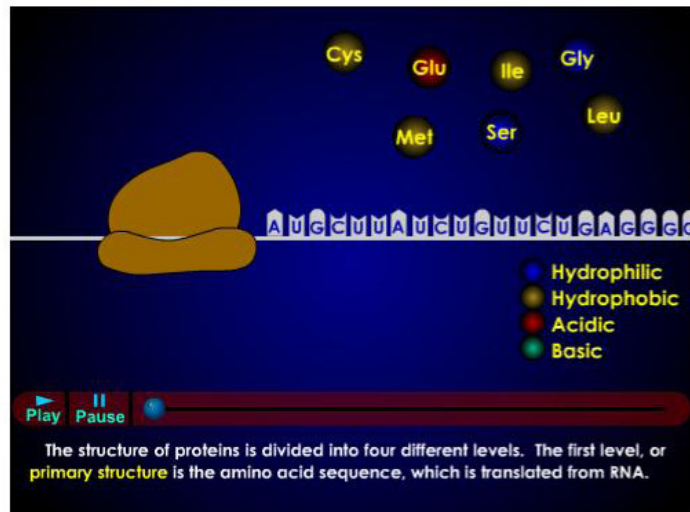
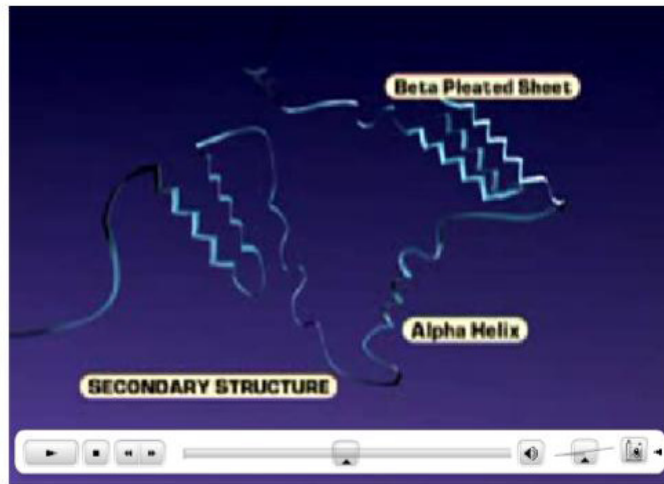


Proteins (7.5 & C.1)

# Protein folding and structure animations:

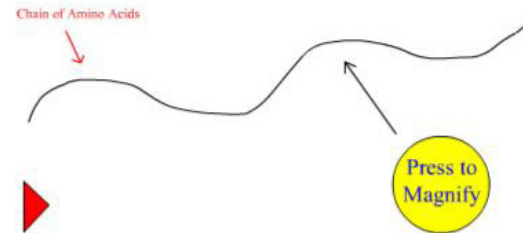


<https://mywebspace.wisc.edu/jonovic/web/proteins/Proteins.swf>

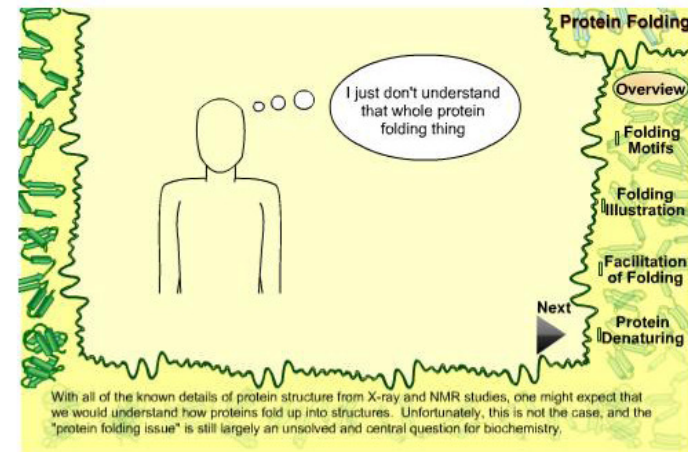


<http://www.youtube.com/watch?v=lijQ3a8yUYQ>

## Primary Structure



<http://www.stolaf.edu/people/giannini/flashanimat/proteins/protein%20structure.swf>



[http://www.wiley.com/legacy/college/boyer/0470003790/animations/protein\\_folding/protein\\_folding.swf](http://www.wiley.com/legacy/college/boyer/0470003790/animations/protein_folding/protein_folding.swf)

- <http://www.youtube.com/watch?v=lijQ3a8yUYQ>
- <http://www.stolaf.edu/people/giannini/flashanimat/proteins/protein%20structure.swf>
- <http://www.wiley.com/legacy/college/boyer/0470003790/>
- <http://www.stolaf.edu/people/giannini/flashanimat/proteins/hydrophobic%20force.swf>

Proteins are the stable, folded 3D structure of polypeptides.

Polypeptides are the chain of amino acids that fold into proteins.

There are **four levels of protein structure**.

