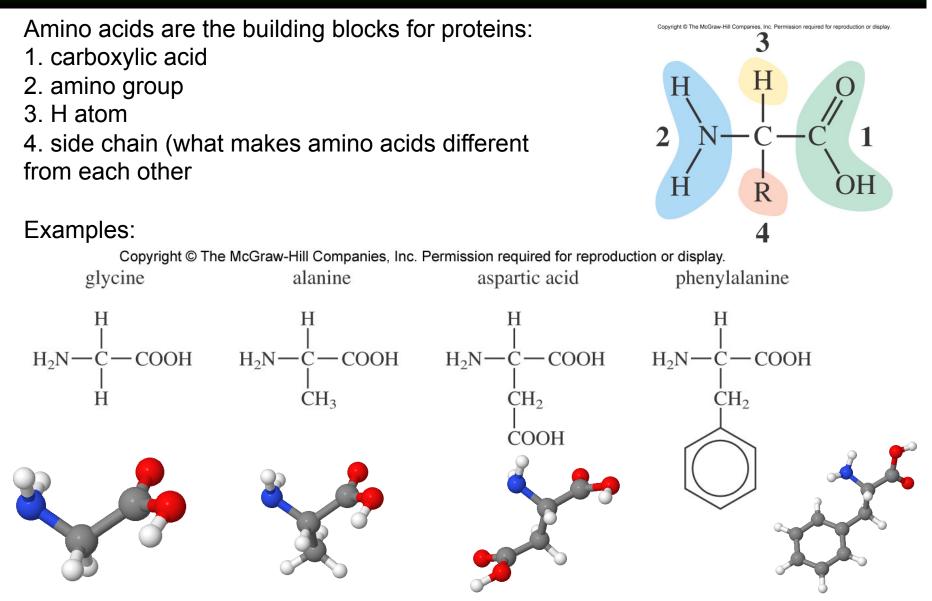
# Proteins

# Amino Acids (Ch. 11.4)



20 naturally occurring amino acids; our bodies can make only 11 of them...

# **Proteins from Amino Acids (Ch. 11.4)**

#### Yet another example of a condensation reaction:

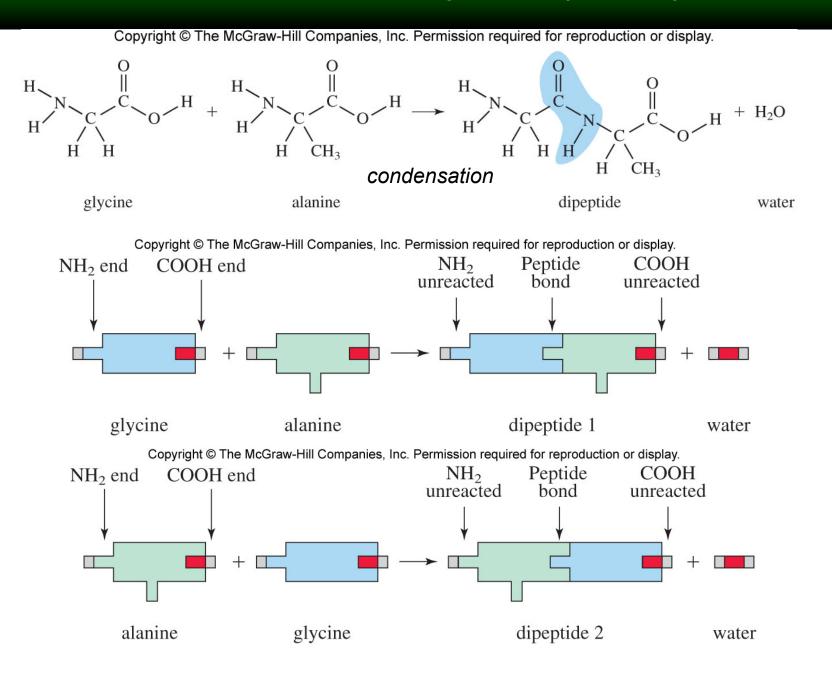
Copyright © The McGraw-Hill Companies, Inc. Permission required for reproduction or display.  $H \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{H} \xrightarrow{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{H} \xrightarrow{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{H} \xrightarrow{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{H} \xrightarrow{H} \xrightarrow{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{O}_{H} \xrightarrow{H} \xrightarrow{H} \xrightarrow{O}_{H} \xrightarrow{O}_{$ 

Two amino acids combine to form a **dipeptide**. The order matters: gly-ala is not the same molecule as ala-gly.

