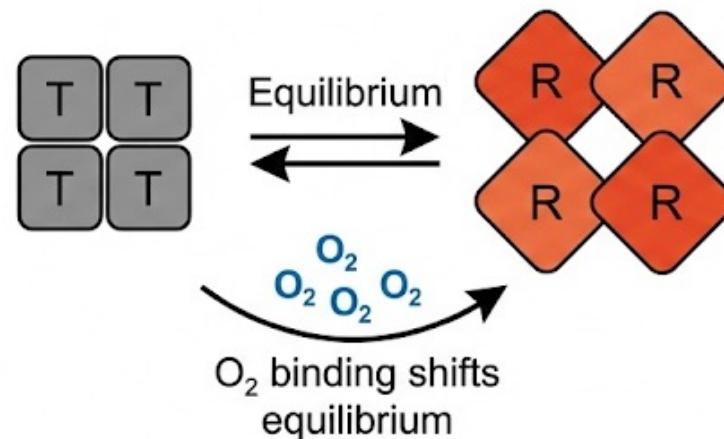
- The transition from deoxyhemoglobin (T state) to oxyhemoglobin (R state) occurs upon oxygen binding.
- The iron ion moves into the plane of the heme when oxygen binds. The proximal histidine, which is a component of an α helix, moves with the iron.
- The resulting structural change is communicated to the other subunits so that the two $\alpha\beta$ dimers rotate with respect to each other, resulting in the formation of the R state.



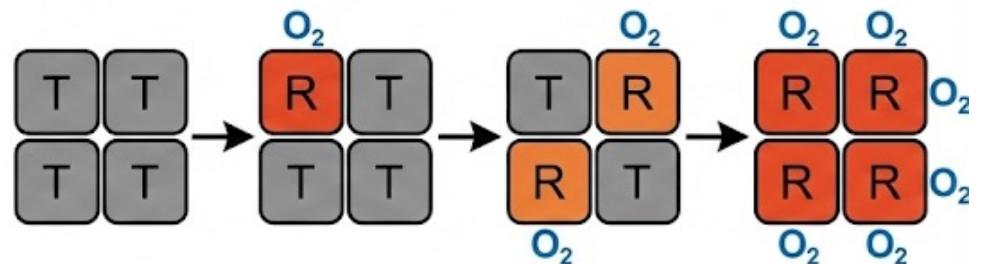
Concerted vs Sequential Models (MWC vs KNF)



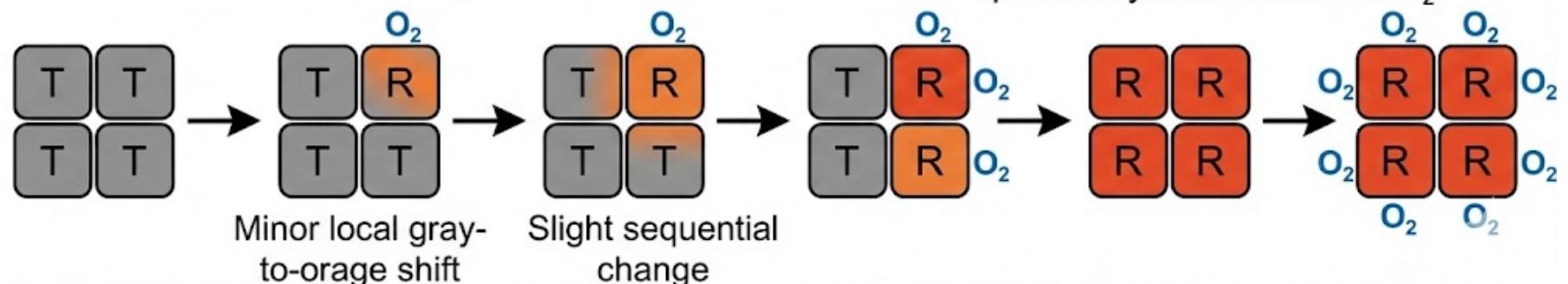
A Concerted Model (MWC)



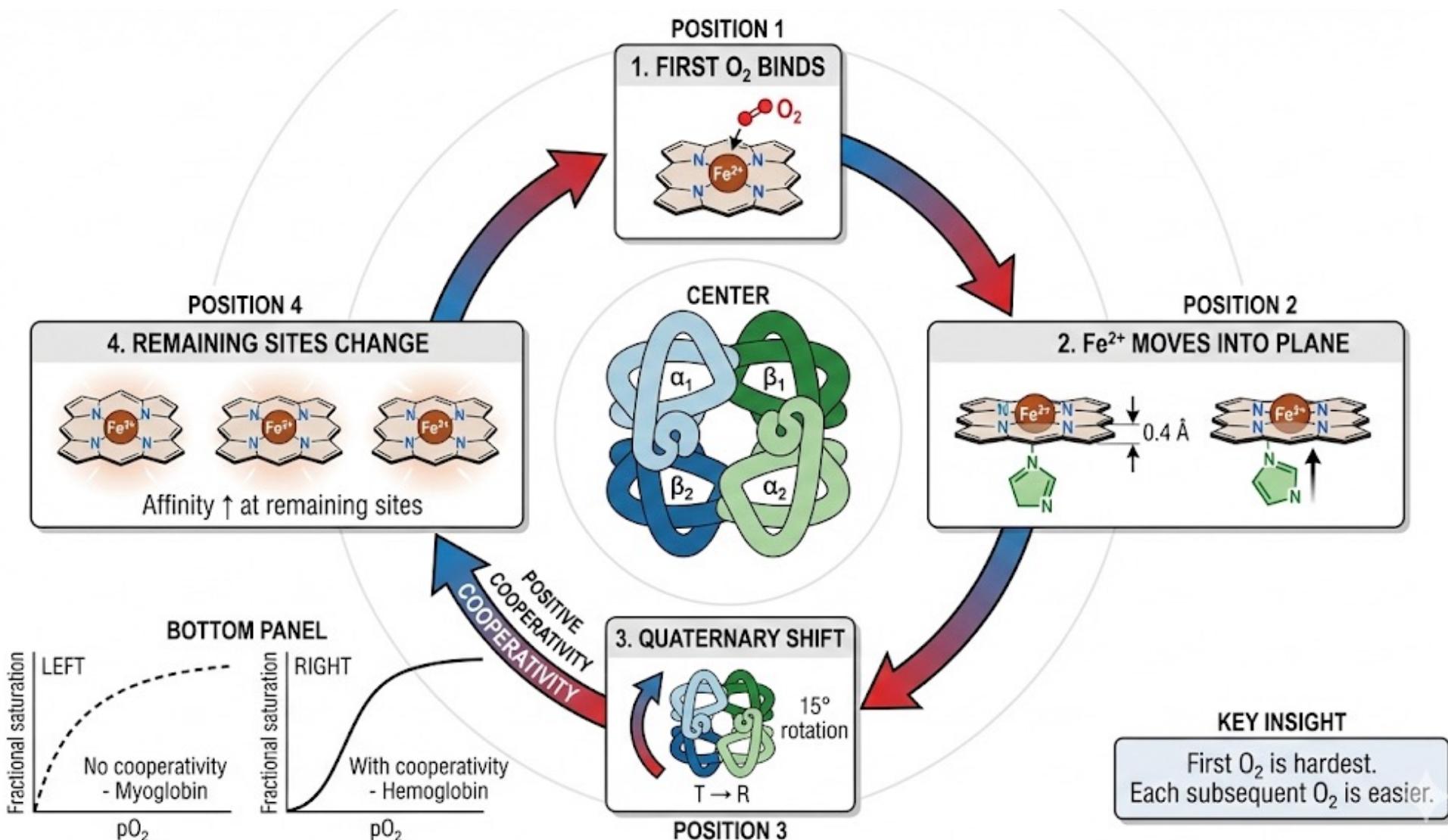
B Sequential Model (KNF)



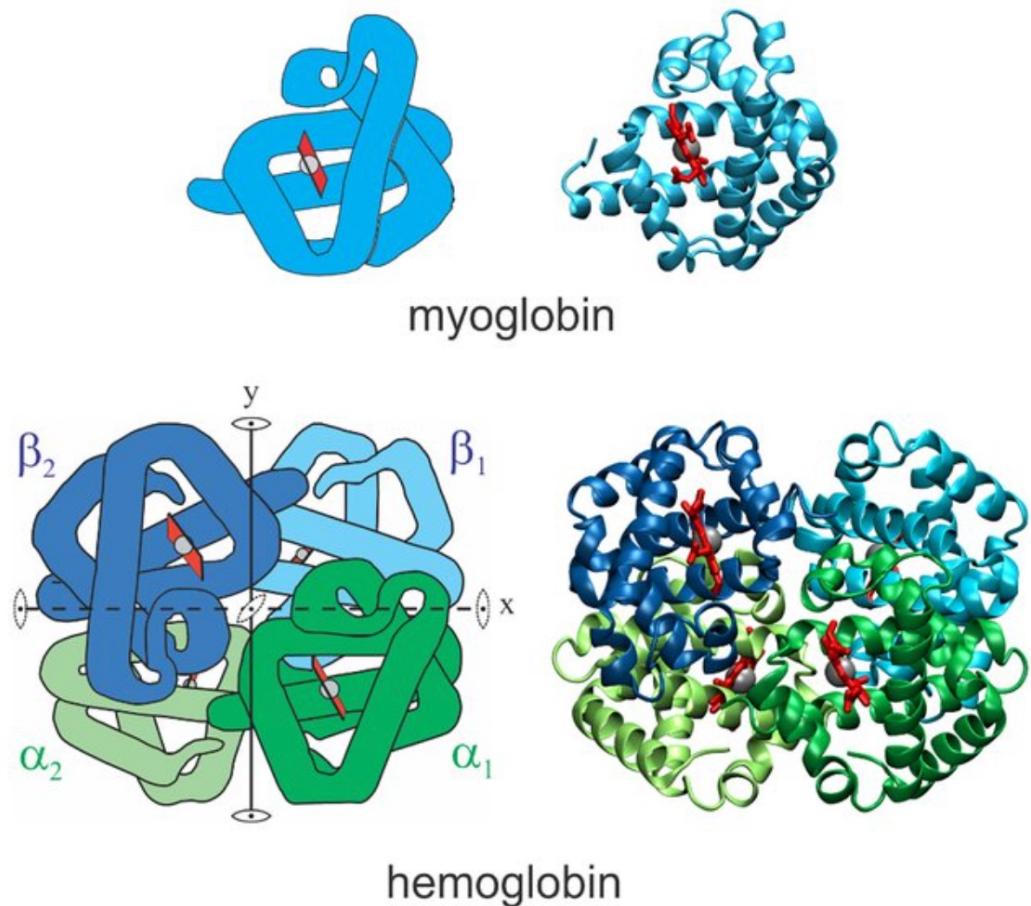
C Reality (Hybrid Model)



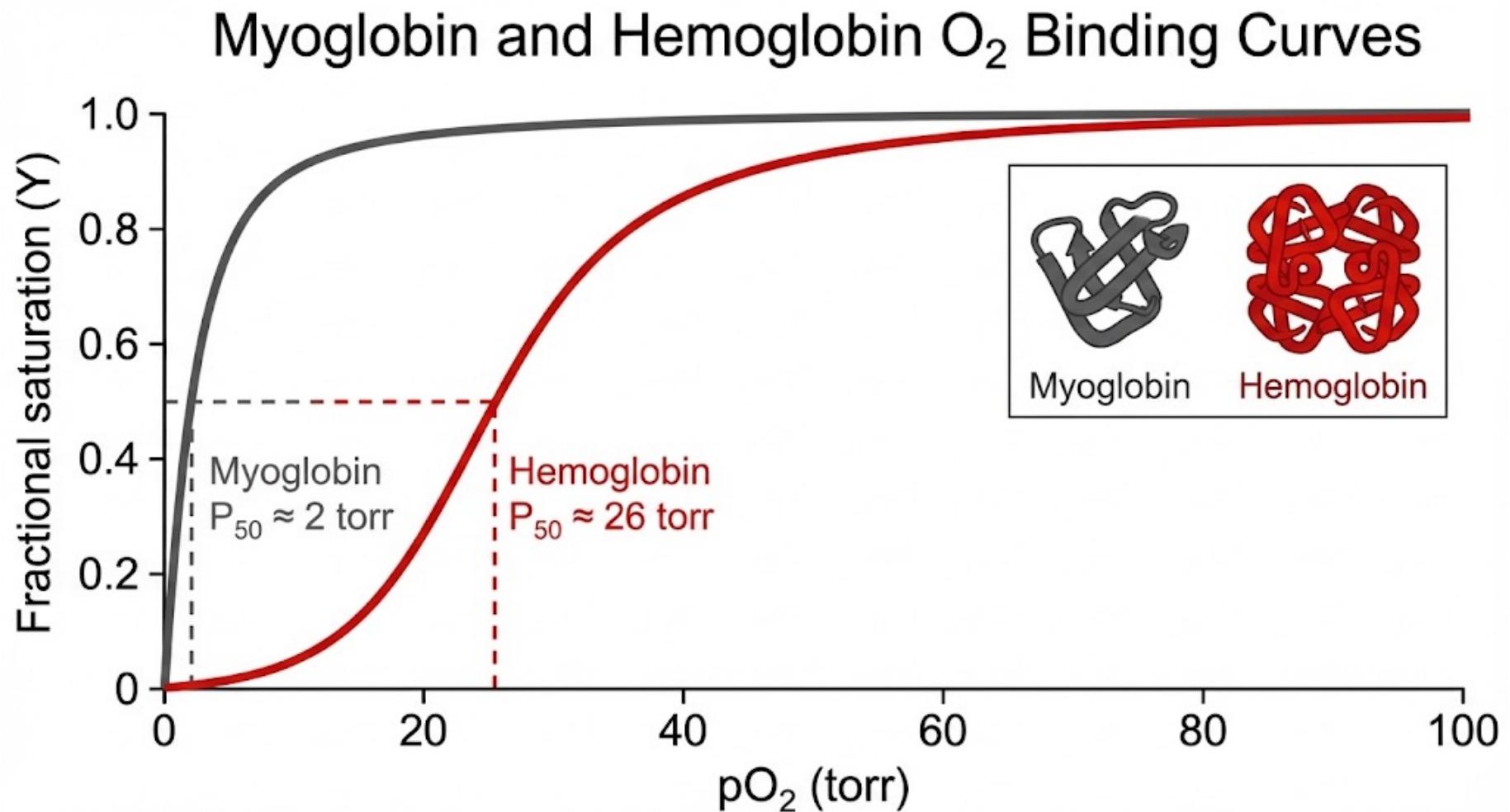
Cooperativity: The Complete Picture



What do we
imagine the O₂
binding curves
would look like
between
myoglobin and
hemoglobin?



