

Chem 452 – Lecture 3

Hemoglobin & Myoglobin

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Hemoglobin (Hb) and Myoglobin (Mb) function as oxygen transport and storage molecules in higher organisms. Their functions have been long studied and, together, provide a wealth of examples of how the structure and function of proteins are related.

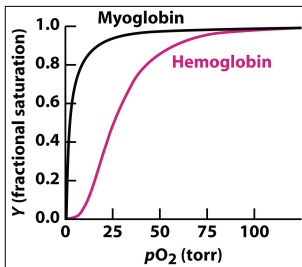
Introduction

- † Together, Hb and Mb provide an excellent example of structure-function relationships in proteins.
- † They illustrate the substrate binding portion of an enzyme catalyzed reaction.
- † They illustrate **allosteric regulation**.

Chem 452, Lecture 3 – Hb and Mb 2

Introduction

- † The cooperative binding of oxygen by Hb, compared to Mb.



Chem 452, Lecture 3 – Hb and Mb 3

Introduction

- † Hb also provided one of the first examples for the molecular basis of genetic diseases.
- † Sickle-cell anemia.

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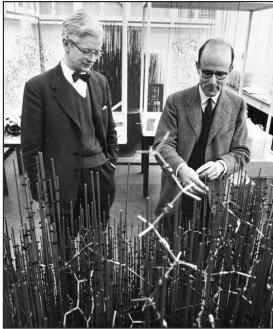
Introduction

- ✦ Mb and Hb were also the first proteins to have their 3-dimensional structures determined.

Introduction

- ✦ The crystal structure of Mb was determined by John Kendrew in 1957 using X-ray diffraction.
- ✦ This was closely followed by the crystal structure for Hb, which was determined by Max Perutz in 1958.

Introduction



John Kendrew Max Perutz

The Medical Research Council (MRC) at Cambridge University

- ✦ Kendrew
- ✦ Perutz
- ✦ Sanger
- ✦ Watson
- ✦ Crick

Nobel Prizes in 1962

Introduction



Nobel Prizes in 1962

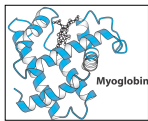
Introduction

- † Hb and Mb provide an excellent example of how proteins have evolved to most efficiently carry out a particular function.
- † Hb binds oxygen in the lung, where the O_2 concentration is high, and delivers it to the tissues where the O_2 concentration is low.
- † Mb accepts the O_2 from the Hb in the tissues where the O_2 concentrations are low.

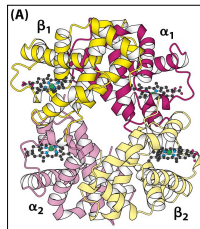
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Introduction

- † Structures of Mb and the α and β subunits of Hb are very similar



Mb



Hb

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Introduction

- † The amino acid sequences for Mb and the α and β chains of Hb are **homologous** (Chapter 6.2-6.4)

```

Hemoglobin  $\alpha$   VLSPADKTNKKAKGGKVGQHAGEYSAALCLRMPLSPFTTKTYPPIH-----
Myoglobin      CLSEGEWQLLNKGGKVFDIPGHSEYMLIRLKGHGHLELEDKQHLKS
               FLSHGDAQVKKKKKADRTNVAHVDDMPNALSASDILHAKLRVDPV
               EEEMKASGEDIKMGATULALGGILKKKGHHEAIKPLAQSHATQKIPVK
               NFKLLSHCLLVTAARHPAETTPAVHASLQRFASVSTVLTQNR
               YLEFIISGLTQVLDZSKHPGDSGADAQGAMNQLFRKDMASQKELGFQG

```

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Introduction

- † The amino acid sequences for Mb is also homologous to the sequence for the plant protein leghemoglobin

```

Hemoglobin  $\alpha$   VLSPADKTNKKAKGGKVGQHAGEYSAALCLRMPLSPFTTKTYPPIH-----
Myoglobin      CLSEGEWQLLNKGGKVFDIPGHSEYMLIRLKGHGHLELEDKQHLKS
               FLSHGDAQVKKKKKADRTNVAHVDDMPNALSASDILHAKLRVDPV
               EEEMKASGEDIKMGATULALGGILKKKGHHEAIKPLAQSHATQKIPVK
               NFKLLSHCLLVTAARHPAETTPAVHASLQRFASVSTVLTQNR
               YLEFIISGLTQVLDZSKHPGDSGADAQGAMNQLFRKDMASQKELGFQG

