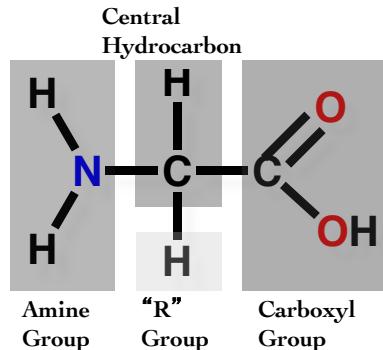
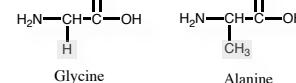
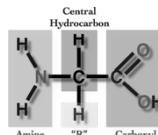


Proteins are composed of monomers called **amino acids**

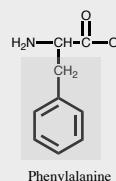
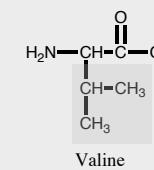
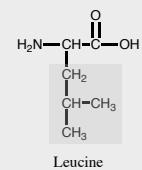


• ALL amino acids have the exact same structure *except* for the "R" Group

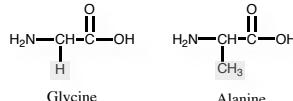
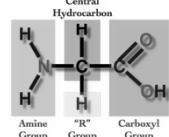
There are **20** different amino acids



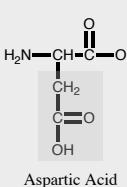
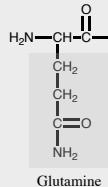
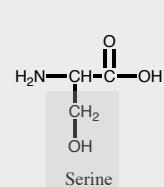
### Hydrophobic Amino Acids



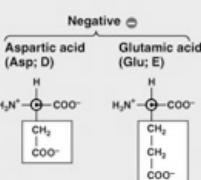
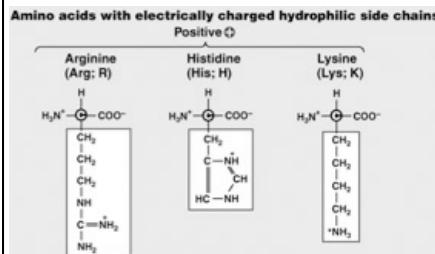
There are **20** different amino acids



### Hydrophilic Amino Acids

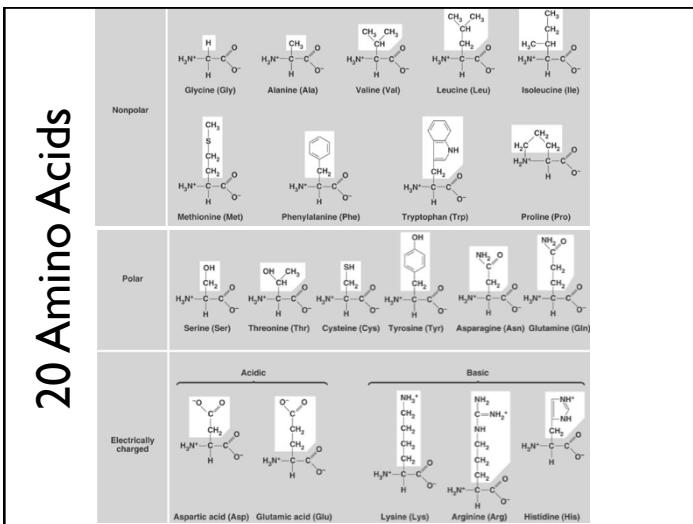


### Charged amino acids (side-chains)



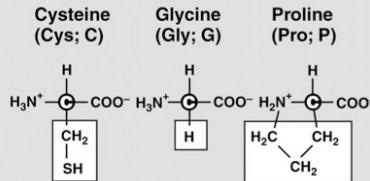
Are these polar or nonpolar side chains??

## 20 Amino Acids



## Special amino acids

### C. Special cases



Cysteine has  $-SH$  that can react with another to form disulfide bridge.  
Controls peptide folding.

Glycine- small side chain allows it to pack tightly and fit in tight corners

Proline- ring structure limits its ability to rotate and form bonds.  
Stabilizes bends and loops.