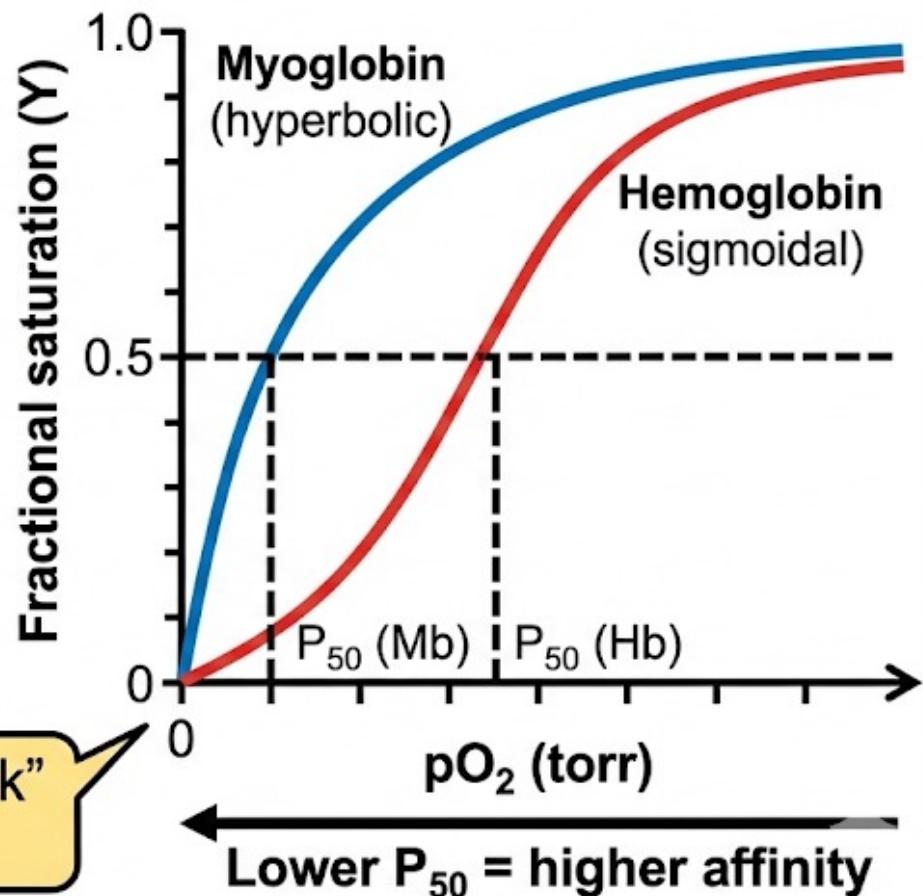
Hemoglobin Displays Cooperative Behavior



Reading O₂-binding curves: Y and P₅₀

- Y = fractional saturation (0 to 1): fraction of sites with O₂ bound
- X-axis is pO₂ (partial pressure of oxygen)
- P₅₀ = pO₂ at Y = 0.5 (lower P₅₀ = higher affinity)
- Myoglobin: low P₅₀, hyperbolic curve
- Hemoglobin: higher P₅₀, sigmoidal curve (cooperativity)

P₅₀ is the “half-full tank” oxygen pressure.



The Hill Equation and Hill Coefficient: Quantifying Binding Cooperativity

Mathematical Framework

Hill Equation:

$$Y = \frac{[L]^n}{K_d + [L]^n}$$

n = 1 (Non-cooperative, hyperbolic)

- Sites are independent
- Standard binding
- Hyperbolic curve

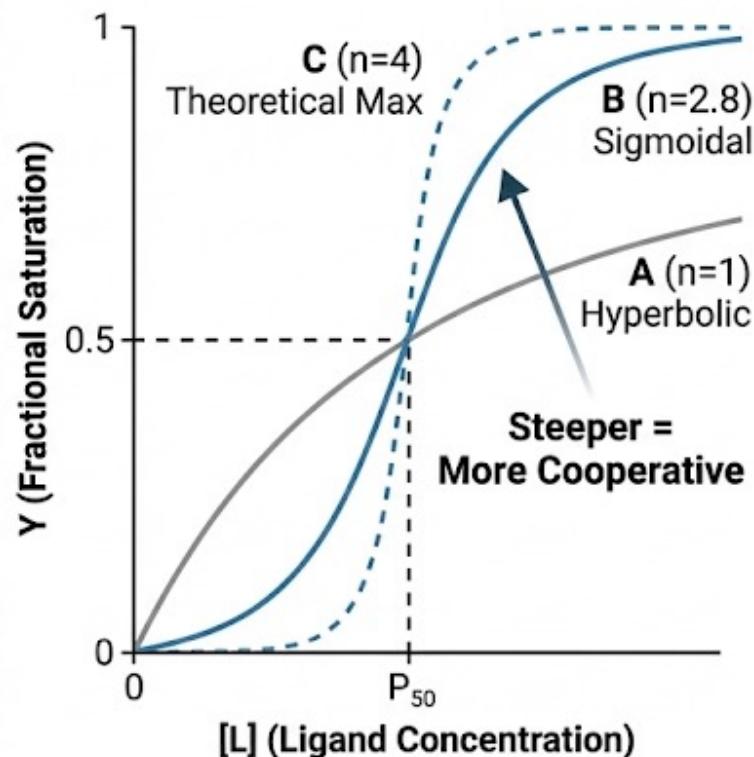
n > 1 (Positive cooperativity, sigmoidal)

- Binding increases affinity
- Sites communicate
- Sigmoidal curve

n < 1 (Negative cooperativity)

- Binding decreases affinity
- Sites communicate
- Sigmoidal curve (shallow)

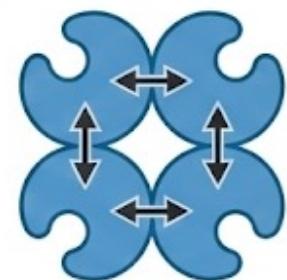
Binding Curves (Y vs [L])



Myoglobin vs. Hemoglobin Summary



Myoglobin
(Mb)



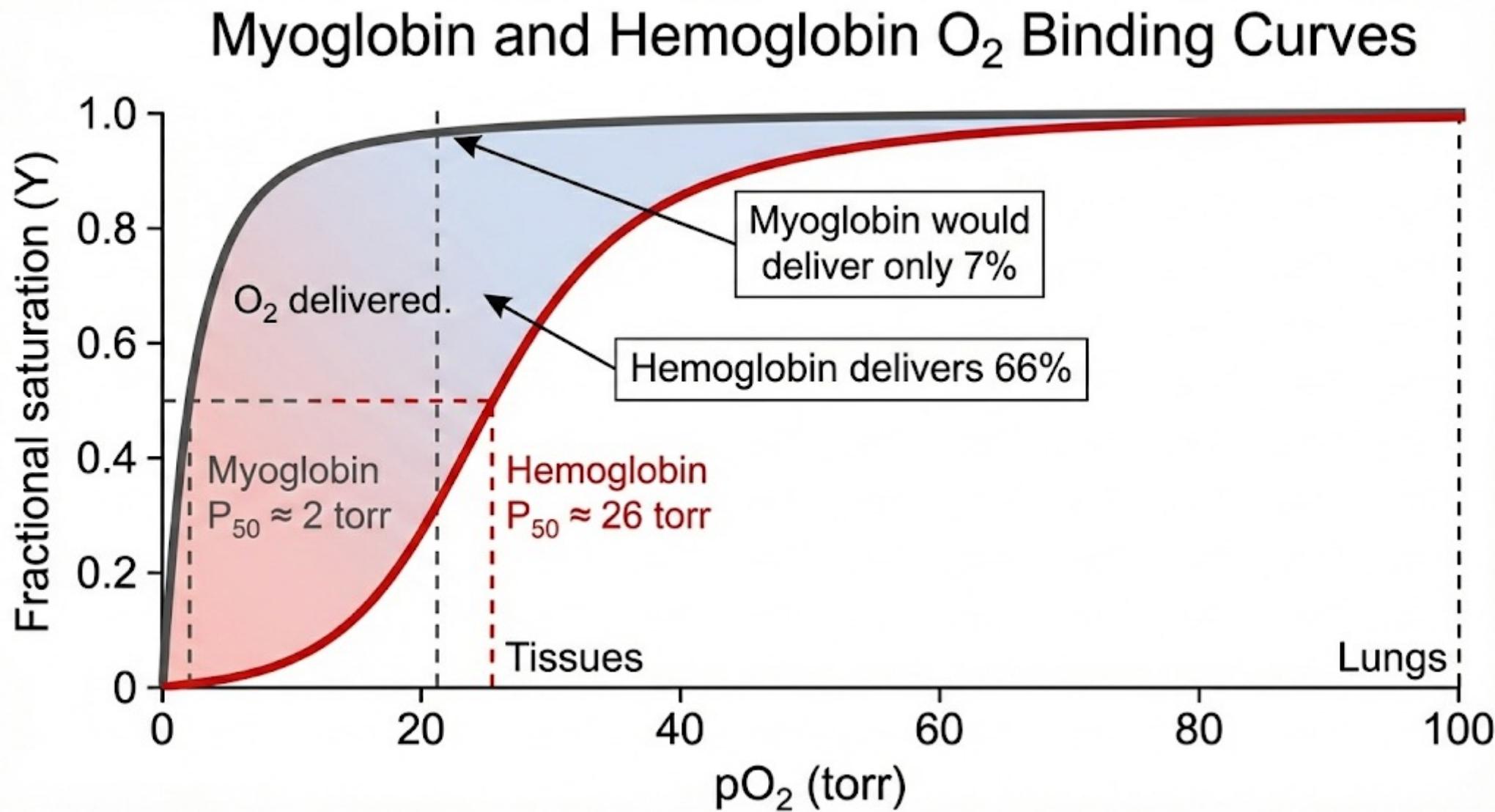
Hemoglobin
(Hb)

$n = 1$
Independent binding

$n \approx 2.8$
Cooperative binding

'The Hill coefficient quantifies how much binding sites communicate.'