```

```

Myoglobin      CLSEGEWQLLNKGGKVFDIPGHSEYMLIRLKGHGHLELEDKQHLKS
Leghemoglobin  GALTESQAAVLYKSDWVFNMLPMTHTRFELVLEIADAK---GLSFLSTSEN
               KASE-GLKMGATLTALGGI---LKKKH---HEERIRPDQSHATQKIPVKYLE
               PGNPFLDQKGGPRLVYEAALQEVTEGVVTESTLQHSVYSVSG-VADANRP
               FLSKDTIQVLQSKHPGDFGADAQGAMNQLFRKDMASQKELGFQG
               VVKNALIKTKIEV-----VGAKWSEELNGLTATDELNVTHQDDAA

```

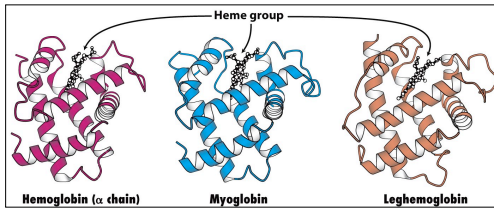
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Introduction

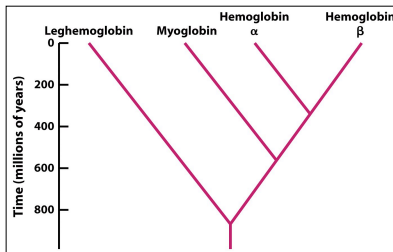
- † These three proteins also have very similar 3-dimensional structures.
- † The tertiary structure appear to be more highly conserved than the primary structure.



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Introduction

- † The amino acid sequences can be used to create an evolutionary tree.



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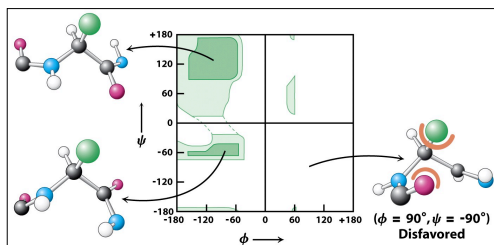
Introduction

- † Number games.
- † We have seen how the Levinthal's Paradox suggests that protein folding is directed.
- † There is not enough time to fold a small protein by a brute force approach.
- † A similar numbers game with amino acid sequence reveals the same directed nature to evolution.

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Predicting a Protein's Fold

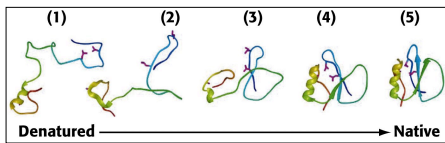
- † The Levinthal Paradox



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Predicting a Protein's Fold

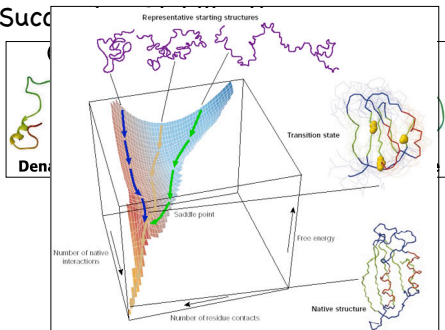
+ Successive Stabilization



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Predicting a Protein's Fold

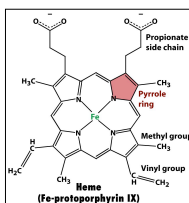
+ Successive Stabilization



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The O₂ binding site for Hb & Mb

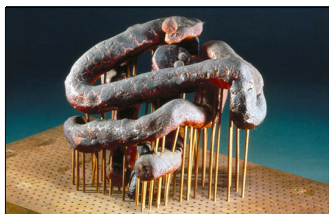
- + Analogous to the active site of enzymes.
- + Heme group is an example of a protein **cofactor**.



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The O₂ binding site for Hb & Mb

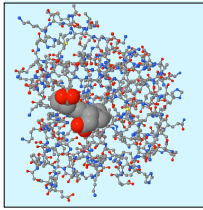
- + Analogous to the active site of enzymes.
- + Heme group provides an example of a **cofactor**.



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The O₂ binding site for Hb & Mb

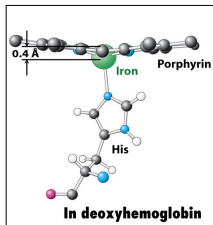
- † Analogous to the active site of enzymes.
- † Heme group provides an example of a **cofactor**.



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The O₂ binding site for Hb & Mb

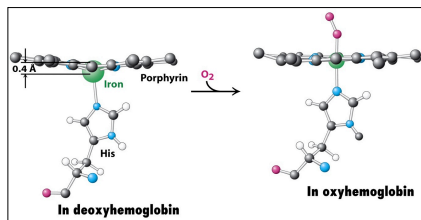
- † The heme Fe²⁺ is ligated by the heme nitrogens and the nitrogen on the **proximal histidine**.