A protein is a polypeptide built up from 100s-1000s of amino acids.

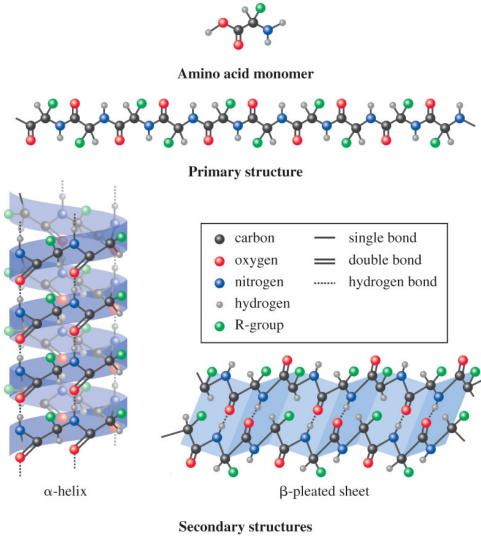
An **enzyme** is a protein that can act as a catalyst.

#### Active Sites: Protein Synthesis (Ch. 11.4)



### Active Sites: Protein Structure (Ch. 12.4)

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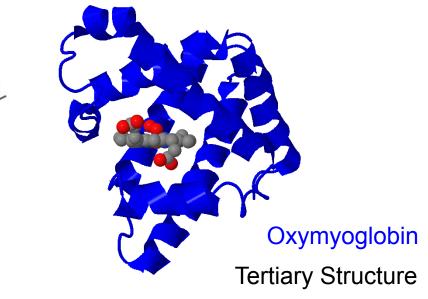


**Primary Structure**: sequence of amino acids (comes from DNA)

**Secondary Structure**: alpha helix, beta sheet, ... (close interactions)

**Tertiary Structure**: 3-D arrangement of chain (distant interactions within chains)

**Quaternary Structure**: 3-D arrangement of chains (interactions between chains)



Myoglobin contains 153 amino acids:

VAL LEU SER GLU GLY GLU TRP GLN LEU VAL LEU HIS VAL TRP ALA LYS VAL GLU ALA ASP VAL ALA GLY HIS GLY GLN ASP ILE LEU ILE ARG LEU PHE LYS SER HIS PRO GLU THR LEU GLU LYS PHE ASP ARG PHE LYS HIS LEU LYS THR GLU ALA GLU MET LYS ALA SER GLU ASP LEU LYS LYS HIS GLY VAL THR VAL LEU THR ALA LEU GLY ALA ILE LEU LYS LYS LYS GLY HIS HIS GLU ALA GLU LEU LYS PRO LEU ALA GLN SER HIS ALA THR LYS HIS LYS ILE PRO ILE LYS TYR LEU GLU PHE ILE SER GLU ALA ILE ILE HIS VAL LEU HIS SER ARG HIS PRO GLY ASP PHE GLY ALA ASP ALA GLN GLY ALA MET ASN LYS ALA LEU GLU LEU PHE ARG LYS ASP ILE ALA ALA LYS TYR LYS GLU LEU GLY TYR GINGIY

# **Primary Structure**: sequence of amino acids (comes from DNA)

**Secondary Structure**: alpha helix, beta sheet, ... (close interactions)

**Tertiary Structure**: 3-D arrangement of chain (distant interactions)

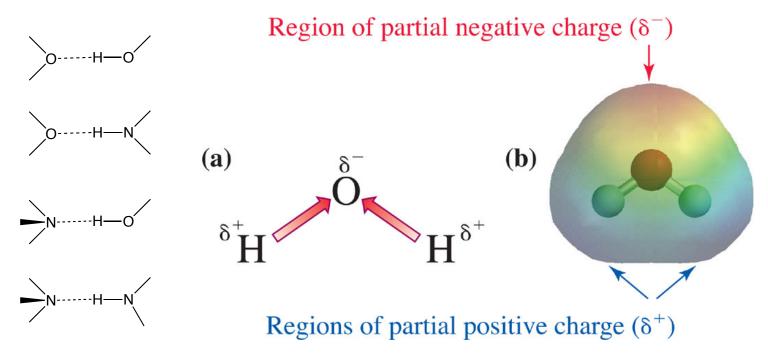
**Quaternary Structure**: 3-D arrangement of chains (interactions between chains)