### Primary (1°)

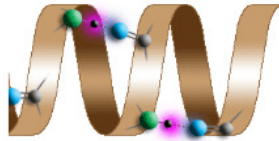
- amino acid sequence
- peptide bonds



### Secondary (2°)

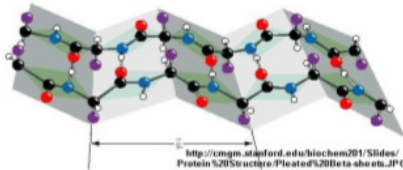
- repeating local structures
- held by H-bonds

$\alpha$ -helix



[http://www.brooklyn.cuny.edu/bc/ahp/LAD/C4b/C4b\\_proteinShape.html](http://www.brooklyn.cuny.edu/bc/ahp/LAD/C4b/C4b_proteinShape.html)

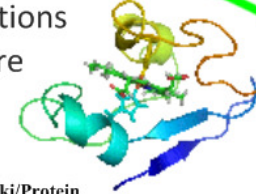
$\beta$ -pleated sheet



<http://cmgm.stapford.edu/biochem201/Slides/Protein%20Structure/Pleated%20Sheet%20to%20the%20em.JPG>

### Tertiary (3°)

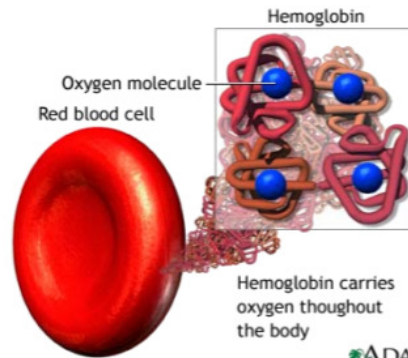
- folding of a single protein
- R-group interactions
- hydrophobic core



<http://en.wikipedia.org/wiki/Protein>

### Quaternary (4°)

- protein complex
- made of two or more subunits

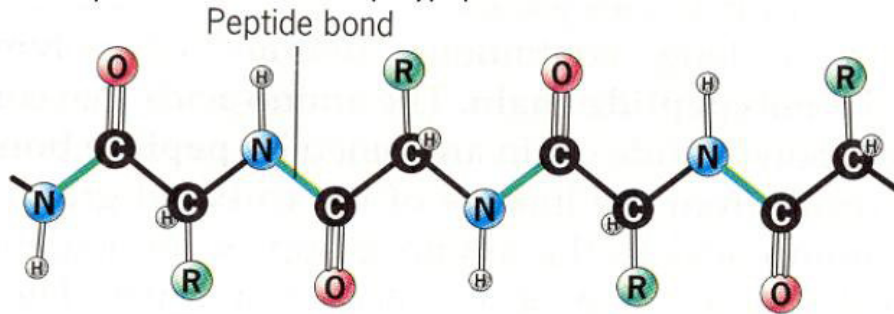


Hemoglobin carries oxygen throughout the body

ADAM

# Primary (1°) Structure of proteins: The polypeptide

How many amino acids in this polypeptide?



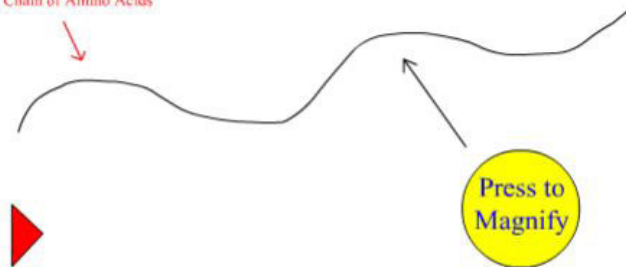
<http://www.bio.miami.edu/~cmallery/255/255amino/255aminoacids.htm>

- Amino acid sequence
- Peptide bonds



## Primary Structure

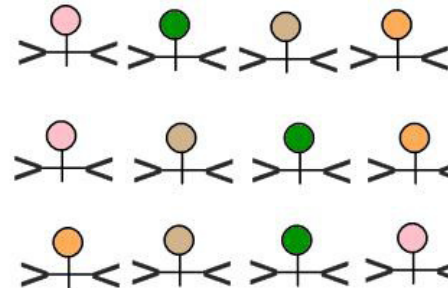
Chain of Amino Acids



<http://www.stolaf.edu/people/giannini/flashanimat/proteins/protein%20structure.swf>