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The O₂ binding site for Hb & Mb

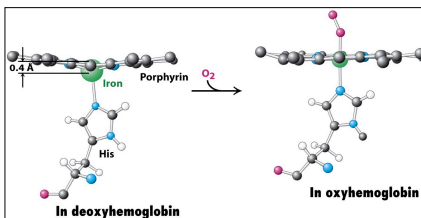
- † When bound, O₂ provides the sixth ligand for the heme Fe²⁺



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The O₂ binding site for Hb & Mb

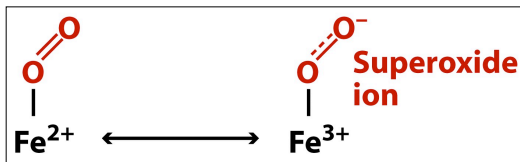
- † When O₂ binds, the heme Fe²⁺ gets smaller and moves into the plane of the heme.



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The O₂ binding site for Hb & Mb

- † The heme Fe²⁺ reduces the bound O₂ to a superoxide ion, O₂⁻.



Chem 452, Lecture 3 - Hb and Mb 23

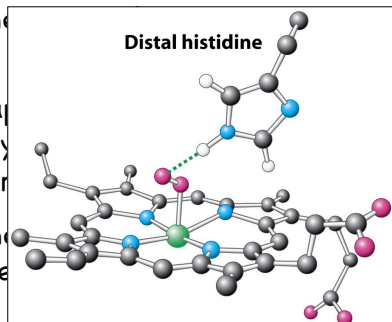
The O₂ binding site for Hb & Mb

- † The heme Fe²⁺ reduces the bound O₂ to a superoxide ion, O₂⁻.
- † Superoxide, like other reactive oxygen species (ROS's), is very damaging.
- † It is the **distal histidine** that helps to prevent the release of the superoxide.

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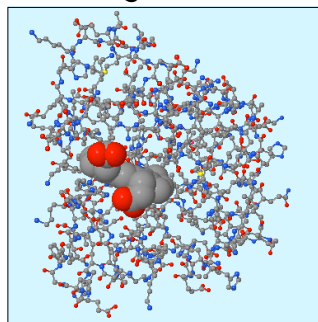
The O₂ binding site for Hb & Mb

- † The heme Fe²⁺ reduces the bound O₂ to a superoxide ion, O₂⁻.
- † Superoxide, like other reactive oxygen species (ROS's), is very damaging.
- † The distal histidine helps to prevent the release of the superoxide.



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The O₂ binding site for Hb & Mb



Oxymyoglobin

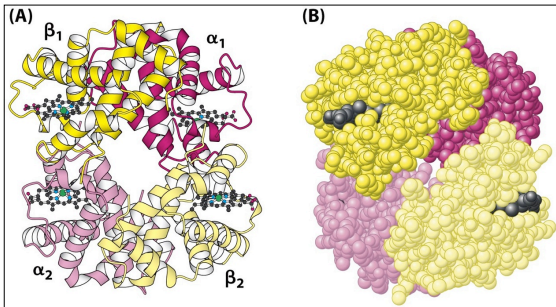
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Hb is a Tetramer

- + Hb's quaternary structure causes it to bind O_2 differently than Mb
- + Hb is a tetramer of myoglobin-like subunits
 - + Two α subunits
 - + Two β subunits
- + Combine as two $\alpha\beta$ dimers
 - + $\alpha_1\beta_1$ and $\alpha_2\beta_2$

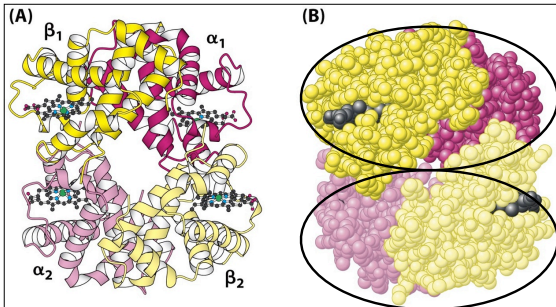
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Hb is a Tetramer