# Hydrogen Bonding (5.5 & 5.6)

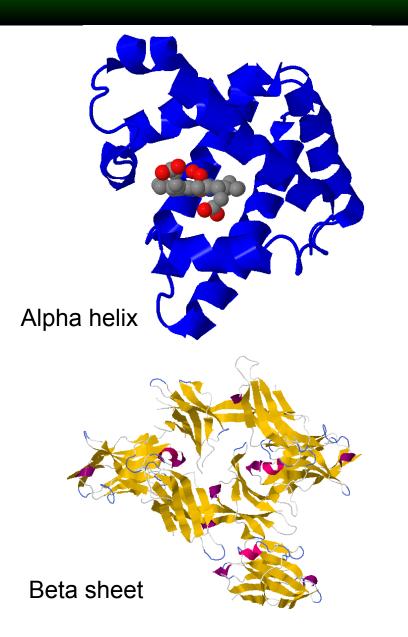
So far, we've talked about shared pairs of electrons

The sharing doesn't have to be equal later electronegativity will tell us which atoms have the stronger pull

For now: N-H and O-H bonds are polarized to make the H partially positive & the O or N partially negative



#### Secondary structure



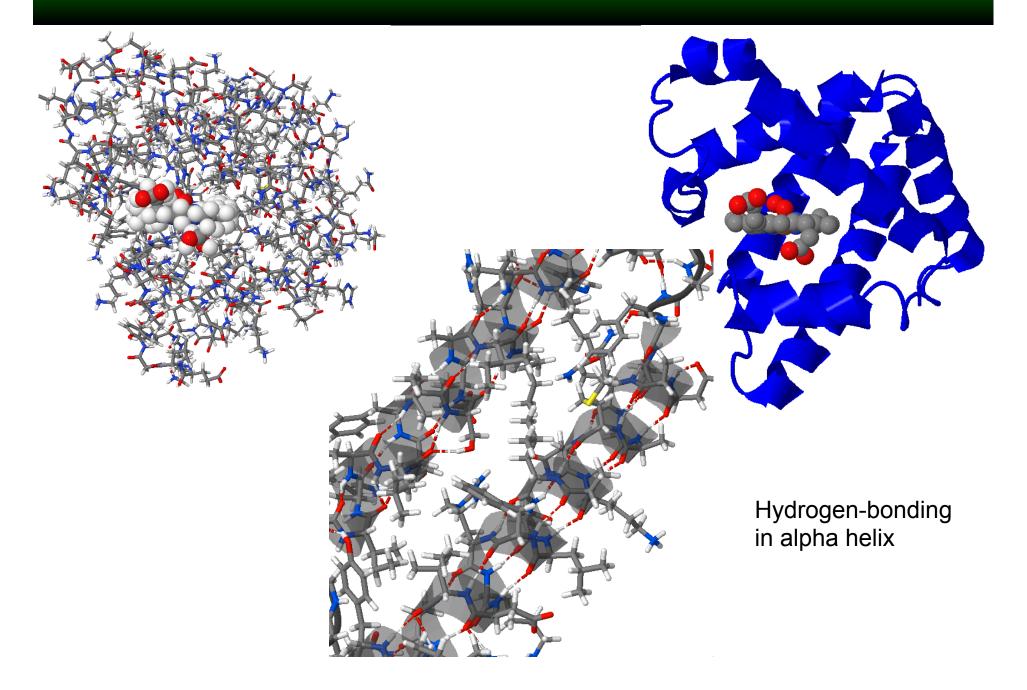
**Primary Structure**: sequence of amino acids (comes from DNA)

