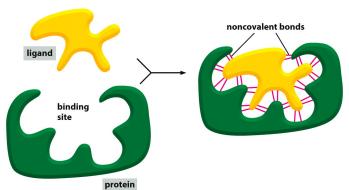
Protein Binding: Ligands, Affinity and Specificity

James Hebda 9/30/11

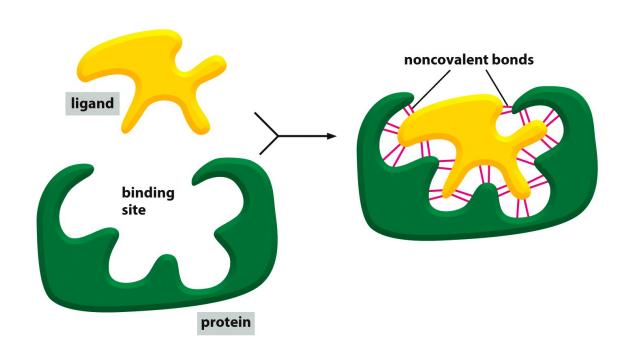


Key terms

Ligand

Affinity

Specificity



Proteins don't act alone

Myglobin binds Heme and O₂

Hemoglobin?

- Proteins act on things
 - transport and catalysis
- Or work together to make things
 - Structural proteins like microtubules and collagen

What can be a protein's ligand?

- Small molecules
 - For example O₂ in Hb and Mb

- Larger molecules
 - Complex sugar
- Other proteins or even more copies of itself

What governs protein-ligand interaction?

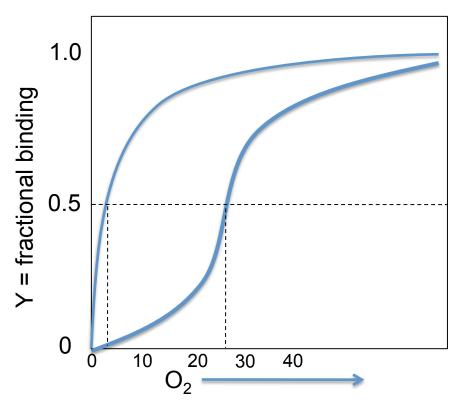
 What types of bonds determine the final structure of a protein?

 These same interactions determine how strongly a protein can hold onto its ligand.

The number and specific location of interactions

Mb and Hb

- Ligand: 0₂
- Affinity: Kd
 - Low Kd, high affinity
 - High Kd, low affinity



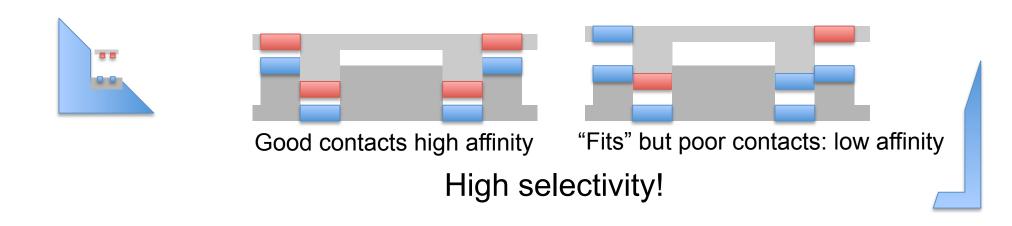
 Specificity: The distal histidine helps select for O₂ over CO

Strong binding when...

The protein and ligand fit together



The chemical contacts are right



Covalent vs Noncovalent bonds

		STRENGTH IN kcal/mole	
BOND TYPE	LENGTH (nm)	IN VACUUM	IN WATER
Covalent	0.15	90 (377)**	90 (377)
Noncovalent: ionic bond*	0.25	80 (335)	3 (12.6)
hydrogen bond	0.30	4 (16.7)	1 (4.2)
van der Waals attraction (per atom)	0.35	0.1 (0.4)	0.1 (0.4)

^{*}An ionic bond is an electrostatic attraction between two fully charged atoms.

Can you draw an example of each of these?

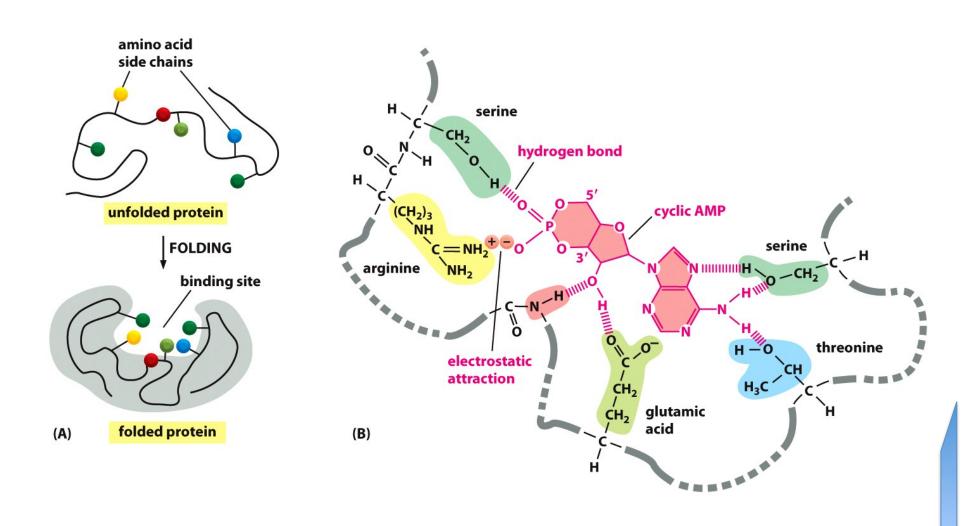
Can you easily recognize them in a figure?

