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# Biochemistry Sheet 10

Globular proteins



### Globular proteins

• Examples → Myoglobin and hemoglobin → they are hemoproteins

#### • Hemoproteins:

- > a group of specialized proteins <u>containing heme</u> as a tightly bound **prosthetic** group
- > Both myoglobin and hemoglobin are hemoproteins
- $\blacktriangleright$  Heme has many functions  $\rightarrow$  dictated by the protein's environment

#### • Heme Group:

- > It is a prosthetic group (a non- protein group covalently attached to a protein)
- Heme group belongs to porphyrins family (it has a ring structure)
- Heme is a **flat** (planar) & rigid molecule
- It has <u>4 cyclic groups</u> known as pyrrole rings with a Ferrous ion (Fe<sup>+2</sup>) in the middle
  - $\checkmark$  Every pyrrole ring has a **nitrogen**, all the <u>4 nitrogen binds with the **Iron**</u>
  - ✓ Every ring has **2 side chains** that are exposed to the outside (methyl/vinyl/propionate)
- ▶ Heme is synthesized by several steps  $\rightarrow$  having different types of porphyrins  $\rightarrow$  until getting heme
- Heme is also called Fe-Protoporphyrin IX
- > Upon absorption of light, heme gives a deep red color

#### • Iron

- It presents in the Fe<sup>+2</sup> (ferrous) state <u>not</u> in the ferric state (Fe<sup>+3</sup>)
- ▶ If it is in the ferric state, we call it Hemin  $\rightarrow$  Incapable of normal O<sub>2</sub> binding  $\rightarrow$  No O<sub>2</sub> transport
- ➤ Fe in the ferrous state (Fe<sup>+2</sup>) can form 6 bonds:
  - ✓ <u>4 with the nitrogen</u> of the pyrrole rings
    - ✓ <u>1 with N of the imidazole in Proximal His</u> → 5<sup>th</sup> coordination
    - ✓ <u>1 with  $O_2$ </u> → 6<sup>th</sup> coordination
    - One (known as the fifth coordinate) with the nitrogen of a histidine imidazole (known as proximal His)
    - ✓ One with O2 (the sixth coordinate)

# Functions of myoglobin and hemoglobin

- Myoglobin → storage of O2 in <u>muscles</u>
  During periods of oxygen <u>deprivation</u>, oxymyoglobin releases its bound oxygen
- Hemoglobin → transport of O2 and CO2 & blood buffering (Due to His residues)
  → Hemoglobin transports O2 to cells to use it in Cellular respiration and combustion reactions

## Myoglobin

- A monomeric protein (1 subunit) has a heme group (interior) → found in the muscle tissue
- The tertiary structure (overall structure)  $\rightarrow 8 \alpha$ -Helices (designated from A-H) connected by short non-helical regions
- It can present in 2 forms:
  - ➤ Oxymyoglobin → Oxygen-bound form
  - ➤ Deoxymyoglobin → Oxygen-free form (not bound to oxygen)
- In Myoglobin & other polar other globular protein, amino acids with polar R-groups are exposed on the surface (hydrophilic), while those in the interior are predominantly hydrophobic
  - - ✓ E7 → The 7<sup>th</sup> residue in Helix E / F8 → the 8<sup>th</sup> residue in Helix F
    - ✓ F8 Histidine → is also called proximal His → binds to iron
    - ✓ E7 Histidine → is also called Distal His → It is a gate that opens and closes as  $O_2$  enters the hydrophobic pocket to bind to the iron of heme → then distal His forms H-bond to  $O_2$





Hemoglobin is an allosteric protein → have many shapes (conformations) → Have 2 forms T & R states

- So, proximal His binds always to Fe, and the distal His stabilizes the interaction (O<sub>2</sub>-Fe) by H-bonds
- In low pO<sub>2</sub> or smoking conditions → CO will bind to Fe (Mb:CO complex) <u>irreversibly</u> preventing the binding of O<sub>2</sub>
- Myoglobin-heme interaction is stabilized by hydrophobic attractions between the hydrophobic pocket of the protein with Heme
  - > The heme group <u>stabilizes the tertiary structure</u> of myoglobin
- The **hydrophobic region** surrounding the Heme group **prevents** the oxidation of iron from  $\underline{Fe^{+2} \text{ to } Fe^{+3}}$ so when O<sub>2</sub> is released, the iron remains in the Fe(II) state and can bind another O<sub>2</sub>
- Myoglobin binds  $O_2$  with <u>high affinity</u>  $\rightarrow$  so can bind to  $O_2$  in very low concentration (sensitive)
- **P50:** oxygen partial pressure (indicates concentration) required for <u>50% of all</u> <u>myoglobin molecules to be saturated</u>
- P50 for myoglobin ~2.8 torrs or mm Hg
- Given that **O2 pressure in tissues** is normally <u>20 mm Hg</u>, it is almost fully saturated with oxygen at normal conditions
- The Mb-O<sub>2</sub> follows a hyperbolic saturation curve