Chem 452, Lecture 3 - Hb and Mb 28

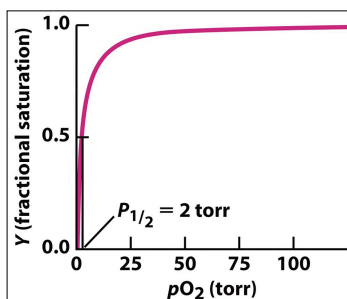
Hb is a Tetramer



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Hb Binds O_2 Cooperatively

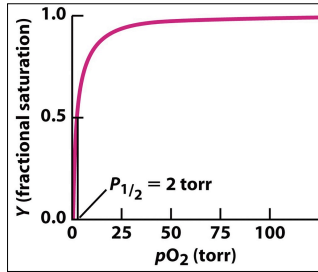
- + Mb has a P_{50} of 2 Torr



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Questions

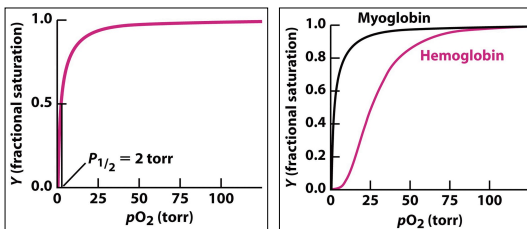
When exposed to air at 1 atm pressure, what fraction of the myoglobin molecule will be bound with O_2 ?



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Hb Binds O_2 Cooperatively

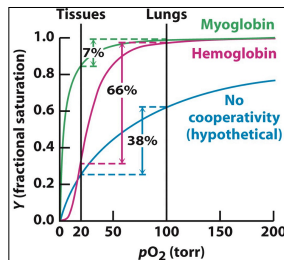
• Hb binds O_2 more weakly than Mb



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Hb Binds O_2 Cooperatively

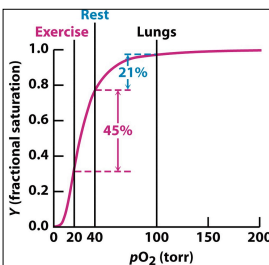
• Cooperative binding makes Hb a more efficient transporter of O_2 than Mb.



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Hb Binds O_2 Cooperatively

• Hb efficiently delivers O_2 to the tissues during stress or exercise.



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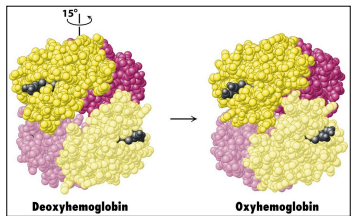
Problem 7.12 & 7.14

For Friday, work Problems 12 and 14 at the end of Chapter 7 and be ready to discuss them in class.

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Hb Binds O_2 Cooperatively

- Cooperativity is associated with changes in the quaternary structure of Hb

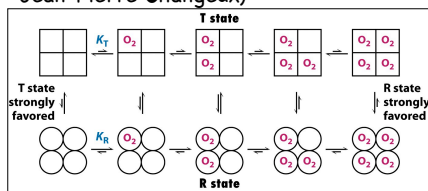


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Hb Binds O_2 Cooperatively

- Models to explain the cooperativity:
 - MWC Model

(Jacques Monod, Jeffries Wyman & Jean-Pierre Changeux)

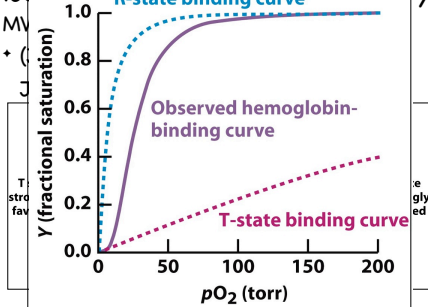


Concerted Model

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Hb Binds O_2 Cooperatively

- Models to explain the cooperativity:
 - MWC Model



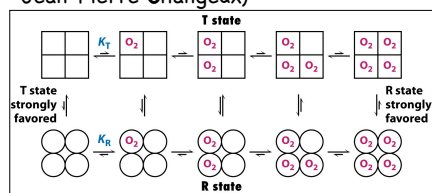
Chem 452, Lecture 3 - Hb and Mb 36

Hb Binds O₂ Cooperatively

Models to explain the cooperativity:

MWC Model

(Jacques Monod, Jeffries Wyman & Jean-Pierre Changeux)



Concerted Model

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Hb Binds O₂ Cooperatively

Models to explain the cooperativity:

Sequential Model



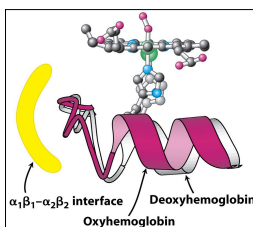
Sequential Model

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Hb Binds O₂ Cooperatively

At the molecular level.