**Secondary Structure**: alpha helix, beta sheet, ... (close interactions)

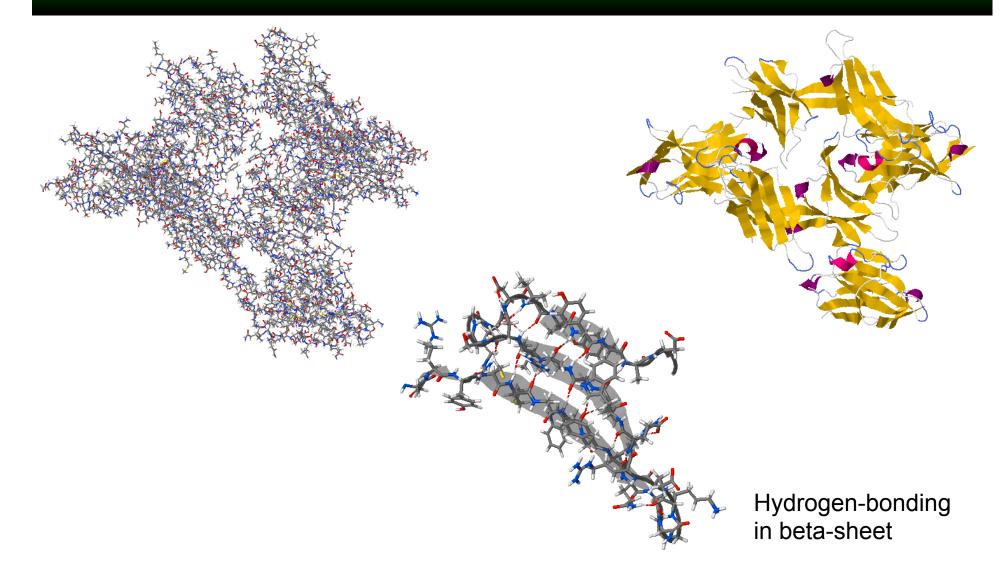
**Tertiary Structure**: 3-D arrangement of chain (distant interactions)

**Quaternary Structure**: 3-D arrangement of chains (interactions between chains)

# Myoglobin, representations



### **Beta-Sheets**



Beta2-microglobulin, thought to be involved in amyloid formation (Alzheimer's disease, Prion disease (Mad Cow))

### **Binding Sites: Quaternary Structure in Hemoglobin**

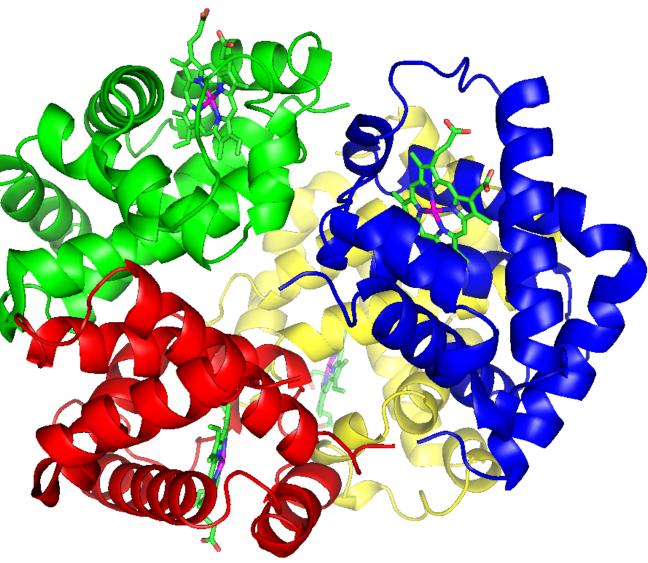
Hemoglobin

4 subunits

 $4 O_2$  binding sites

Binding of 1 molecule of  $O_2$  is slow, but causes the other 3 sites to open up

CO binds more slowly than  $O_2$ , but is more difficult to remove from the Fe-containing binding site



# **Protein Folding**

So far we've talked about the role that hydrogen bonding plays in protein structure

**Primary Structure**: sequence of amino acids (comes from DNA)

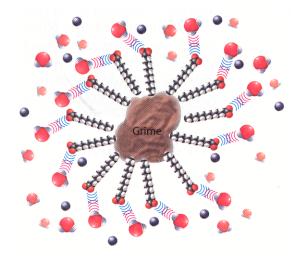
**Secondary Structure**: alpha helix, beta sheet, ... (close interactions)

**Tertiary Structure**: 3-D arrangement of chain (distant interactions within chain)

**Quaternary Structure**: 3-D arrangement of chains (interactions between chains)

Remember: General rule: Like dissolves like Hydrocarbon dissolves in Hydrocarbon C-H bonds oils, fats Nonpolar Hydrophobic