Cooperativity Enhances Oxygen Delivery



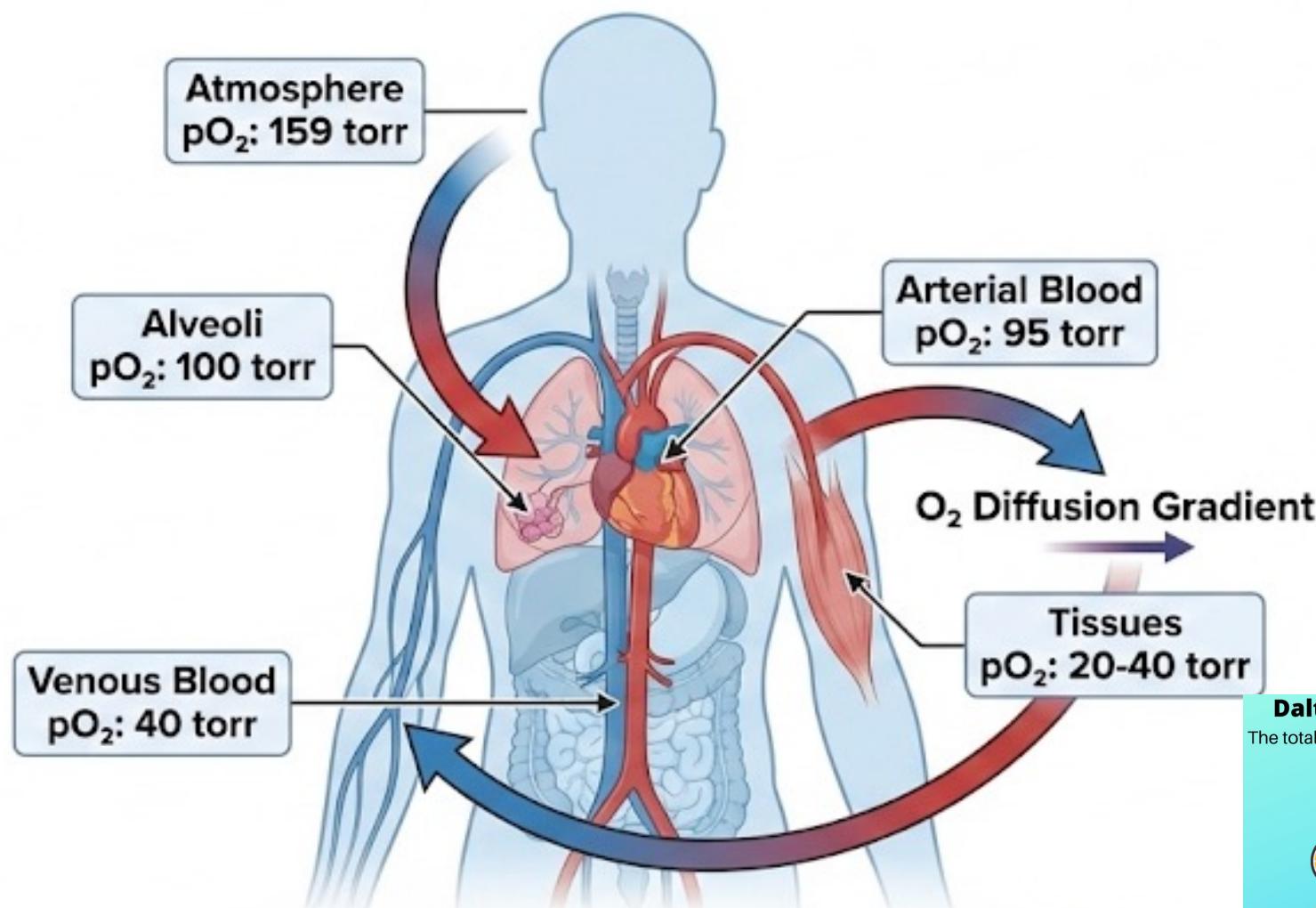
Dalton's Law & Physiological Oxygen Gradient

ATMOSPHERIC COMPOSITION & PARTIAL PRESSURES

78% N₂
(pN₂ = 593 torr)

21% O₂
(pO₂ = 159 torr)

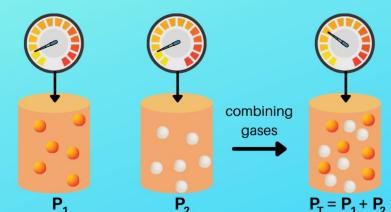
1% Other
(p_{Other} = 8 torr)



Dalton's Law of Partial Pressure

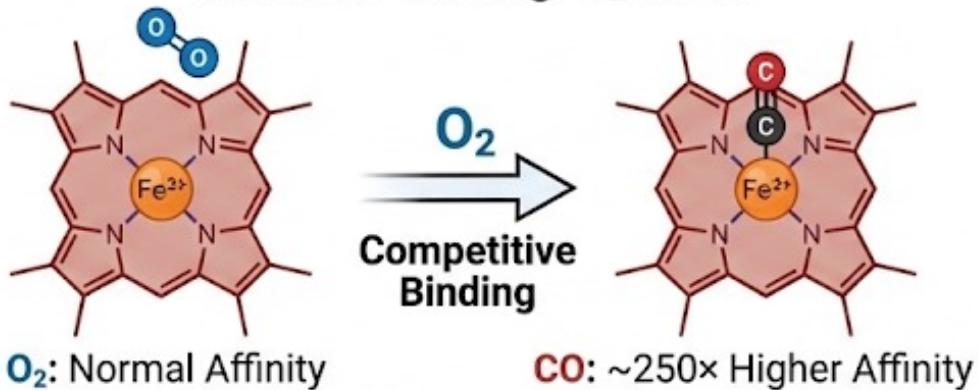
The total pressure of a mixture of gases is the sum of the partial pressures of each gas.

$$P_T = P_1 + P_2 + P_3 + \dots$$

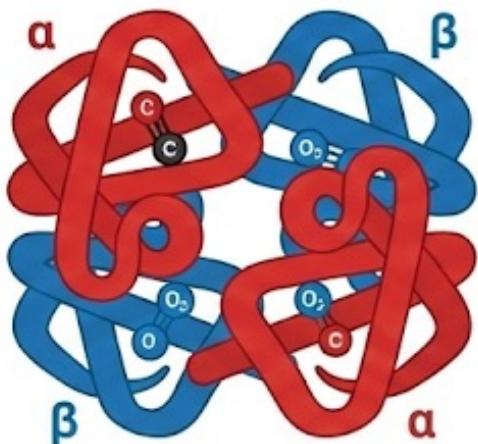


Carbon Monoxide (CO) Poisoning & Hemoglobin

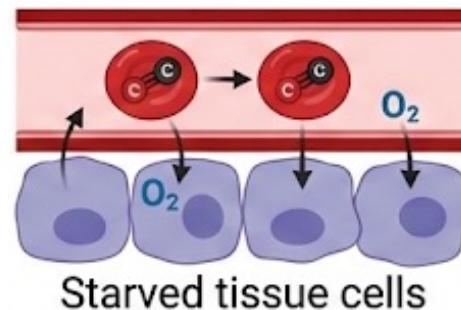
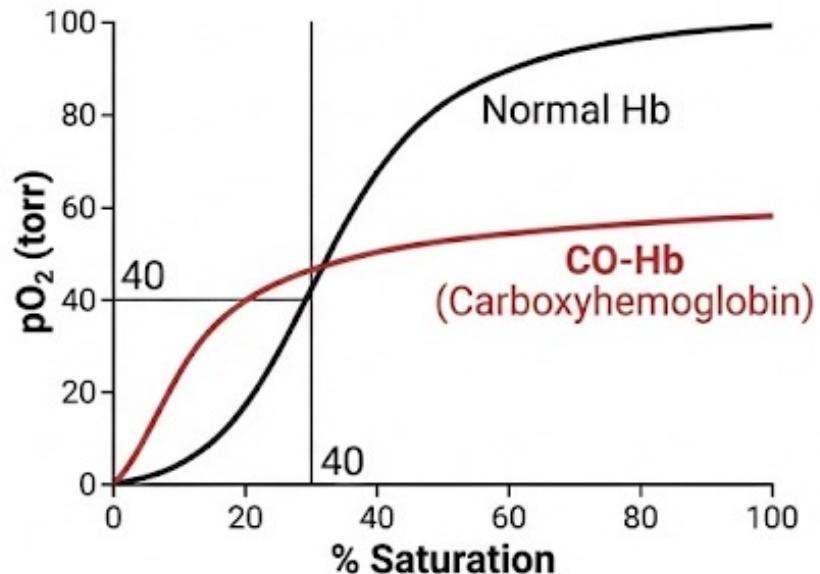
Molecular Binding: O₂ vs. CO



Allosteric Effect & R-State Stabilization



Dissociation Curve & Tissue Hypoxia



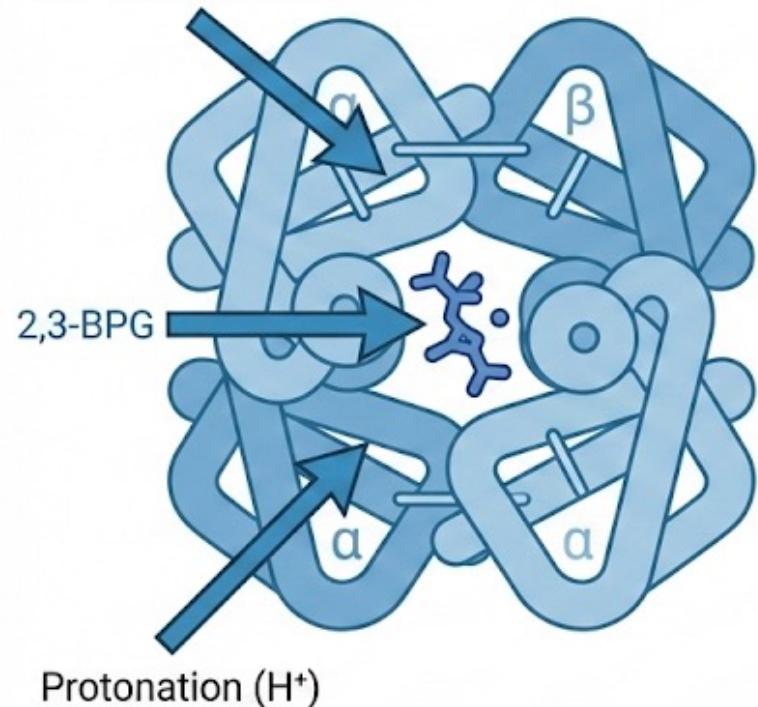
Break



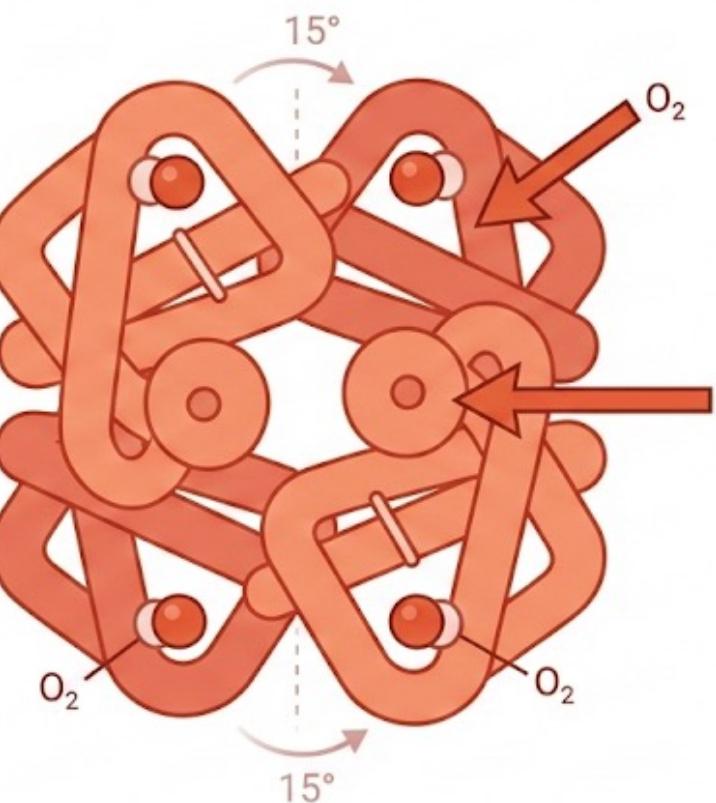
What stabilizes the T state?