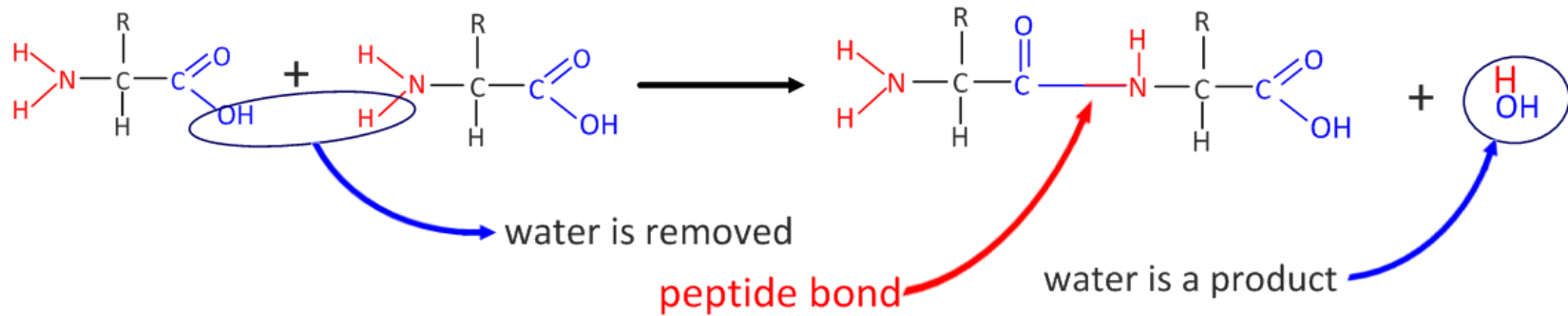
Why are there **infinite possibilities** of **polypeptides**?

- 20 different amino acids
- could be any length
- amino acids can be in any order



} different properties

# Polypeptides are chains of amino acids joined by peptide bonds:



<p>Arginine (Arg / R)</p>	<p>Glutamine (Gln / Q)</p>	<p>Phenylalanine (Phe / F)</p>	<p>Tyrosine (Tyr / Y)</p>	<p>Tryptophan (Trp / W)</p>
<p>Lysine (Lys / K)</p>	<p>Glycine (Gly / G)</p>	<p>Alanine (Ala / A)</p>	<p>Histidine (His / H)</p>	<p>Serine (Ser / S)</p>
<p>Proline (Pro / P)</p>	<p>Glutamic Acid (Glu / E)</p>	<p>Aspartic Acid (Asp / D)</p>	<p>Threonine (Thr / T)</p>	<p>Cysteine (Cys / C)</p>
<p>Methionine (Met / M)</p>	<p>Leucine (Leu / L)</p>	<p>Asparagine (Asn / N)</p>	<p>Isoleucine (Ile / I)</p>	<p>Valine (Val / V)</p>

There are 20 different amino acids.

These can be combined in any order.

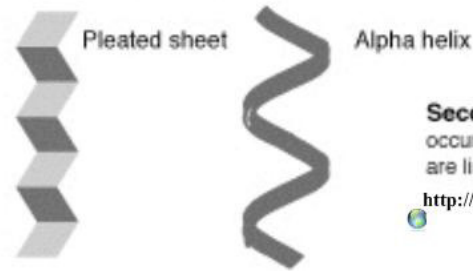
Each amino acid has unique properties:

- some are polar (hydrophilic)
- some are non-polar (hydrophobic)
- some are positively or negatively charged
- some contain sulphur