### Hemoglobin

- Hemoglobin is a hetero-tetramer that is made of 4 subunits (2 alpha, 2 beta)
- Each subunit consists of multiple  $\alpha$ -helices ( $\alpha$  subunits have 7 helices &  $\beta$  subunits have 8 helices), with a **heme group** in the interior of the protein with Fe<sup>+2</sup>
- The chains (subunits) interact with each other via hydrophobic interactions
  - Therefore, hydrophobic amino acids are <u>not only present in the interior</u> of the protein chains, but also on the surface
- Electrostatic interactions (salt bridges) and hydrogen bonds also exist between the 2 different chains
- Hemoglobin must bind oxygen efficiently and become saturated at the <u>high oxygen pressure found in</u> <u>lungs</u> (approximately 100 mm Hg) → forming Oxyhemoglobin (Oxygen-bound)
- Then, it releases oxygen and become unsaturated in tissues where the oxygen pressure is low (about 20 mm Hg) → Deoxyhemoglobin (Oxygen-Free)
- The saturation curve:
  - > It has a sigmoidal shape
    - ✓ Each Hemoglobin bind to 4 O₂ but myoglobin binds only to 1 O₂ so it takes longer time for hemoglobin to be saturated



- ➤ At 100 mm Hg (in the lungs) → High affinity state → hemoglobin is 95-98% saturated (oxyhemoglobin)
- ➤ As the oxygen pressure falls in the tissues → Low Affinity state → oxygen is released → unsaturated
- **p50** of hemoglobin is approximately **26** mm Hg (P50 of myoglobin = 2.8)
  - $\checkmark$  That indicates that myoglobin is <u>more sensitive</u> for oxygen than Hemoglobin







50 75 100

Myoglobin

Hemoglobin

1.0

0.8

0.6

0.2

0.0 L

saturation

- Allosteric protein: A protein where binding of a molecule (ligand) to one part of the protein affects binding of a similar or a different ligand to another part of the protein
- The T-state is also known as the "taut" or "tense" state and it has a low-binding affinity to oxygen
  - The **R-state** is known as the "relaxed" state and it has 500 times **higher affinity** to oxygen than as the T
    - Binding of O<sub>2</sub> causes conformational changes in hemoglobin, converting it from the low affinity Tstate to the high affinity R- state
    - > In the T-state the 2  $\alpha\beta$  dimers are bound to each other by **a lot of interactions (salt bridges)**
    - ▶ The binding of O2 → less interactions between dimers → R-state

# • How does the structure change by the binding of O<sub>2</sub>?

## 1) The structure becomes flat pulling the proximal His

- > When heme is free of oxygen, it has a **domed** structure and iron is outside the plane of the heme
- When oxygen binds to an iron atom, heme adopts a **planar** structure and the iron moves into the plane of the heme pulling proximal histidine (F8) along with it

## 2) Breakage of the electrostatic bonds

➤ This movement triggers changes in tertiary structure of individual hemoglobin subunits → breaking the electrostatic interactions at the other oxygen-free hemoglobin chains



- > In myoglobin, movement of the helix does not affect the function of the protein
- So, the sigmoidal curve is caused by the gradual binding of O<sub>2</sub> → due to the cooperativity between the subunits of hemoglobin (allosteric)
- The affinity state (T or R)  $\rightarrow$  depends on the pressure of O<sub>2</sub> in the environment

## • This is a protective mechanism

- A sudden drop in pulmonary capillary oxygen tension does not affect hemoglobin saturation (such in high altitudes), for example:
  - ✓ If pO₂ became 75 or 80 instead of 100 it would remain fully saturated (approximately)
  - ✓ If pO2 became 60 or 50 → it would have a higher effect on the saturation it would be about (60% saturated)