T STATE - Low Affinity

Salt bridges
between subunits



R STATE - High Affinity

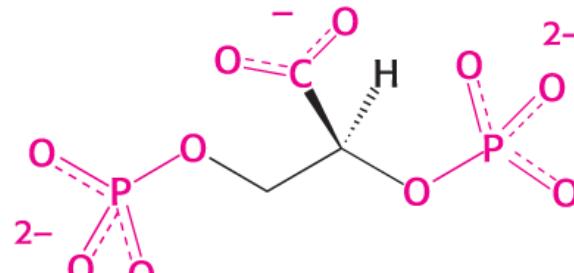


EQUILIBRIUM

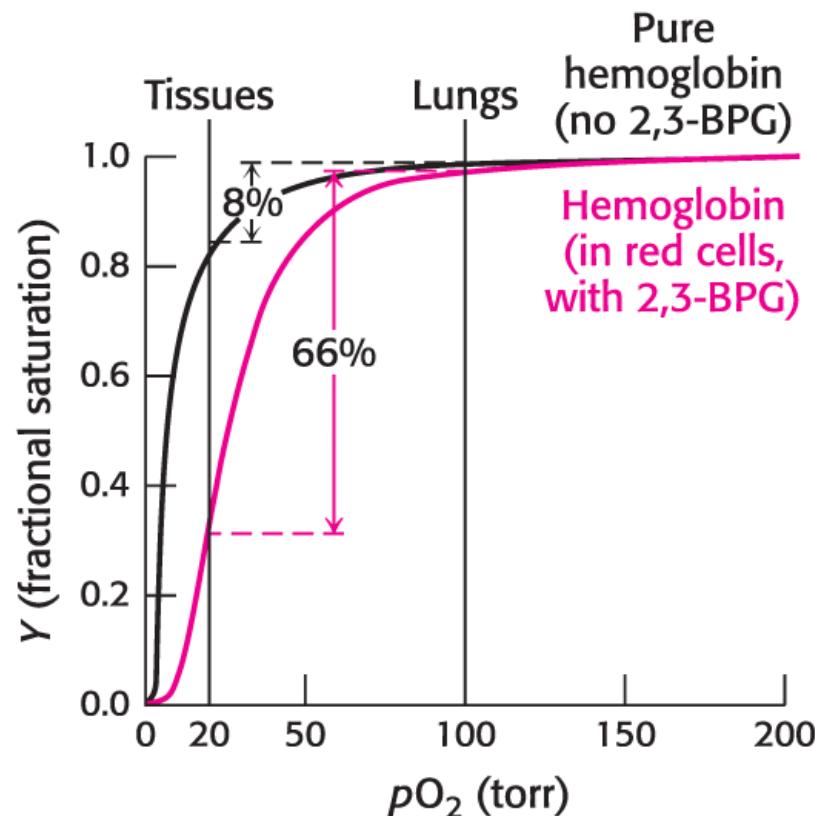
Anything that stabilizes T → Decreases O_2 affinity → Right-shifts curve

Section 5.4 An Allosteric Regulator Determines the Oxygen Affinity of Hemoglobin (1/3)

- Learning objective 8: Identify the key regulators of hemoglobin function.
- 2,3-Bisphosphoglycerate (2,3-BPG) stabilizes the T state of hemoglobin and thus facilitates the release of oxygen.
- 2,3-BPG binds to a pocket in the hemoglobin tetramer that exists only when hemoglobin is in the T state.
- **Tissue Specificity:** In tissues with high metabolic demand or low oxygen levels, 2,3-BPG levels are elevated. This allows hemoglobin to off-load oxygen more efficiently in these tissues.



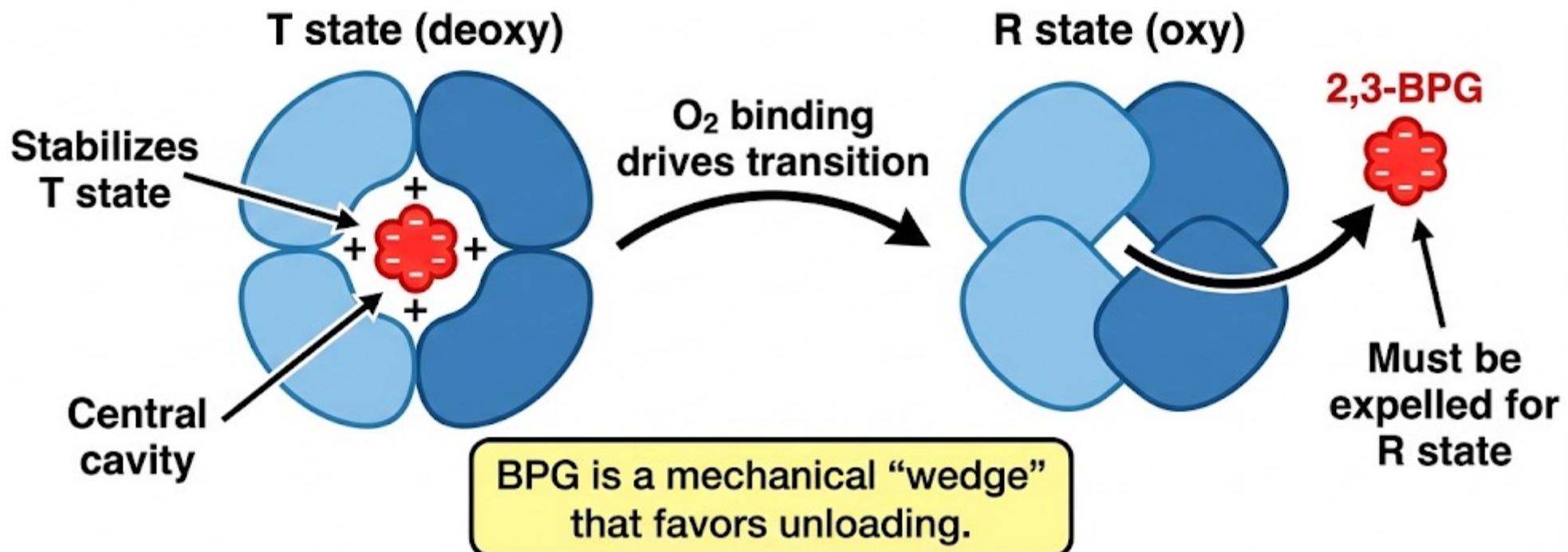
2,3-Bisphosphoglycerate
(2,3-BPG)



Tymoczko et al., *Biochemistry: A Short Course*, 4e, © 2019 W. H. Freeman and Company

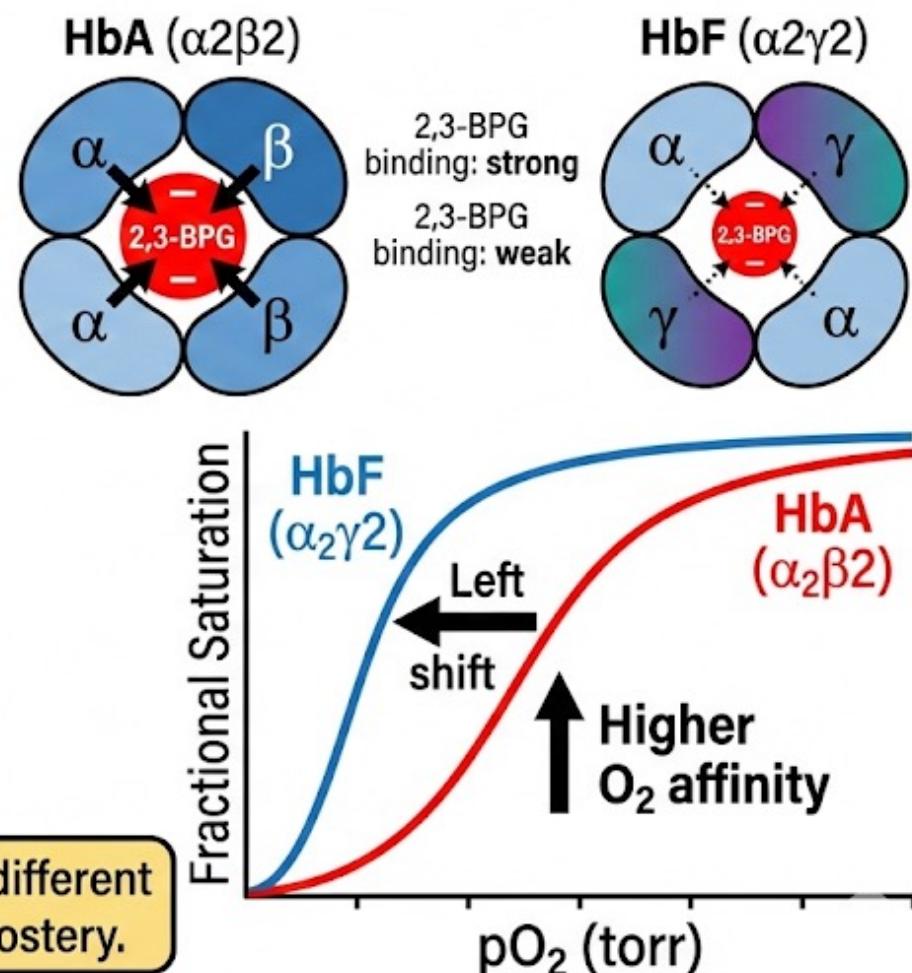
2,3-BPG locks in the T state by occupying the central cavity

- One 2,3-BPG binds the central cavity of deoxyhemoglobin (T state)
- Electrostatic attraction: BPG is highly negative, pocket is positively charged
- Binding stabilizes T state → reduced O₂ affinity → better unloading
- To reach R state, BPG interactions must break and BPG is expelled

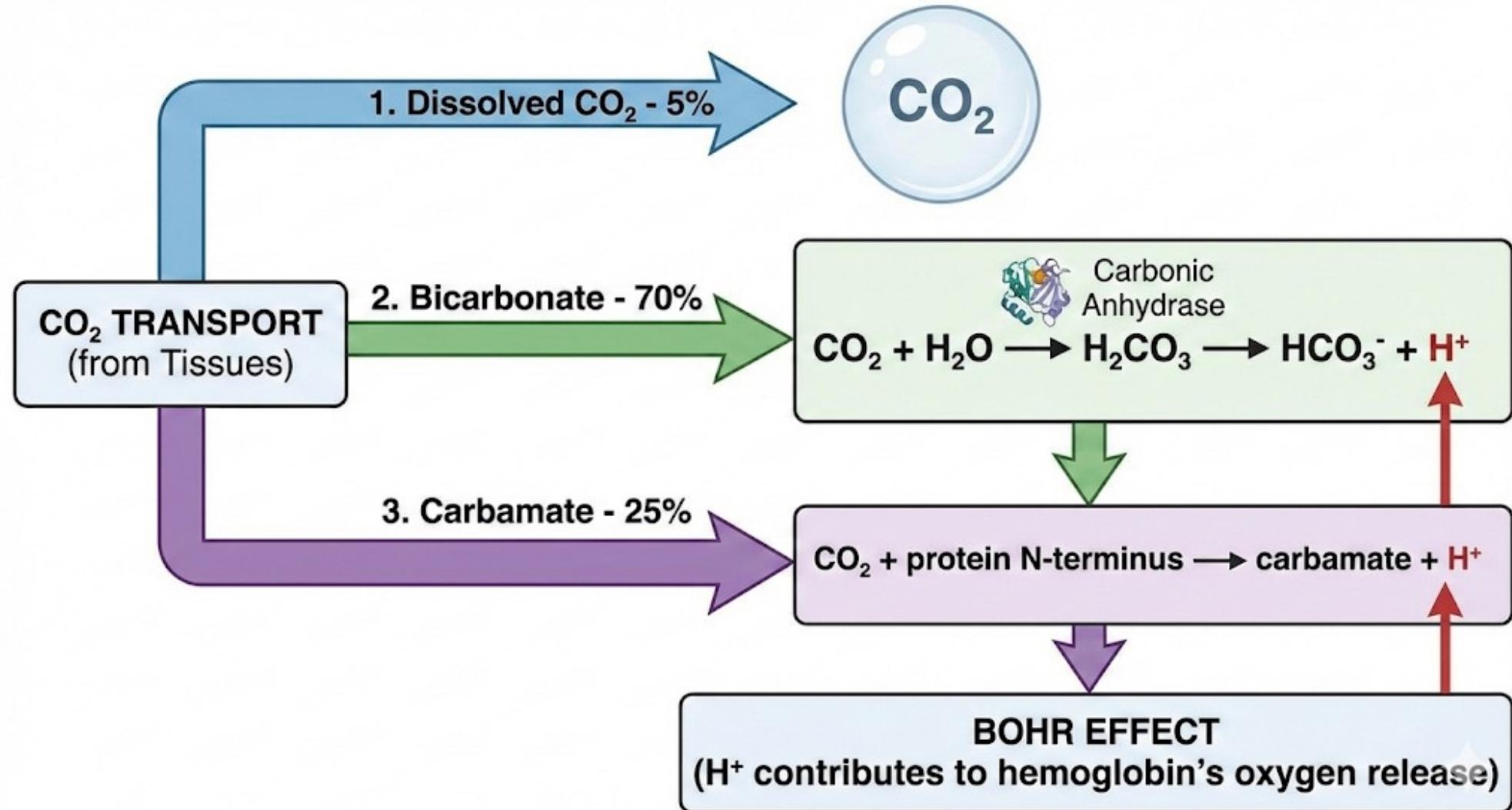


Fetal hemoglobin (HbF) binds O₂ tighter by binding BPG weaker

- Fetus must extract O₂ from maternal **blood** → needs higher O₂ affinity
- HbF tetramer: $\alpha_2\gamma_2$ (not $\alpha_2\beta_2$)
- HbF binds 2,3-BPG more weakly → - less T-state stabilization
- Result: HbF curve is left-shifted relative to HbA**
- Enables efficient O₂ transfer mother → fetus