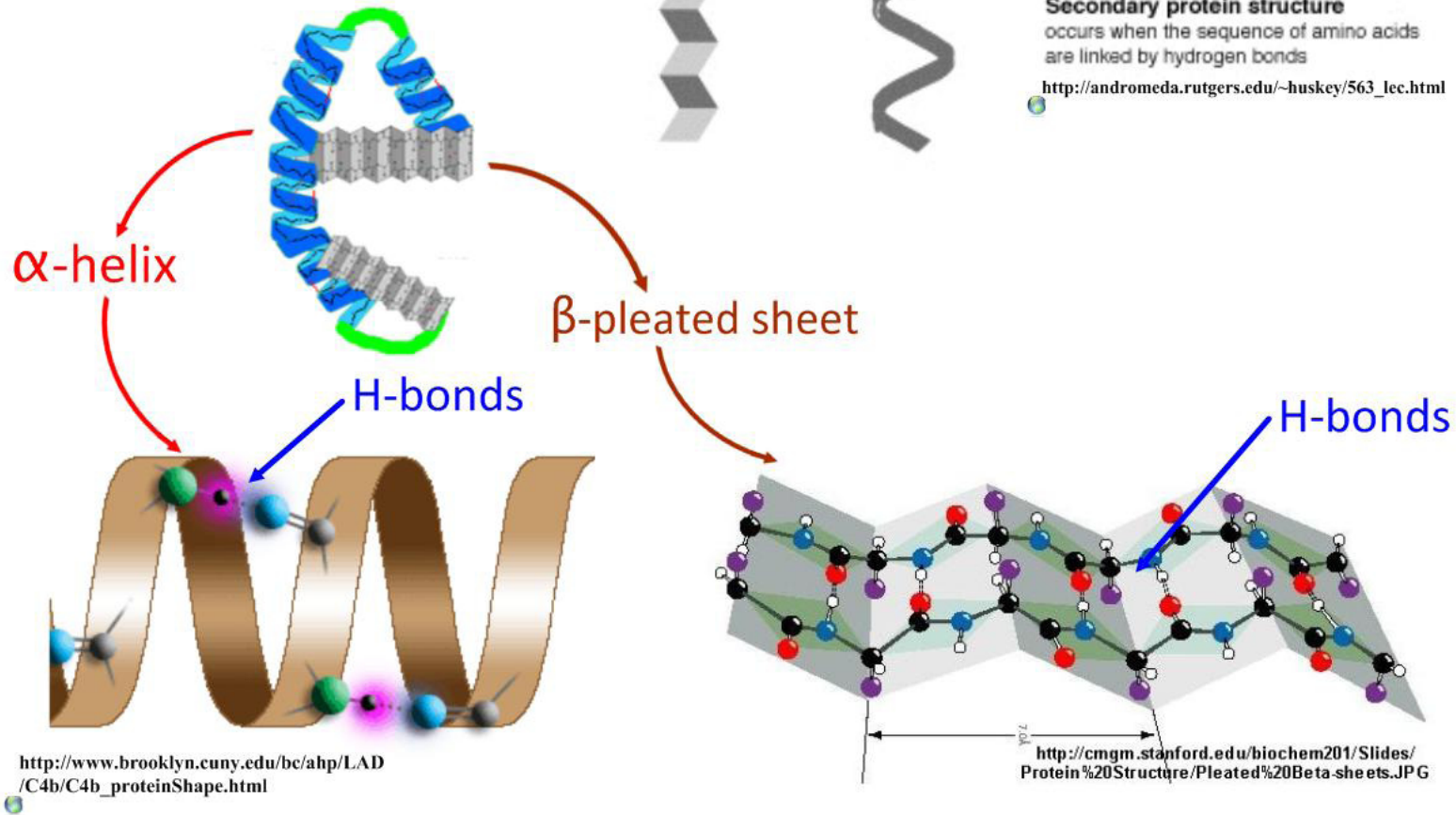
The properties of the amino acids determine how a polypeptide folds up into a protein.

# Secondary (2<sup>o</sup>) structure: Repeating Units

- Repeating local structures
- Held by H-bonds



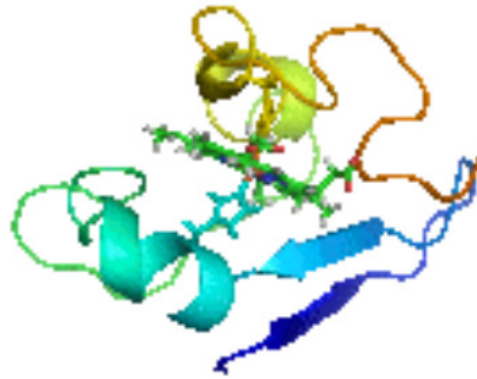
**Secondary protein structure**  
occurs when the sequence of amino acids  
are linked by hydrogen bonds  
[http://andromeda.rutgers.edu/~huskey/563\\_lec.html](http://andromeda.rutgers.edu/~huskey/563_lec.html)



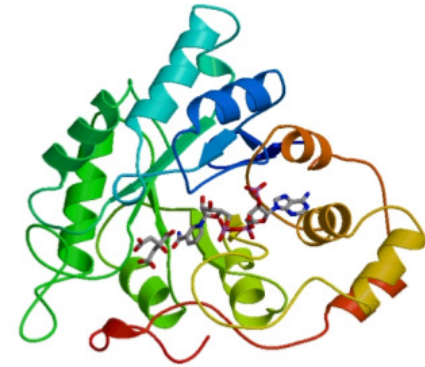
# Tertiary structure (3<sup>o</sup>): 3D Folded Structure

- Folding of a single protein into a 3D structure
- Hydrophobic core

This 3D structure gives proteins their functional properties, such as active sites on enzymes.



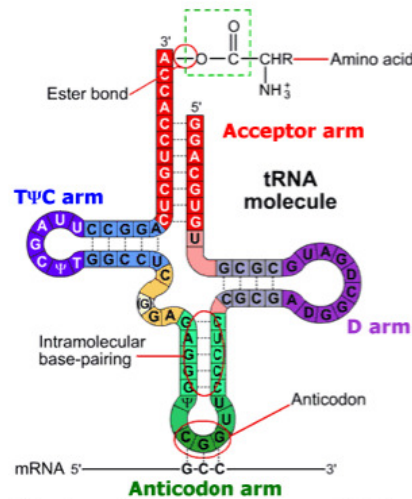
<http://en.wikipedia.org/wiki/Protein>



<http://shelx.uni-ac.gwdg.de/~fabio/endwkcon.htm>

## • R-group interactions:






- hydrophilic polar amino acids orient to the outside
- hydrophobic non-polar amino acids protected in the core
- oppositely-charged ions attract
- disulphide bridges formed between sulphur-containing amino acids



tRNA is an example of a tertiary

<http://www3.interscience.wiley.com:8100/legacy/college/boyer/0471661791/structure/tRNA/trna.htm>

# Modeling protein folding

colour					
number	11	1	1	2	2
e.g.	serine	lysine	aspartic acid	cysteine	valine
	SER	LYS	ASP	CYS	VAL
property	polar	<sup>+</sup> ve	<sup>-</sup> ve	contains S	non-polar






What is the primary structure of this section of a polypeptide?