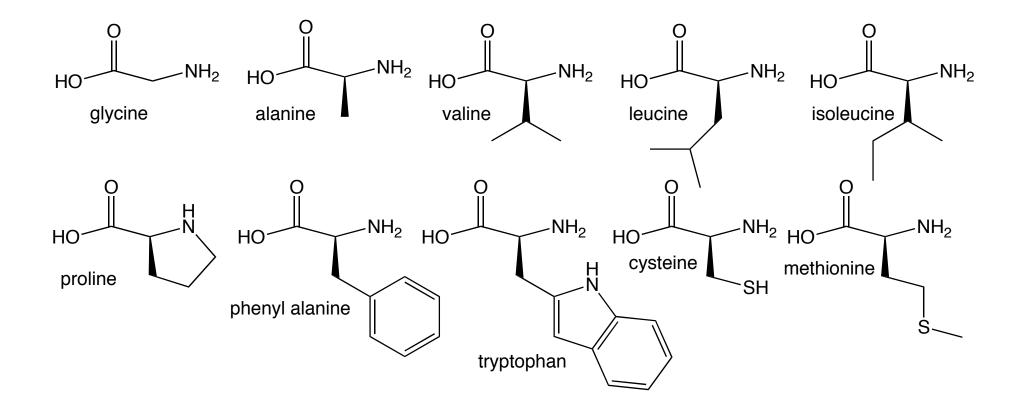
#### & how soap works



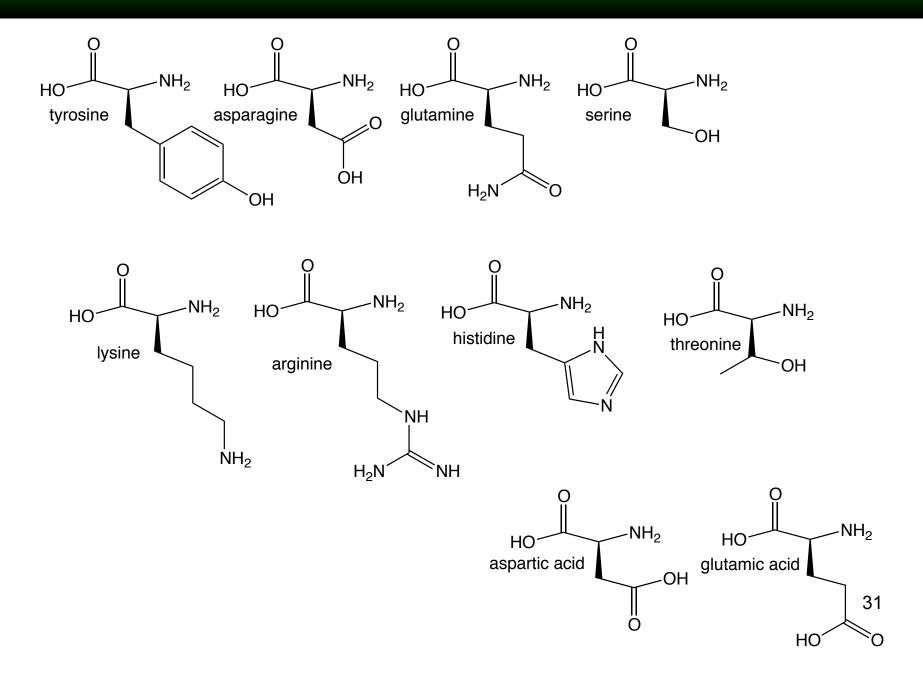
Hydroxyl dissolves in Hydroxyl O-H bonds water, alcohols, sugars Polar Hydrophilic