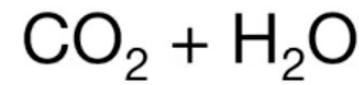
3 modes of CO₂ Transport



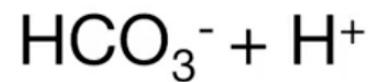
pH Regulation

Peripheral tissues

*From
respiration*



Lower pH



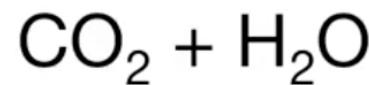
Carbonic
Acid

Bicarbonate

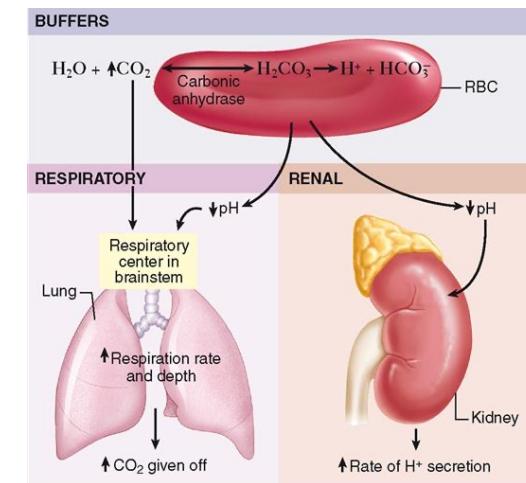
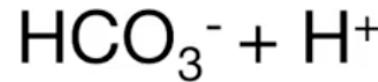
*From
glycolysis*

Lactic acid

Lungs

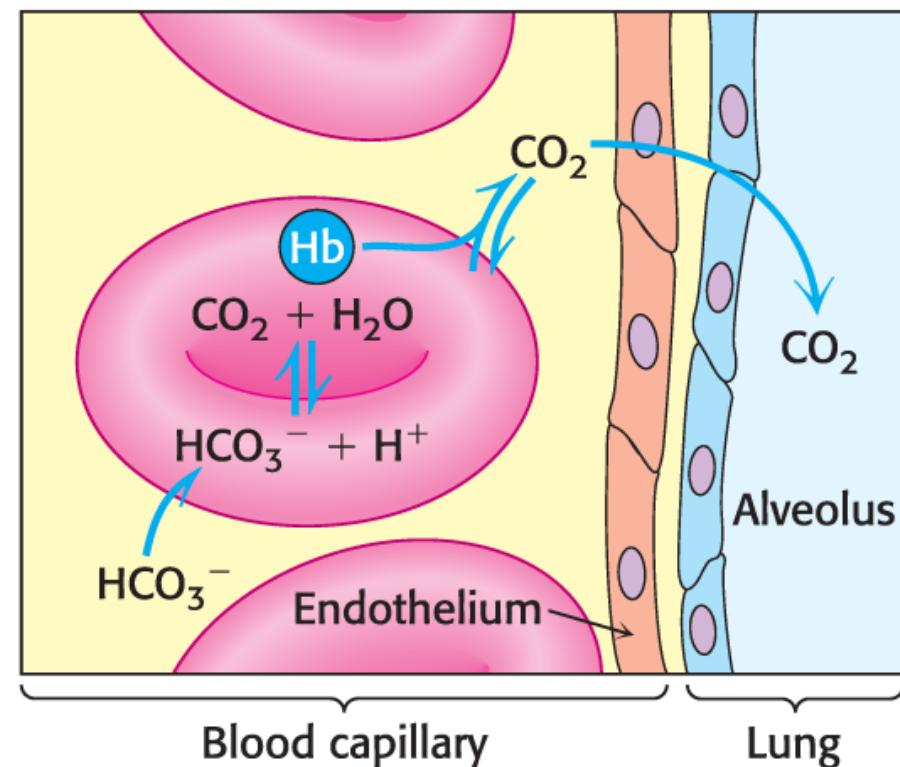
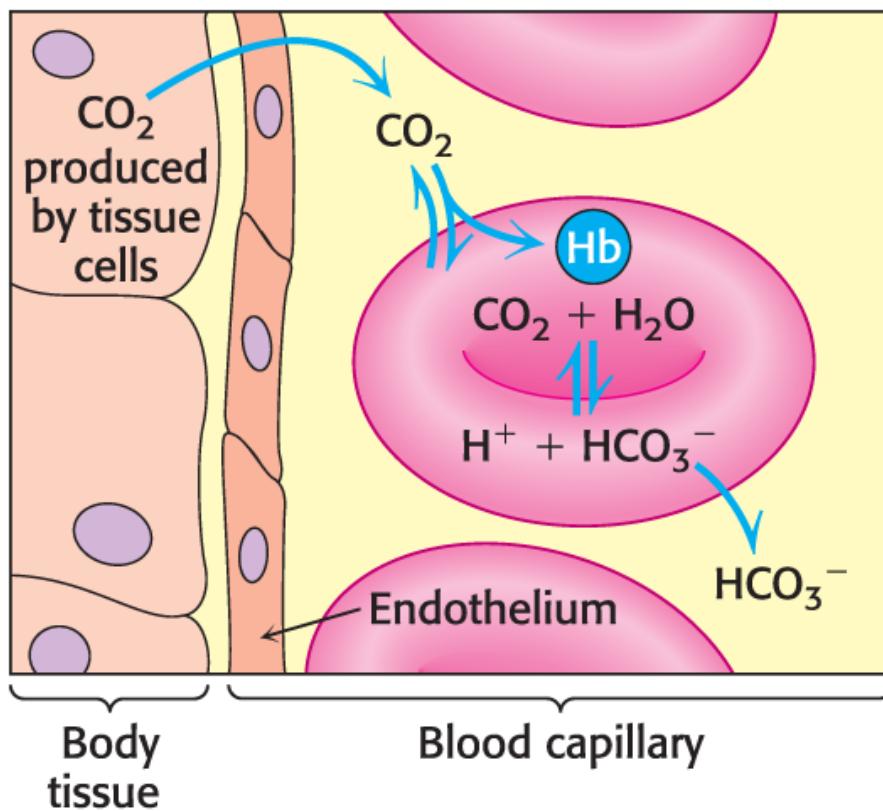


Higher pH



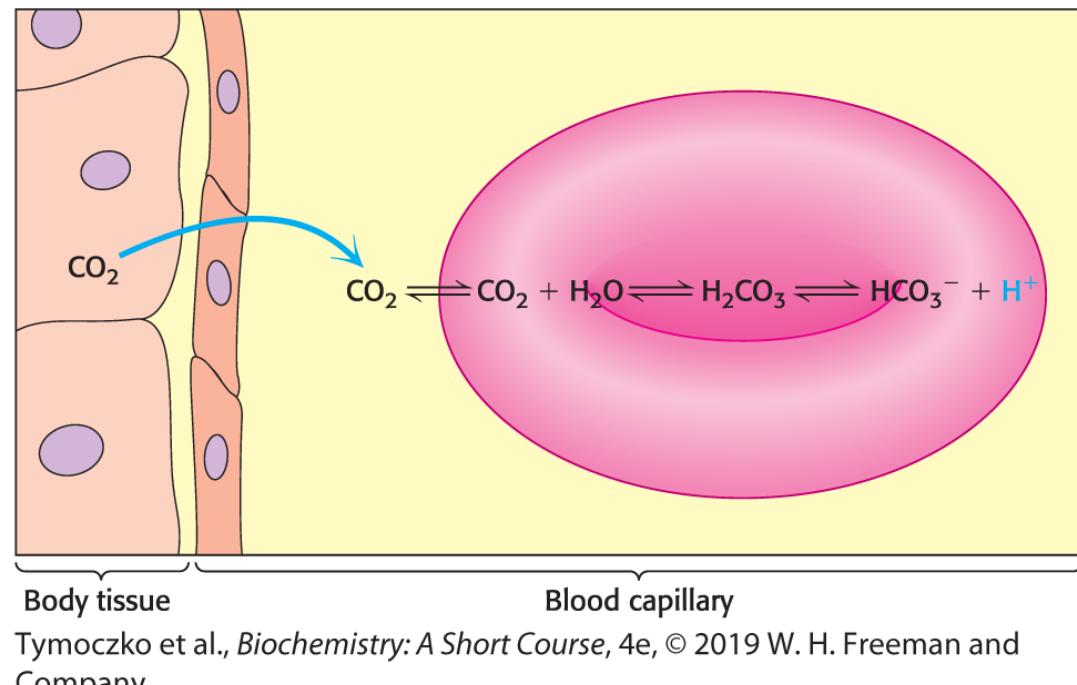
Section 5.5 Hydrogen Ions and Carbon Dioxide Promote the Release of Oxygen (3/3)

- Carbon dioxide is transported to the lungs as bicarbonate.
- Carbonic anhydrase facilitates the formation of bicarbonate ions.



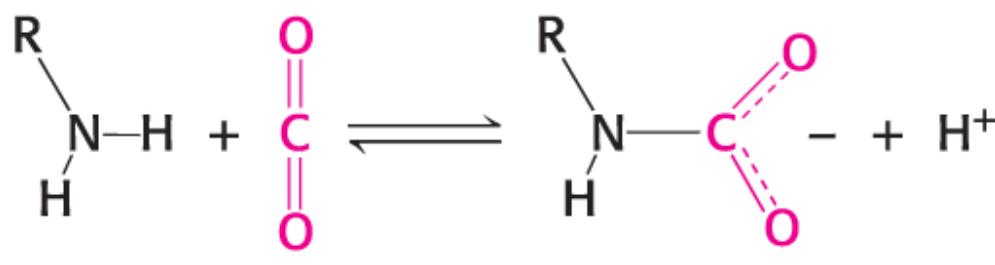
Section 5.5 Hydrogen Ions and Carbon Dioxide Promote the Release of Oxygen (1/3)

- Carbon dioxide and H^+ , produced by actively respiring tissues, enhance oxygen release by hemoglobin.
- The stimulation of oxygen release by carbon dioxide and H^+ is called the Bohr effect.
- carbon dioxide reacts with water to form carbonic acid in a reaction catalyzed by the enzyme carbonic anhydrase

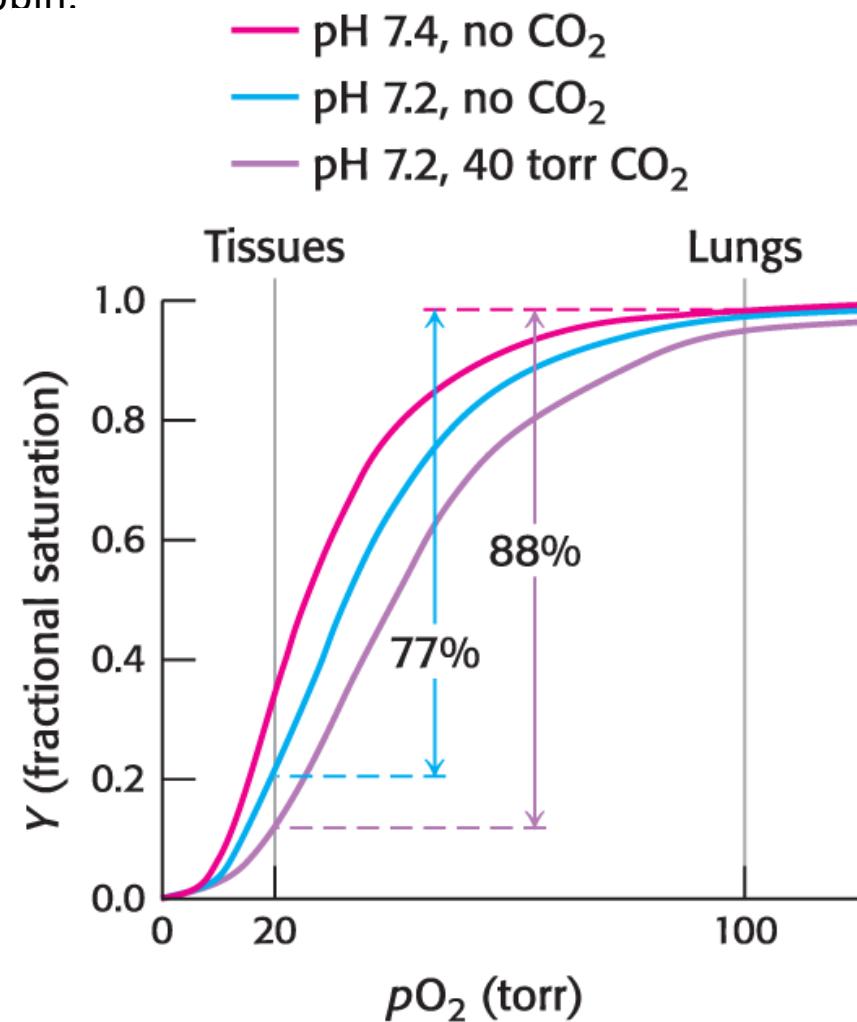
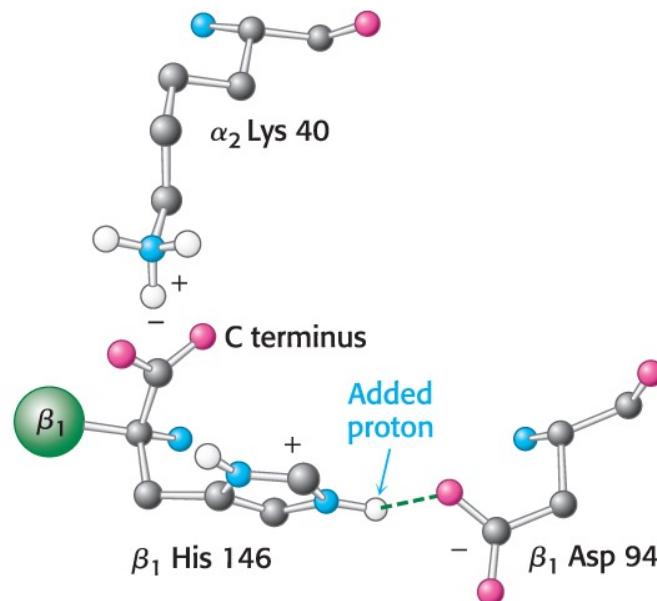


Section 5.5 Hydrogen Ions and Carbon Dioxide Promote the Release of Oxygen

- Low pH allows the formation of ionic interactions that stabilize the T state of hemoglobin, enhancing oxygen release.
- Carbon dioxide reacts with terminal amino groups to form negatively charged carbamate groups. The carbamate forms salt bridges that stabilize the T state.
- Carbon dioxide and H^+ are heterotropic regulators of oxygen binding by hemoglobin.