# Modeling protein folding

try it with plasticene

colour					
number	11	1	1	2	2
e.g.	serine	lysine	aspartic acid	cysteine	valine
	SER	LYS	ASP	CYS	VAL
property	polar	<sup>+</sup> ve	<sup>-</sup> ve	contains S	non-polar

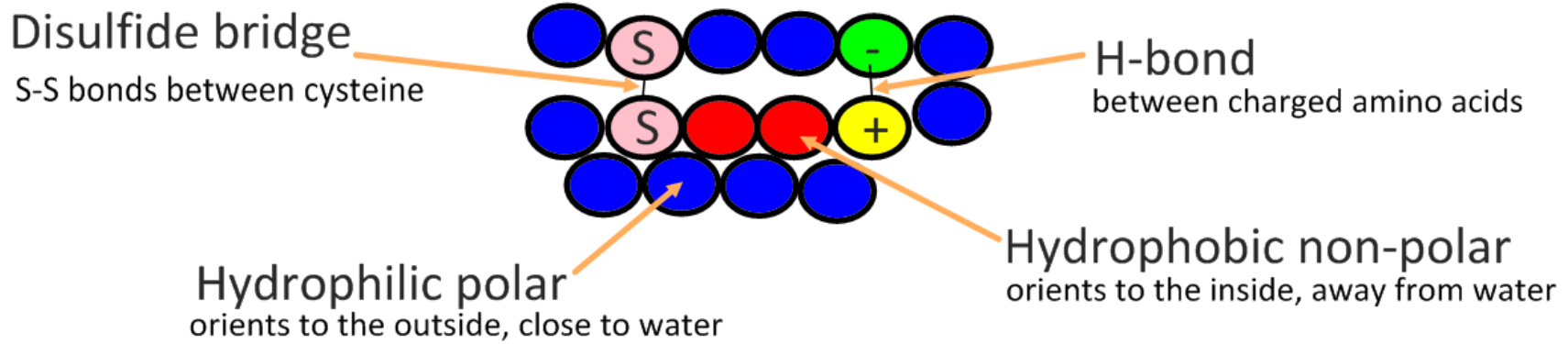
What is the primary structure of this section of a polypeptide?








How would the polypeptide fold if placed in water?

1. polar groups outside
2. non-polar groups inside
3. positive near negative
4. disulfide bridges

# Modeling protein folding:



colour					
number	11	1	1	2	2
e.g.	serine	lysine	aspartic acid	cysteine	valine
	SER	LYS	ASP	CYS	VAL
property	polar	+ <sub>ve</sub>	- <sub>ve</sub>	contains S	non-polar

# Quaternary structure (4<sup>o</sup>): Multiple Subunits

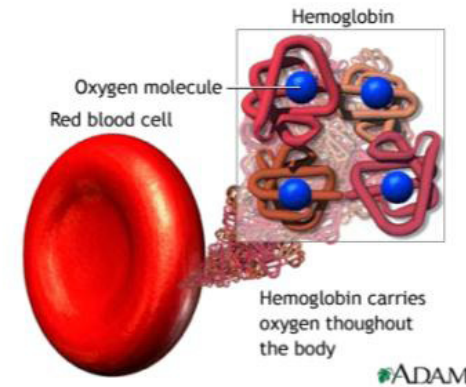
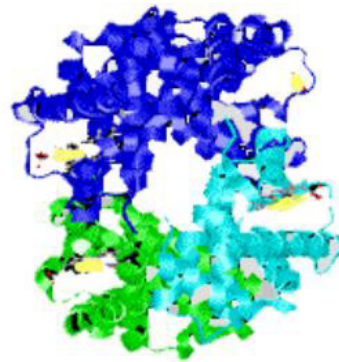
- Protein complex
- Made of two or more subunits

e.g. DNA Polymerase



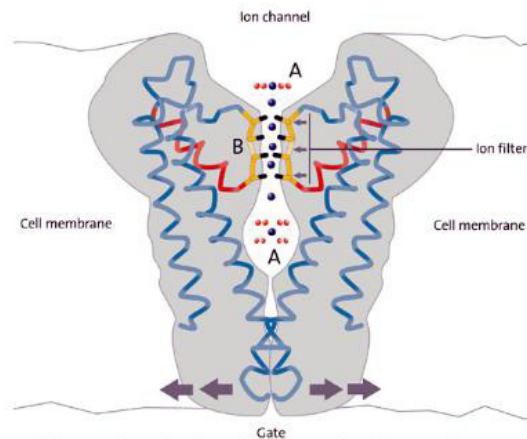
[http://en.wikipedia.org/wiki/DNA\\_polymerase](http://en.wikipedia.org/wiki/DNA_polymerase)

e.g. haemoglobin



<http://www.bmb.uga.edu/wampler/tutorial/prot4.html>

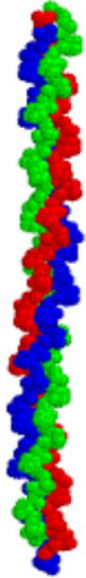
e.g. Ion channels in the plasma membrane