^{**}Values in parentheses are kJ/mole. 1 calorie = 4.184 joules.

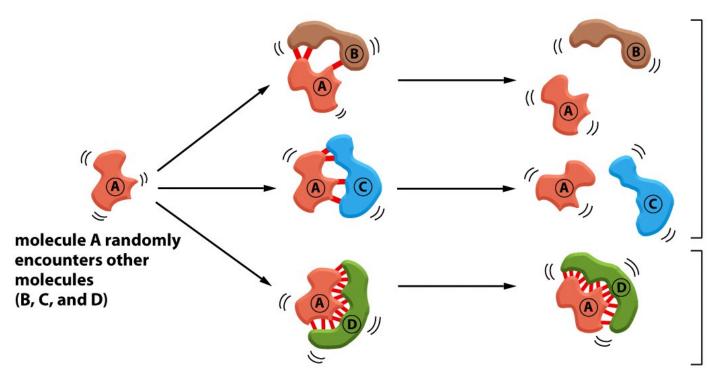
Binding affinity vs specificity

- The strength of each bond adds to the binding affinity
 - Many noncovalent bonds can result in "tight binding"
- The more bonds formed the more specific the protein is to for ligand
 - Especially true for ionic interactions

Visualizing specificity



Specificity and Function



the surfaces of molecules A and B, and A and C, are a poor match and are capable of forming only a few weak bonds; thermal motion rapidly breaks them apart

the surfaces of molecules A and D match well and therefore can form enough weak bonds to withstand thermal jolting; they therefore stay bound to each other

What you already know, and why.

 Each protein can bind more than one different ligand at a time

Binding of a ligand can change protein structure

 A change in protein structure can be passed along by non-covalent bonds

What would happen?

- If a lysozyme increased sugar affinity at the cost of specificity?
 - Lysozyme breaks down complex sugars that make up bacterial cell walls.

A fun example

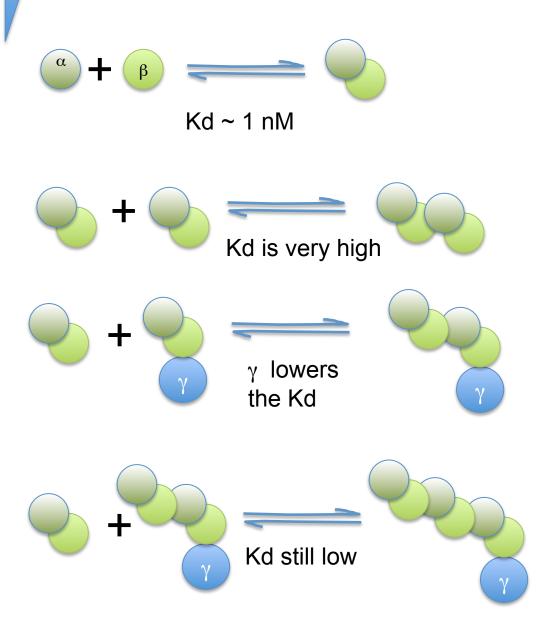
Microtubules

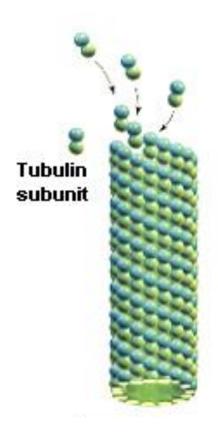
Part of the cytoskeletal network

Divide chromosomes during mitosis

Large fibers made up of dimers of two related proteins

A fun example





Gartner and Hiatt, 1994

Dynamic Instability

Tubulin monomers have a ligand

 Over time, Tubulin in the microtubules hydrolyze GTP $-GTP \rightarrow H_2PO_4 +$

$$-GTP \rightarrow H_2PO_4 +$$

 This causes a structural rearrangement that lowers the stability of dimer-dimer contacts

What happens?

• To the board.

- Inner Life of the Cell
 - Worth watching in total, but just a quick clip of microtubule polymerization and depolymerization for now

Summary

Proteins interact with many things

- Which protein interacts with what?
 - The shape and chemical nature of each protein's surface determines specificity

- A protein's affinity for a ligand is based on the summation of non-covalent bonds
 - Low Kd, High affinity