Carbamate

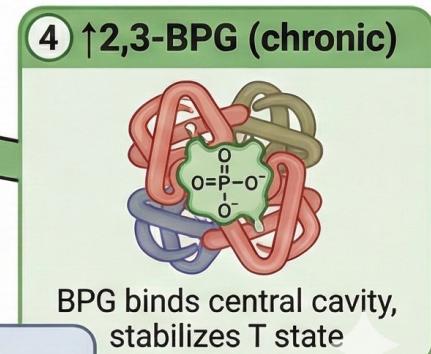
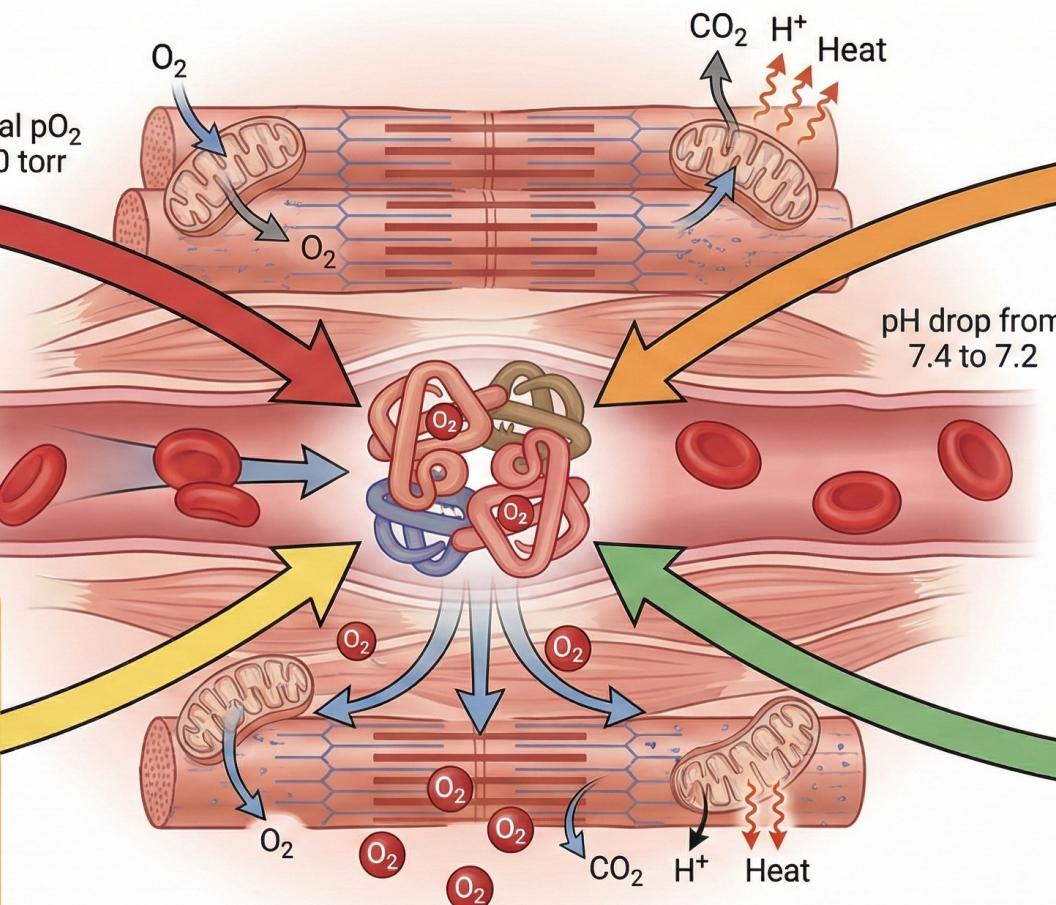
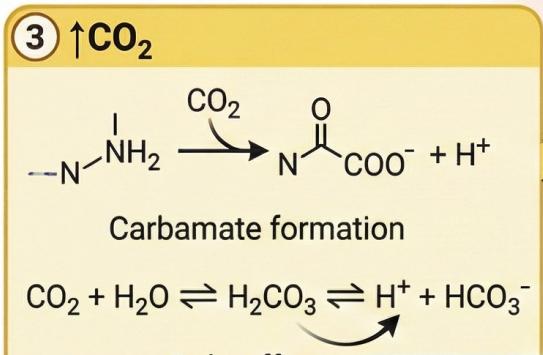
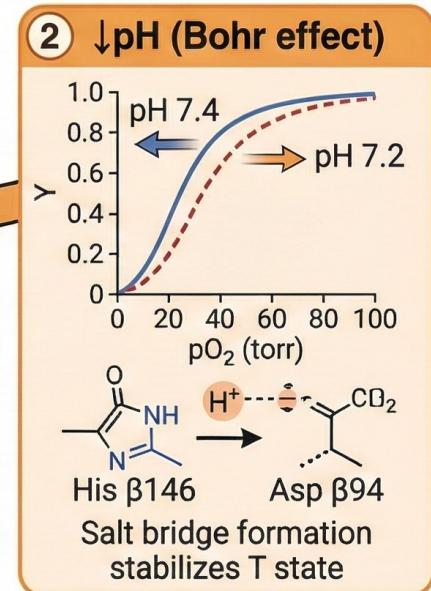
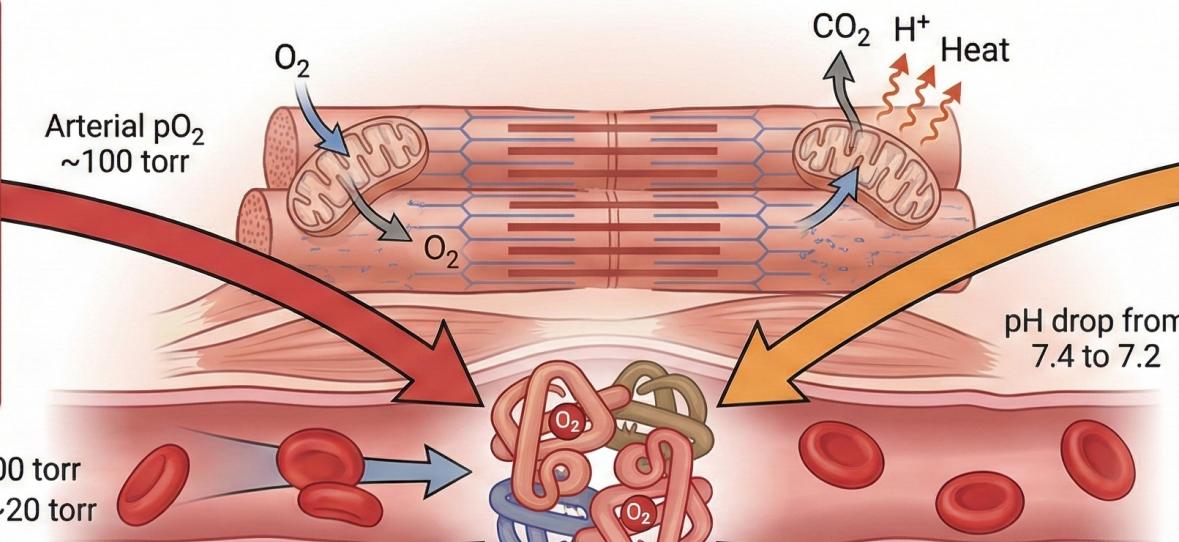
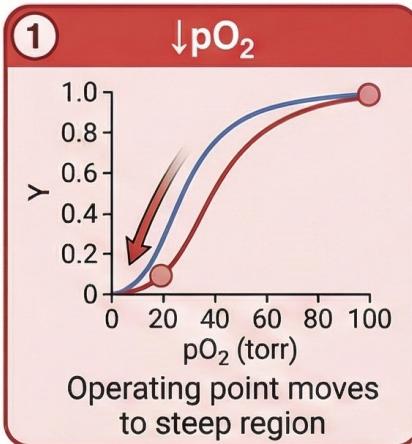


Your exercising muscle drops pO₂ to 20 torr or lower. How does hemoglobin respond—and is cooperativity enough?

Your exercising muscles consume O₂ faster than circulation delivers it—what happens to hemoglobin?

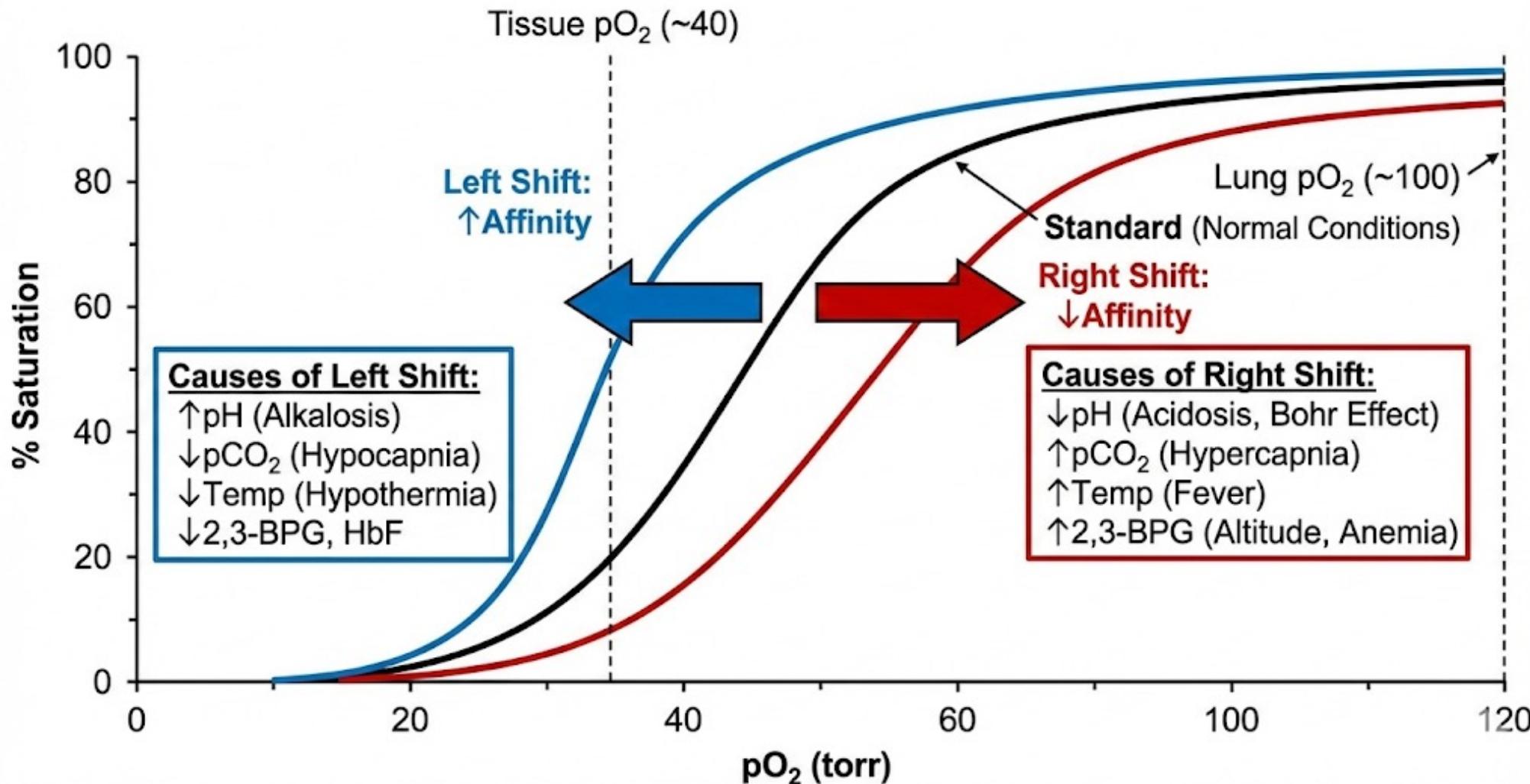


Sinultaneous, Integrated regulation of hemoglobin during exercise



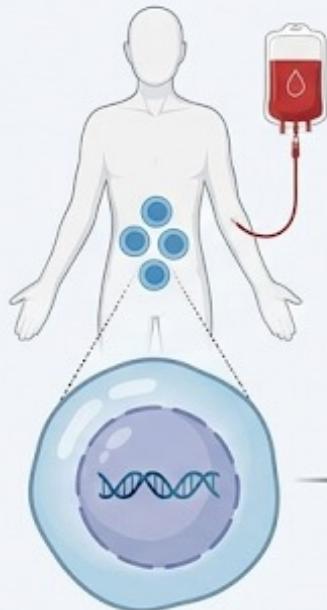
During exercise, **FOUR** mechanisms work together to maximize O₂ delivery:
lower pO₂ + Bohr effect + CO₂ carbamate + BPG all promote T state and O₂ release.