[http://www.bio.miami.edu/~cmallery/150/memb/ion\\_channels.htm](http://www.bio.miami.edu/~cmallery/150/memb/ion_channels.htm)

- The heme is a small but important non-protein molecule, or prosthetic **group**, that binds an **iron** atom in the ferrous (+2) oxidation state.

## Fibrous Proteins



insoluble in water

structural  
(support, strength)

keratin (hair/nails)

elastin (skin)

collagen (tissue  
strengthening)

## Globular Proteins

can be soluble

functional  
(enzymes, antibodies)

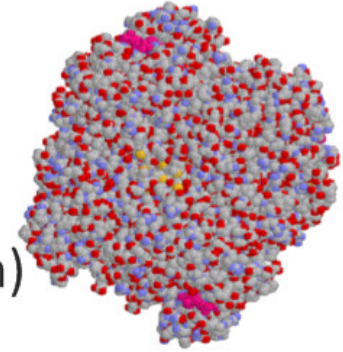
amylase (digests starch)

Insulin (blood sugar regulation)

haemoglobin

immunoglobulins (antibodies)

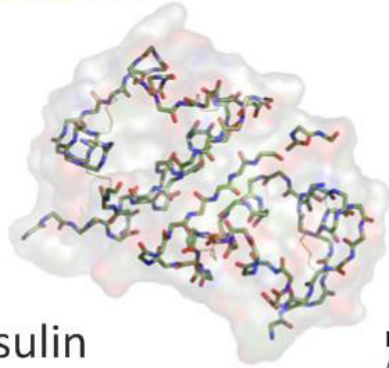
transport ( $\text{Na}^+/\text{K}^+$  pump)



# Uses of proteins:

Proteins can be structural or part of the plasma membrane. They can also perform other diverse functions in an organism:

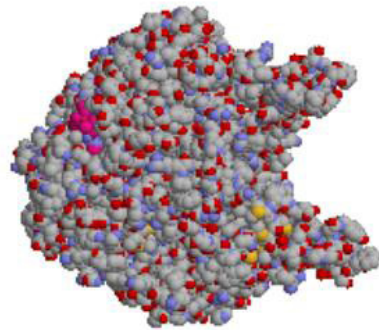
## Hormones



insulin

<http://www.3dchem.com/imagesofmolecules/Insulin2.jpg>

## Enzymes

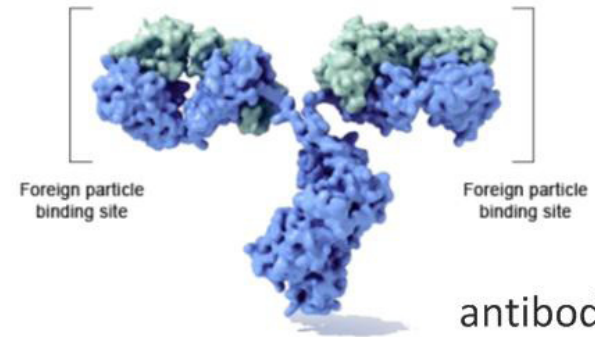


<http://www.steve.gb.com/images/molecules/proteins/catalase.gif>

catalase

Immunoglobulin G (IgG)

## Immunoglobulin



antibodies

<http://ghr.nlm.nih.gov/handbook/illustrations/igg.jpg>

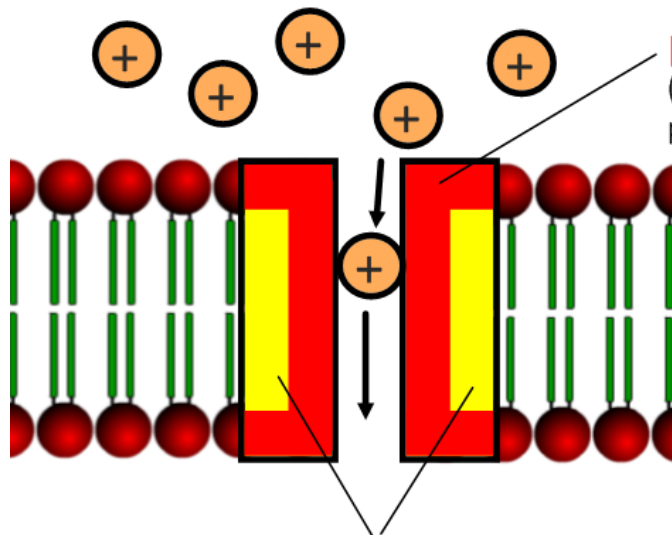
## Gas transport

haemoglobin



<http://chemistry.ewu.edu/jcorkill/biochem/48006.htm>

Polar and non-polar amino acids are used in **membrane channels**:

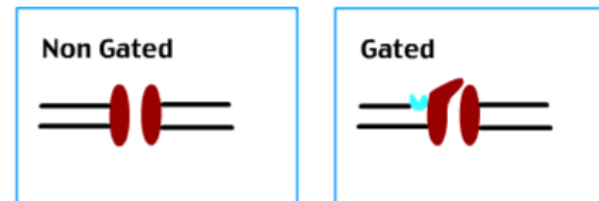


**polar sections** allow ions through  
(charged or polar molecules cannot normally pass through the hydrophobic layers)

**non-polar** sections bond with hydrophobic tails

**polar amino acids** are positioned on external surfaces and line the **protein channels** for **facilitated diffusion**

## CHANNEL

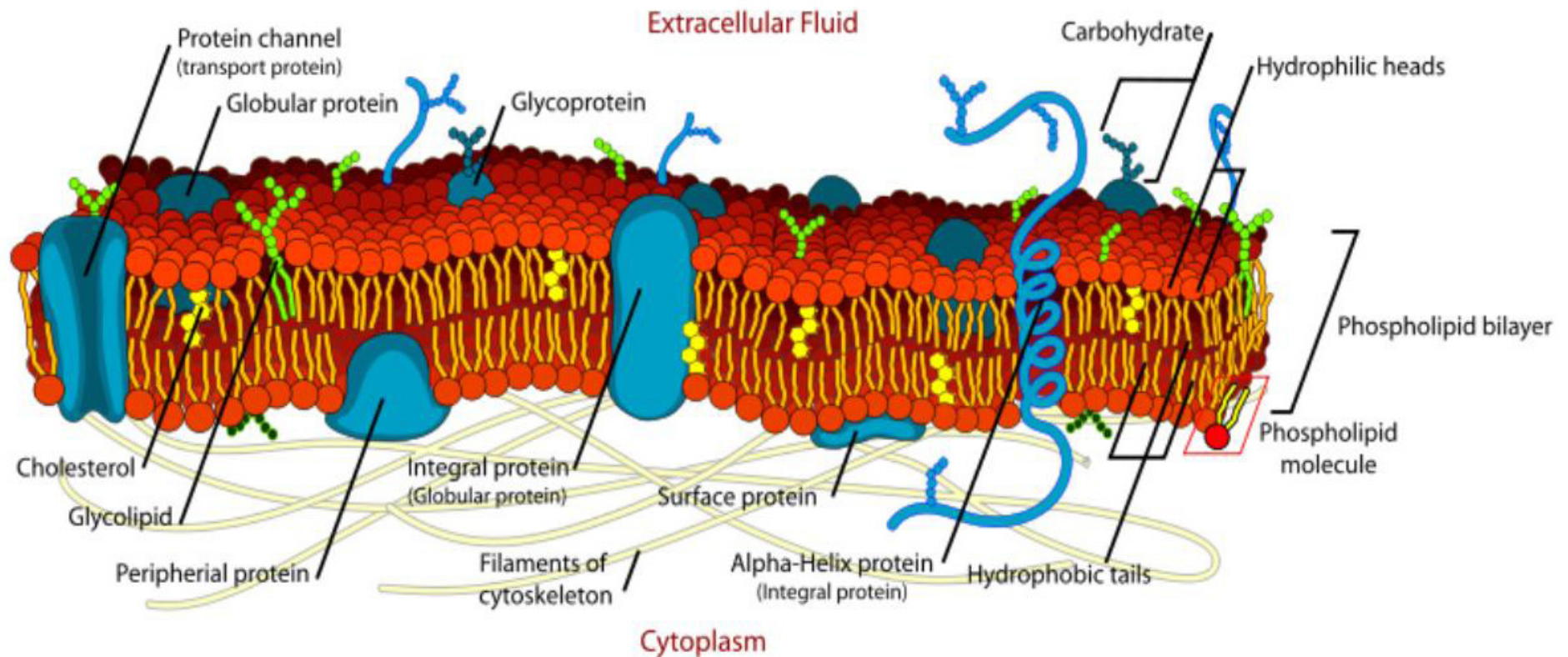


CLICK ONE BOX

<http://www.stolaf.edu/people/giannini/flashanimat/transport/channel.swf>

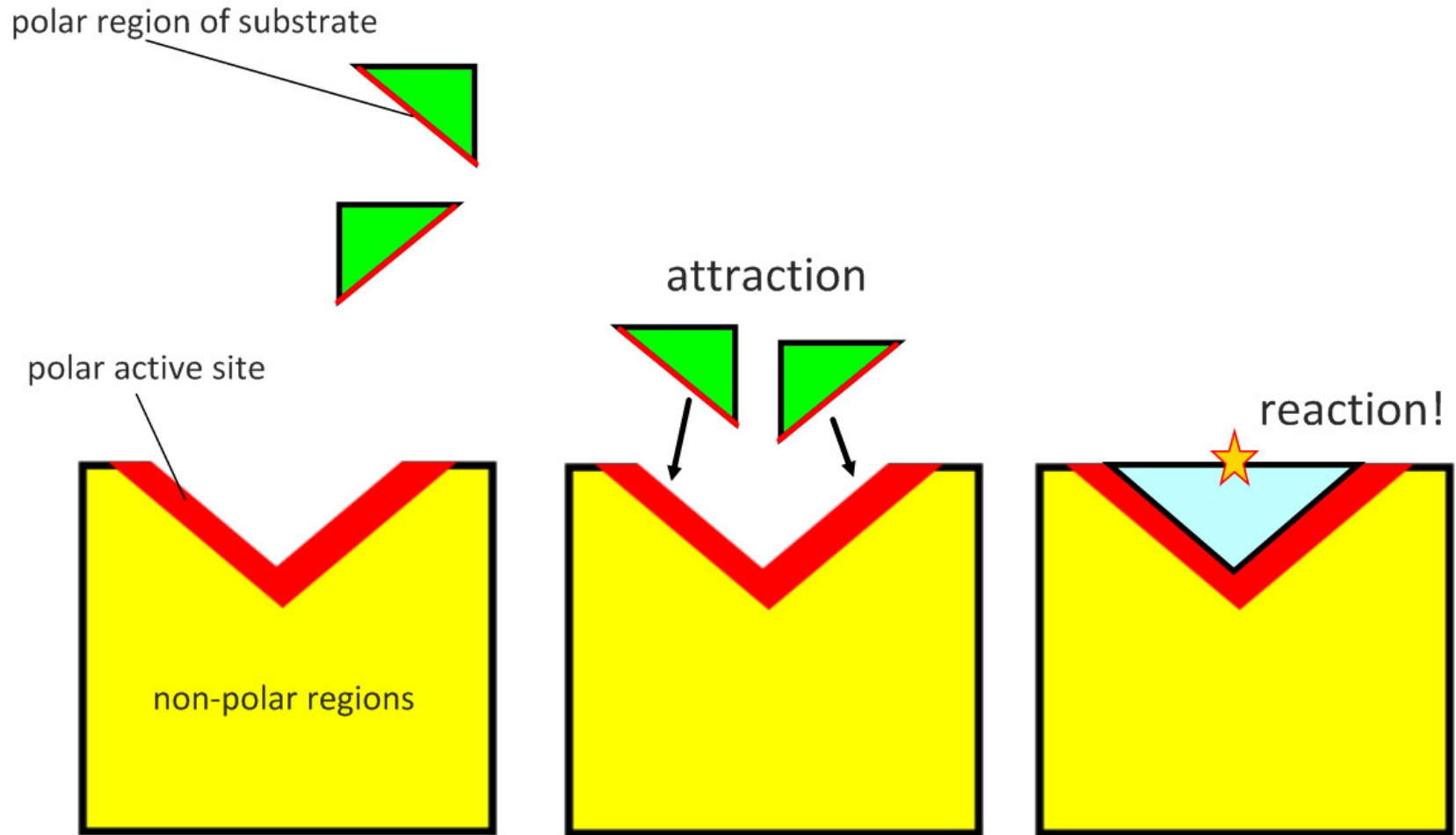
- <http://www.stolaf.edu/people/giannini/flashanimat/transport/channel.swf>

## Non-polar amino acids help bond proteins to the plasma membrane



<http://cellbiology.med.unsw.edu.au/units/science/lecture0803.htm>

Polar amino acids in enzymes help **bind the substrates for reaction:**





# Foldit: Solve puzzles for Science!



Rank: 17 Score: 9092  
48: Pro Peptide

Group Competition	
# Group Name	Score
1 The Lone Folder	9388
2 Street Smarts	9367
3 Illinois	9303
4 Berkeley	9255

Player Competition	
# Player Name	Score
16 psen	9098
17 kathleen	9092
18 versatR2	9091
19 darktorres	9081
20 ccarico	9066
21 mbojkegren	9048
22 sldickerson	9038

Shake sidechains to improve the protein. Hotkey: S

Shake Sidechains Wiggle Backbone Clear Locks and Bands Reset Puzzle Mouse Help

Actions History View File

Download and play:

GET STARTED: DOWNLOAD



Win XP/Vista



Intel OS X 10.4 or later



Linux

<http://fold.it/portal/>



You can see proteins in the protein data bank:

Can you make them in FoldIt?

<http://www.pdb.org/pdb/home/home.do>

- <http://fold.it/portal/>