Conformational changes occurring upon O₂ bonding to one subunit are transmitted to other subunits

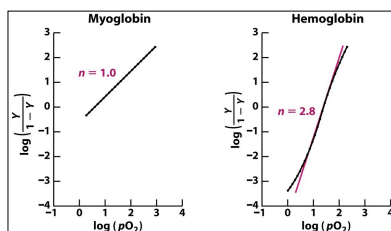


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Hb Binds O₂ Cooperatively

Cooperativity can be assessed with a Hill plot.

n is the Hill coefficient

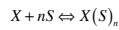


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Hb Binds O₂ Cooperatively

• Cooperativity can be assessed with a **Hill plot**.

• **n** is the **Hill coefficient**



$$Y = \frac{[S]^n}{[S]^n + [S_{50}]^n}$$

$$Y = \frac{pO_2^n}{pO_2^n + P_{50}^n}$$

$$\frac{Y}{1-Y} = \frac{pO_2^n}{P_{50}^n}$$

$$\log\left(\frac{Y}{1-Y}\right) = n \log(pO_2) - n \log(P_{50})$$

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Problem 7.12a & 7.14

For Friday, work Problems 12a and 14 at the end of Chapter 7 and be ready to discuss it in class.

7.12.a Using the Hill equation, plot an oxygen binding curve for a hypothetical two-subunit hemoglobin with $n = 1.8$ and $P_{50} = 10$ torr.

7.14 Oxygen binding for primitive Hb from a lamprey eel is given

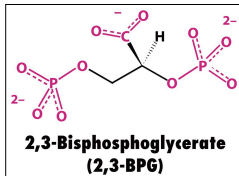
- Plot data and determine P_{50}
- Make Hill plot and determine n
- Propose model to explain cooperativity

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Allosteric Regulation

• Hb provides an example of **allosteric regulation**.

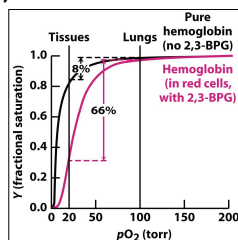
• In red blood cells (RBC's), the metabolite 2,3-Bisphosphoglycerate (2,3-BPG) alters the O₂ binding behavior of Hb.



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Allosteric Regulation

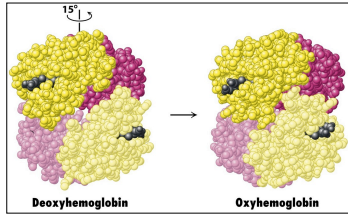
• 2,3-BPG lowers Hb's affinity for O₂, allowing it to release O₂ more efficiently to the tissues.



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Hb Binds O₂ Cooperatively

- Cooperativity is associated with changes in the quaternary structure of Hb

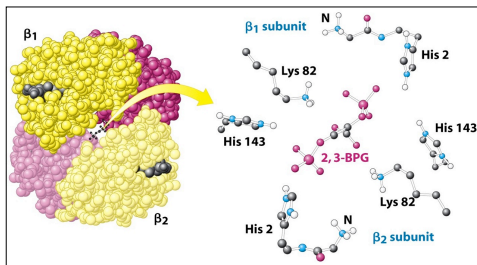


Tense (T) State Relaxed (R) State

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Allosteric Regulation

- 2,3-BPG binds to, and stabilizes, the T-state of Hb.



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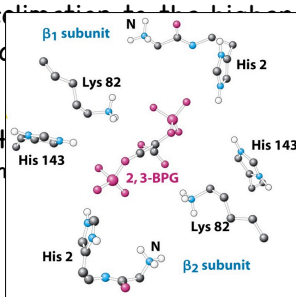
Allosteric Regulation

- The acclimation to the higher elevations involves the production of higher levels of 2-BPG.
- Fetal Hb
 - γ chains are substituted for β chains (H143S)

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Allosteric Regulation

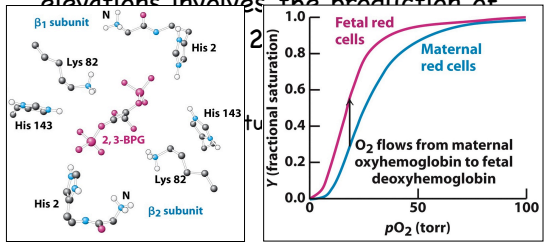
- The acclimation to the higher elevations involves the production of higher levels of 2-BPG.
- Fetal Hb
 - γ chains are substituted for β chains (H143S)



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Allosteric Regulation

- † The acclimation to the higher elevations involves the production of



Next up

- † Hemoglobin and Myoglobin (con'd).
 - † Bohr effect
 - † Sickle-cell Hb
- † Enzymes (Chapter 8)