Summary: Hemoglobin-Oxygen Dissociation Curve Shifters (pH, pCO₂, 2,3-BPG, Temperature)



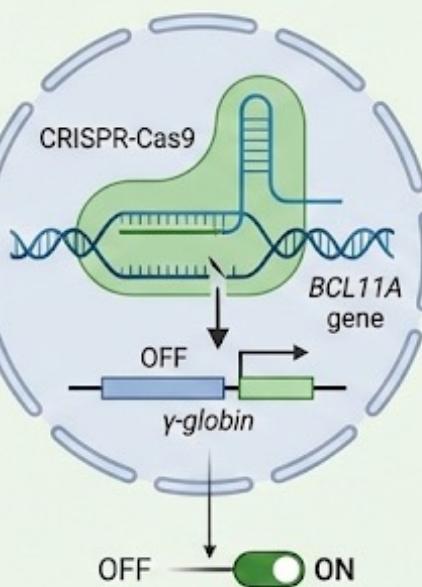
CRISPR GENE THERAPY FOR SICKLE CELL DISEASE: REACTIVATING FETAL HEMOGLOBIN

1. Hematopoietic Stem Cell (HSC) Collection



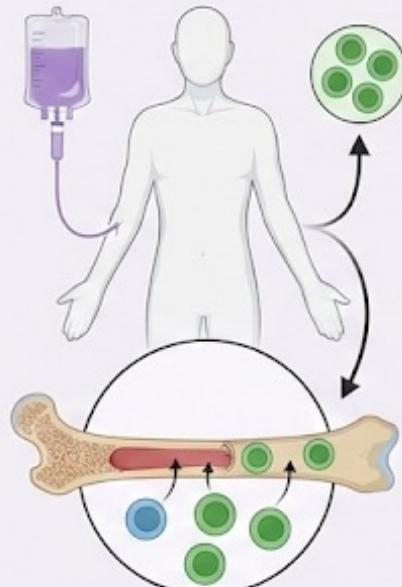
Patient's own stem cells
collected from blood

2. CRISPR Editing (*Ex Vivo*)



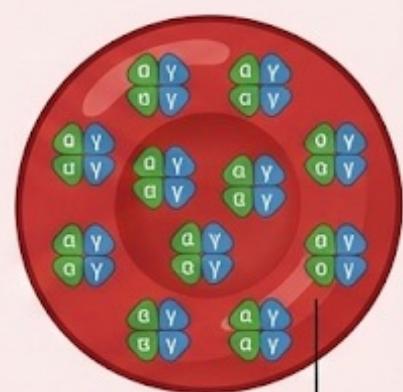
CRISPR-Cas9 edits *BCL11A* gene,
removing the "brake" on fetal
hemoglobin (HbF) production

3. Conditioning & Infusion



Chemotherapy clears patient's
bone marrow, followed by
infusion of edited stem cells

4. Fetal Hemoglobin (HbF) Production



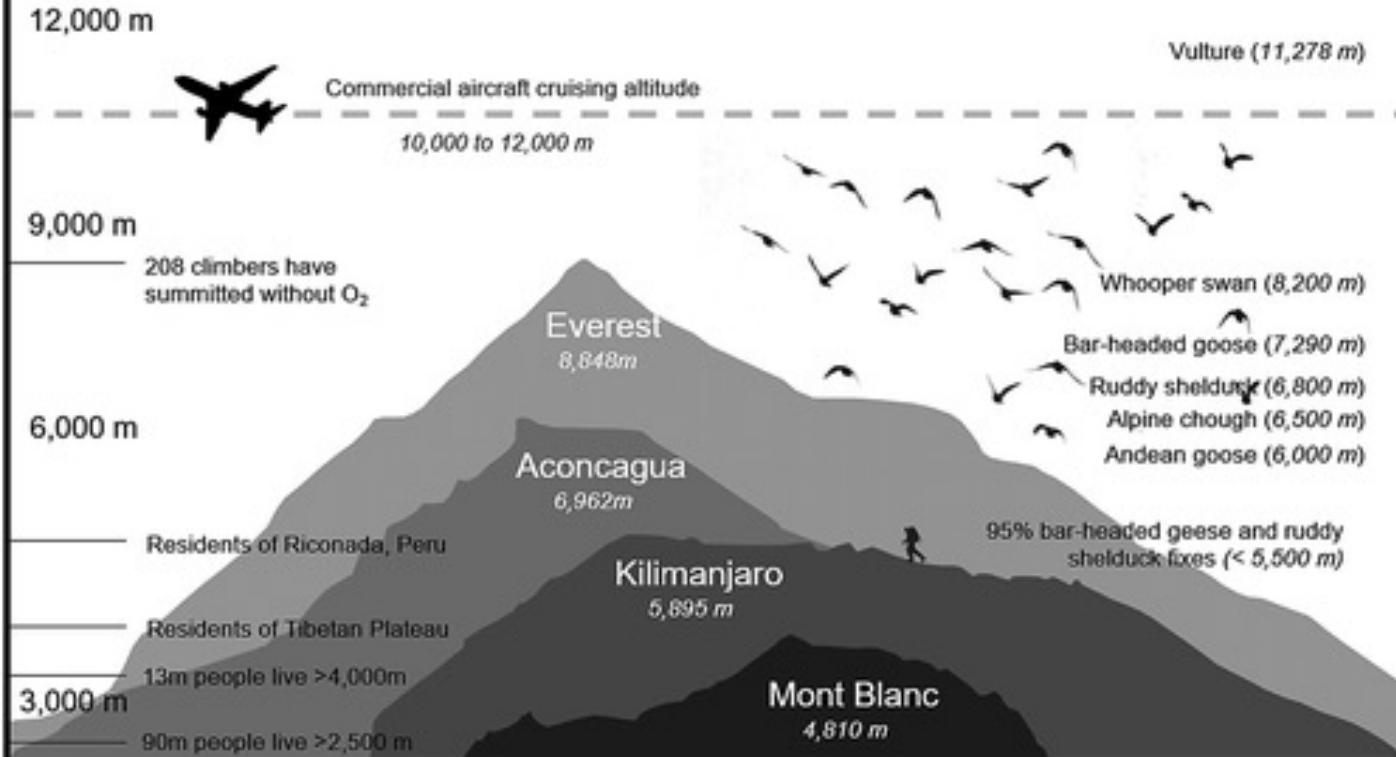
Functional Fetal Hemoglobin
(HbF, $\alpha_2\gamma_2$)

Edited stem cells produce healthy
red blood cells rich in HbF,
preventing sickling

🔗 Directly connects to Fetal
Hemoglobin discussion (Slide 36)

Clinical Translation: Casgevy (exagamglogene autotemcel) demonstrates biochemistry translating to life-changing treatment.

Bar-Headed Goose—High-Altitude Adaptation



- Biological Insight: Hemoglobin Adaptations Allow Oxygen Transport in **Extreme Environments**
- The bar-headed goose can fly over Mount Everest, where the oxygen concentration is low (30% that of sea level).
- Changes in hemoglobin that facilitate the formation of the R state may account in part for this remarkable ability.
- Shifts curve to the left

Apply Your Knowledge: Emergency Department Case



A 34-year-old is brought to the ED after being found in a **running car in a closed garage**. The patient is confused and **lethargic**. Pulse oximetry shows **98% O₂ saturation**, but arterial blood gas reveals **severely low tissue oxygenation**. The patient's lips appear unusually pink.

Questions

1. What molecule is likely bound to the patient's hemoglobin?
2. Why does pulse oximetry show normal saturation despite tissue hypoxia?
3. Using what you know about allosteric regulation, explain the mechanism.



Think time: 60 seconds

Click to Reveal

▶ Hint: Consider what happens when something other than O₂ stabilizes the R state.

Quick Quiz

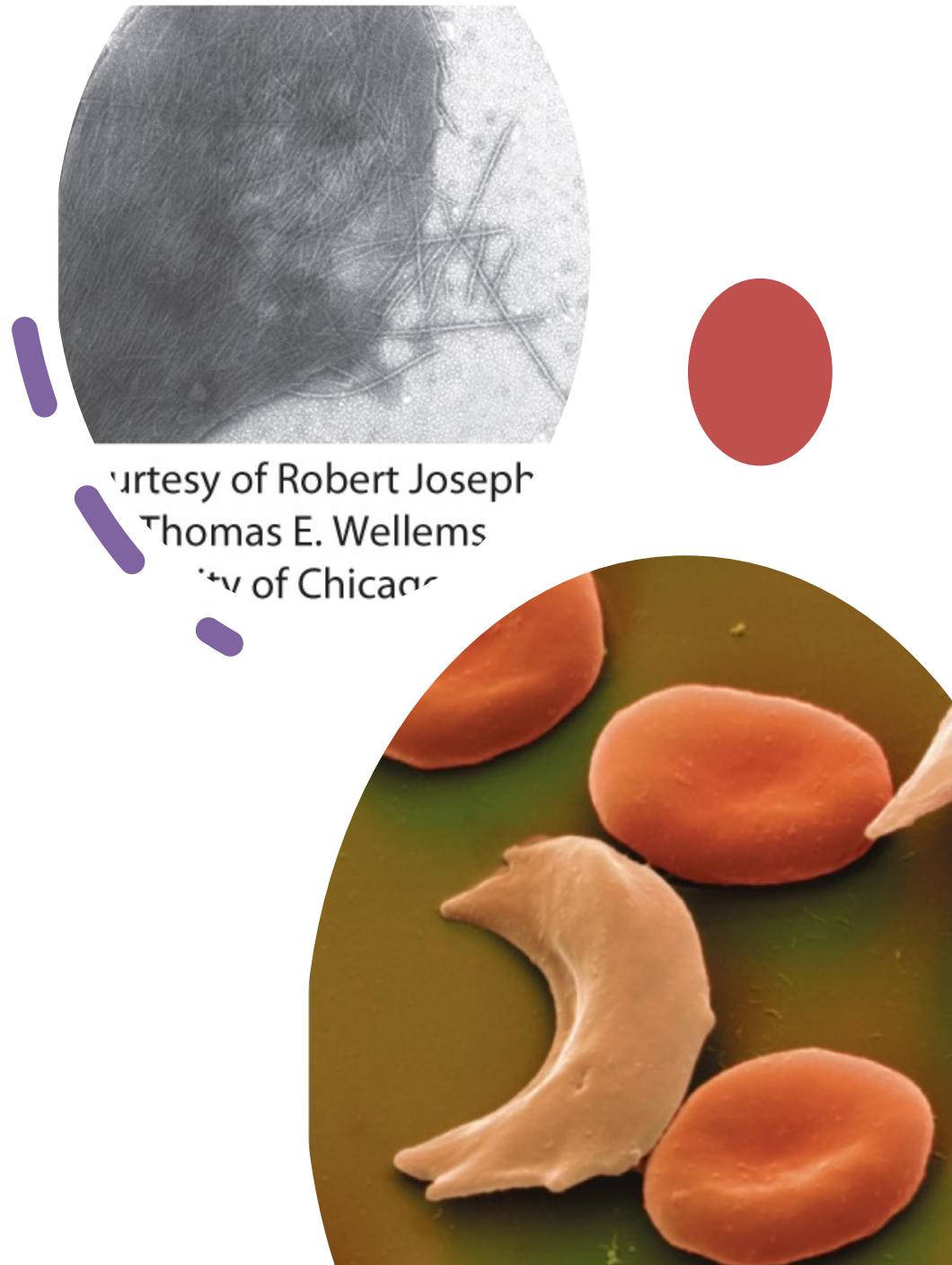


QUICK QUIZ

Name three factors that stabilize the deoxy form of hemoglobin.