Some amino acid side chains are hydrophobic & some are hydrophilic

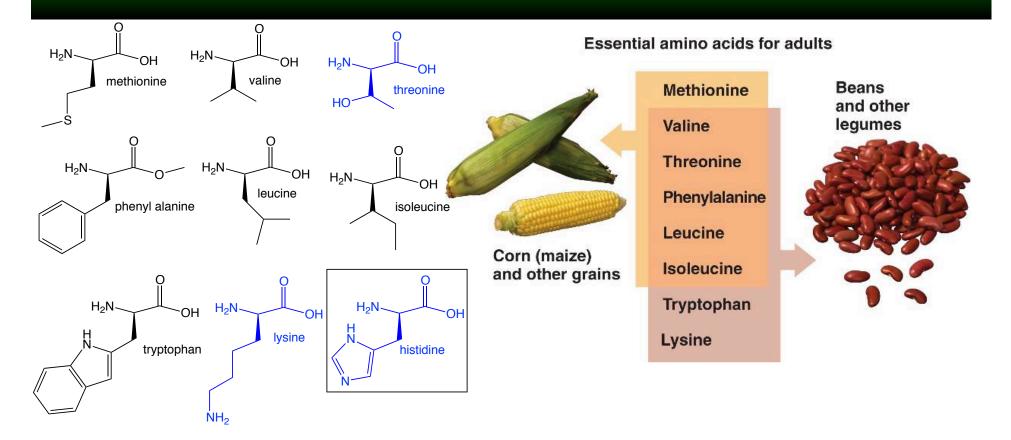
# Hydrophobic amino acids



# Hydrophilic amino acids



# **Essential amino acids**



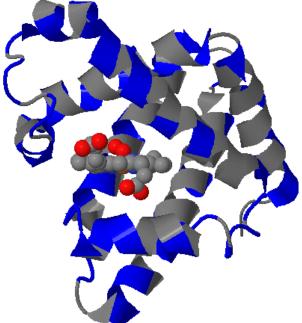
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Table 11.5	The Essential Amino Acids		
histidine		lysine	threonine
isoleucine		methionine	tryptophan
leucine		phenylalanine	valine

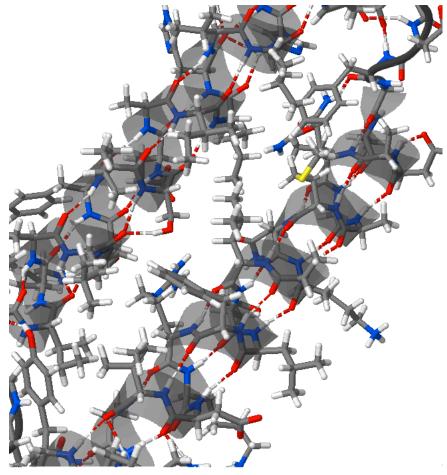
# Myoglobin—solution phase protein

Primary oxygen carrier in muscle tissue

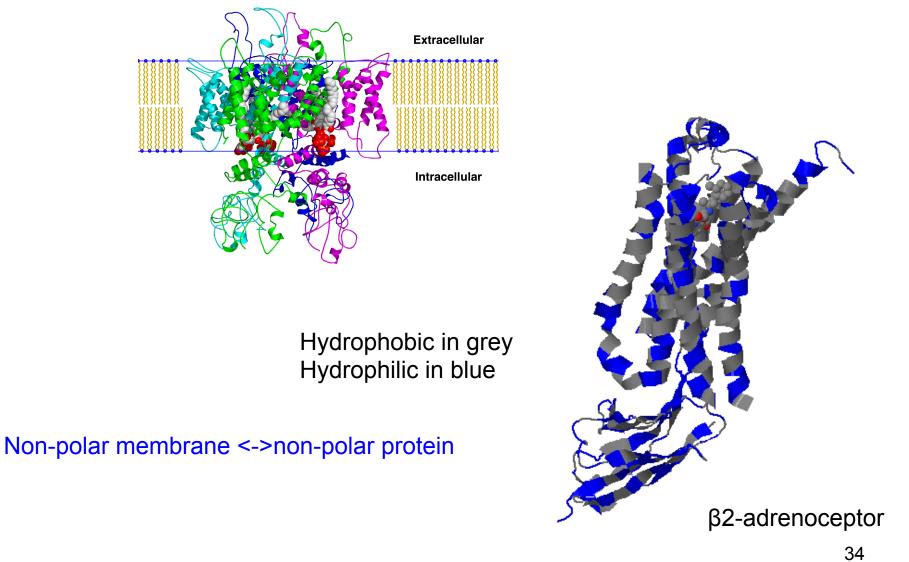
Hydrophobic in grey Hydrophilic in blue



Hydrophobic side chains are on the inside Hydrophobic interactions contribute to folding Non-polar <-> non-polar polar <-> polar



#### **Trans membrane receptor**



Receptor for adrenaline (fight-or-flight response)