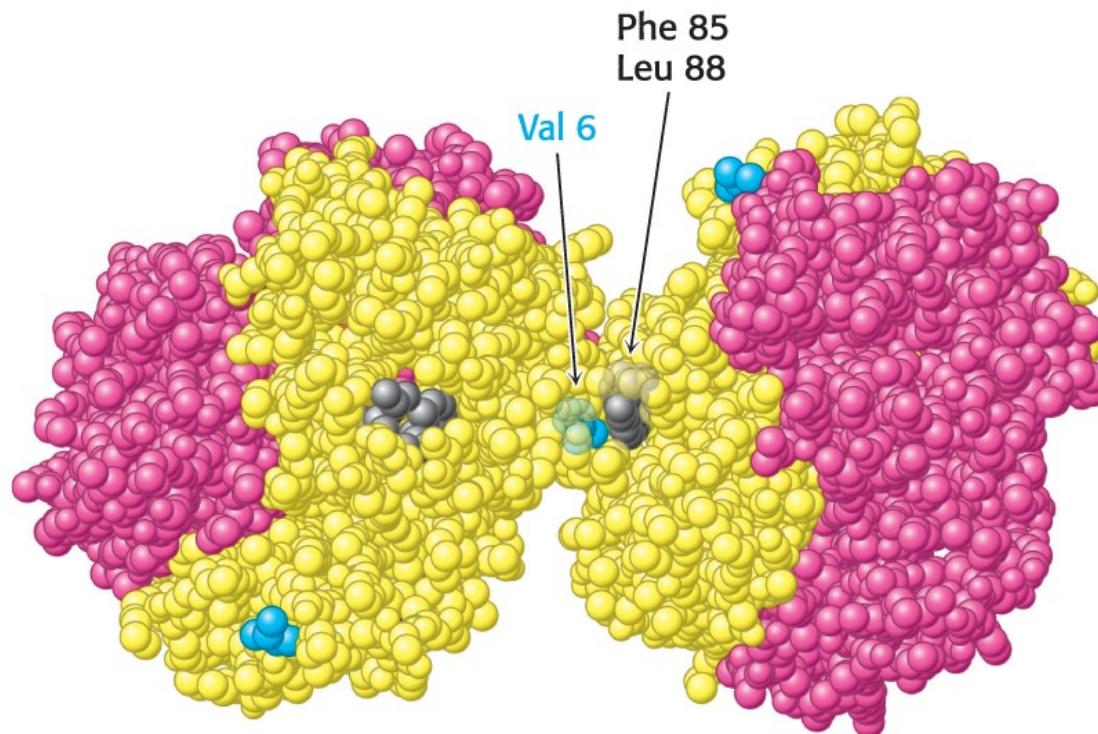
Section 5.6 Mutations in Genes Encoding Hemoglobin Subunits Can Result in Disease

- **Clinical Insight: Sickle-Cell Anemia Is a Disease Caused by a Mutation in Hemoglobin**
- Sickle-cell hemoglobin is called hemoglobin S (HbS). The substituted valine is exposed in deoxyhemoglobin and can interact with other deoxy HbS to form aggregates that deform the red blood cells.
- The sickled cells clog blood flow through the capillaries, leading to tissue damage.
- Sickle-cell anemia is a genetic disease caused by a mutation resulting in the substitution of valine for glutamate at position 6 of the β chains.
- Sickle-cell anemia can be fatal when both alleles of the β chain are mutated.
- In sickle-cell trait, one allele is mutated and one is normal. Such individuals are asymptomatic.
(adaptation to malaria)



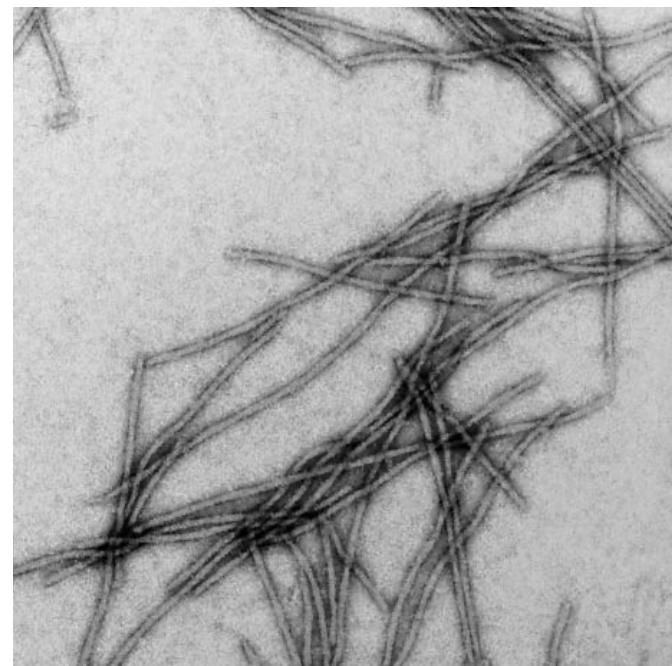
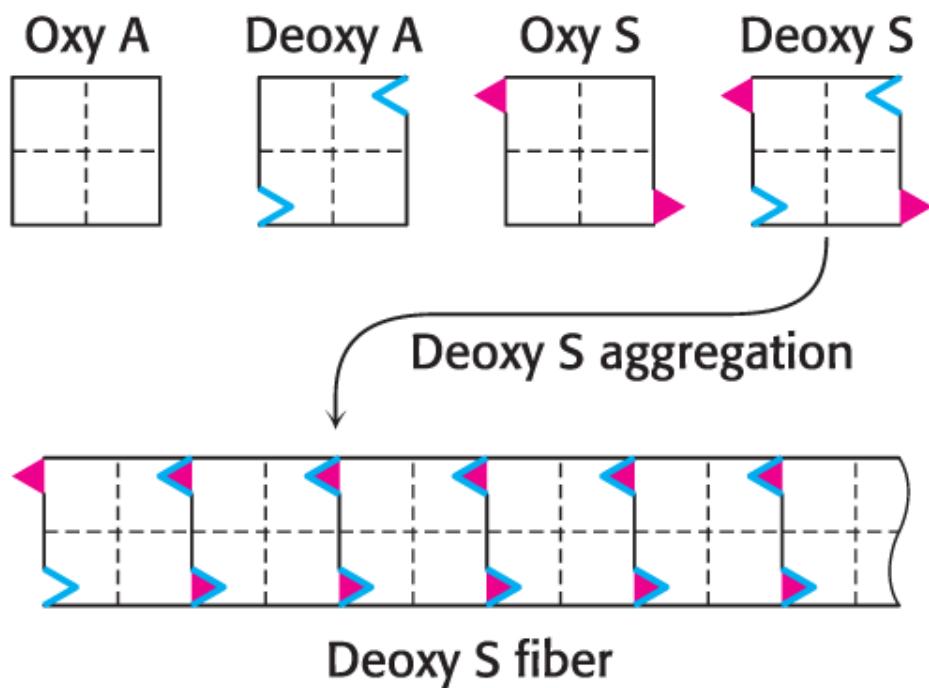
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Structure of Deoxygenated Hemoglobin S



Tymoczko et al., *Biochemistry: A Short Course*, 4e, © 2019
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Diagram of the Formation of Hemoglobin Aggregates

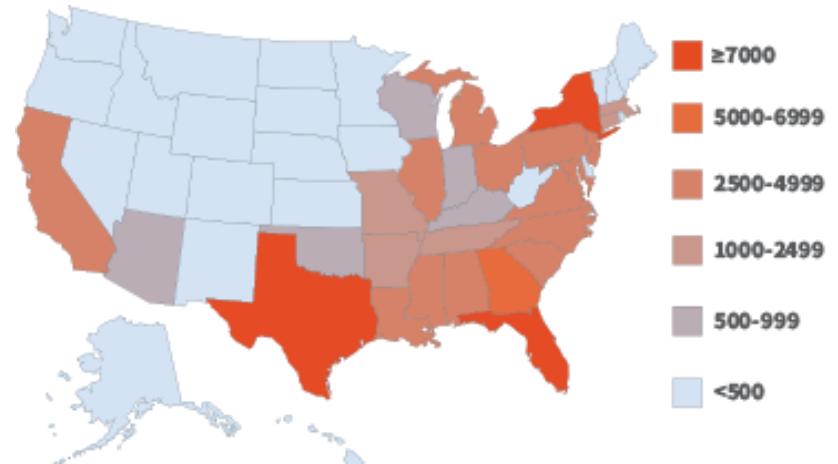


Epidemiology



FACELLY/SIPA/Newscom.

1 in 100 West Africans suffer from Sickle Cell Anemia



MORE THAN **55%** OF PEOPLE WITH SCD RESIDE IN JUST 10 STATES^{2*}

Incidence of SCD¹



Incidence of Sickle Cell Trait¹

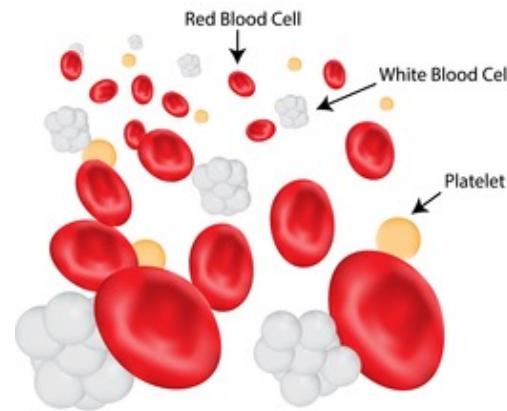


Thalassemia

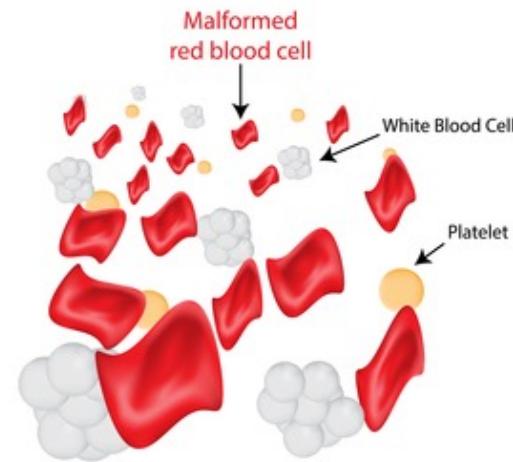
Section 5.6 Mutations in Genes Encoding Hemoglobin Subunits Can Result in Disease (3/3)

- Clinical Insight: Thalassemia Is Caused by an Imbalanced Production of Hemoglobin Chains
- Although sickle-cell anemia is caused by the substitution of a single specific amino acid, thalassemias are caused by a loss or substantial reduction of a single hemoglobin chain.
- In α -thalassemia, the α chain is not produced in sufficient quantity. Tetramers of the β chain form (HbH) and bind oxygen with high affinity but no cooperativity.
- In β -thalassemia, the β chain is not produced in sufficient quantity. The α chains aggregate and precipitate, leading to loss of red blood cells and anemia.

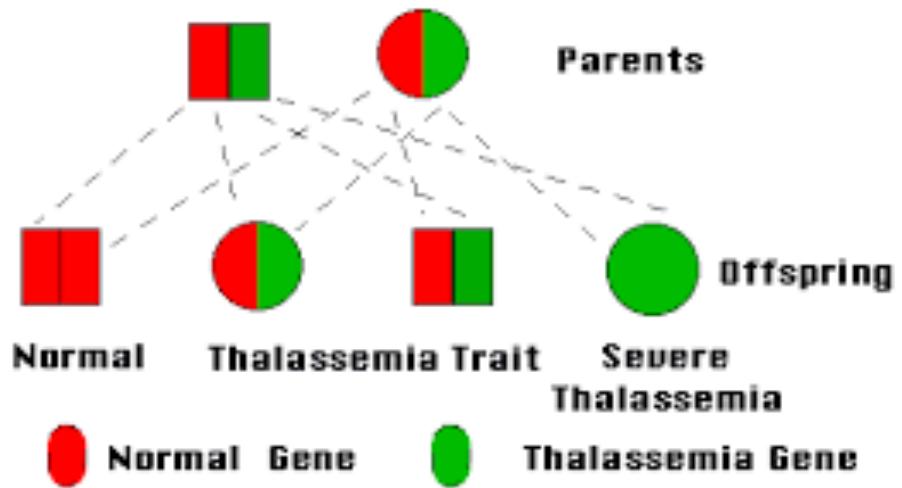
Normal



Thalassemia



Thalassemia Inheritance
Two Carriers



APPENDIX: Biochemistry in Focus

- Additional globins are encoded in the human genome
- The human haploid genome contains various globin genes, two being neuroglobin and cytoglobin.
- Neuroglobin is expressed primarily in the brain and at especially high levels in the retina. It may play a role in protecting neural tissues from hypoxia.
- Cytoglobin is expressed more widely throughout the body.
- Recent research points to an exciting therapeutic role for neuroglobin as a treatment for carbon monoxide poisoning. An altered neuroglobin binds carbon monoxide 500-fold more tightly than hemoglobin, allowing it to strip carbon monoxide from hemoglobin and then be excreted .

	hemoglobin	myoglobin	cytoglobin	neuroglobin	
					
expression sites	red blood cells	skeletal muscle heart smooth muscle	fibroblast cell lineage liver stellate cells CNS/PNS	neurons (CNS,PNS) retina endocrine tissue fish gills	
Fe-atom coordination	penta	penta	hexa	hexa	
oxygen affinity (P₅₀(O₂) in torr)	26	1	1	1	
gene location (human)	α-cluster 16p13 β-cluster 11p15	22q13	14q24	17q25	
locus link	{α} 83587 {β} 64162	4151	114757	58157	
phylogeny	HBA 	HBB 	MB 	CYGB 	NGB 