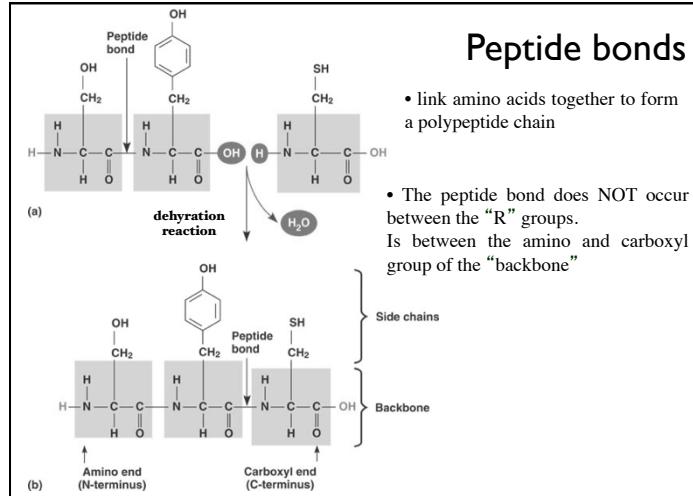
## There are 20 different amino acids

The type of R-Group will affect the 3-dimensional structure of the protein

| AMINO ACID        | SIDE CHAIN | AMINO ACID           | SIDE CHAIN |
|-------------------|------------|----------------------|------------|
| Aspartic acid     | Asp D      | negative             |            |
| Glutamic acid     | Glu E      | negative             |            |
| Arginine          | Arg R      | positive             |            |
| Lysine            | Lys K      | positive             |            |
| Histidine         | His H      | positive             |            |
| Asparagine        | Asn N      | uncharged polar      |            |
| Glutamine         | Gln Q      | uncharged polar      |            |
| Serine            | Ser S      | uncharged polar      |            |
| Threonine         | Thr T      | uncharged polar      |            |
| Tyrosine          | Tyr Y      | uncharged polar      |            |
| Polar amino acids |            | Nonpolar amino acids |            |

## Peptide bonds

- link amino acids together to form a polypeptide chain



**Peptide Bonds Form Chains (Primary Structure)**

**SCHEMATIC**

**SEQUENCE**

The **sequence** of amino acids constitutes the **primary structure** of a protein.

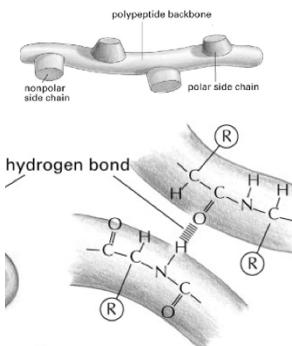
Held together by covalent peptide bonds.

**Amino end**      **Carboxyl end**

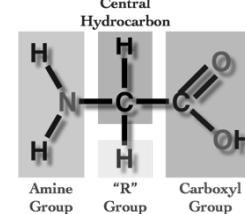
**Amino acid subunits**

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100 101 102 103 104 105 106 107 108 109 110 111 112 113 114 115 116 117 118 119 120 121 122 123

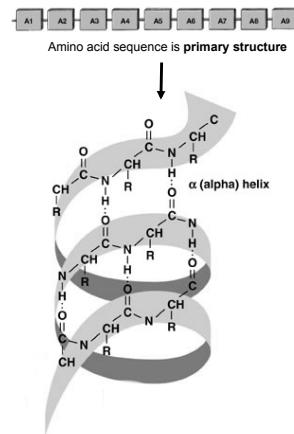
## Protein Secondary Structure



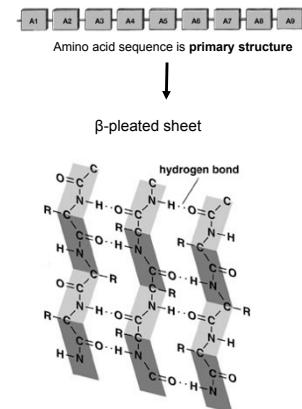
Secondary Structure Arises From the Hydrogen Bonds Between the Peptide Backbone



## Protein Secondary Structure

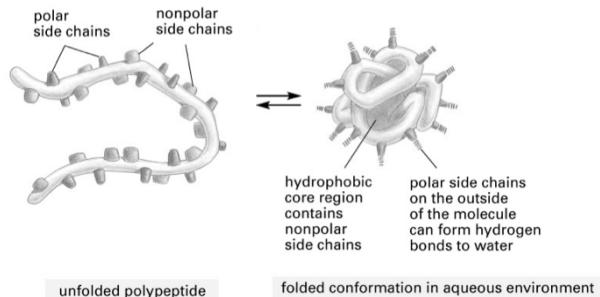


## Protein Secondary Structure

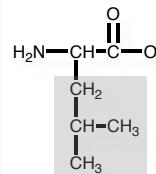


## Protein Tertiary Structure

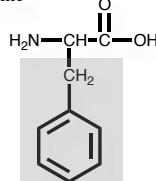
Tertiary Structure Arises From Folding of the Amino Acid Sidechains (R groups)



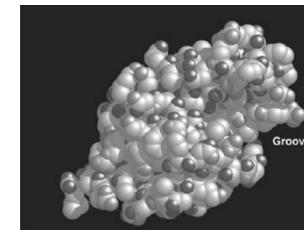
## The Hydrophobic Effect



Leucine

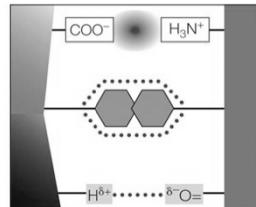


Phenylalanine



Hydrophobic side-chains will be near the center, stabilized by van der Waals forces.

## Protein Tertiary Structure, continued:

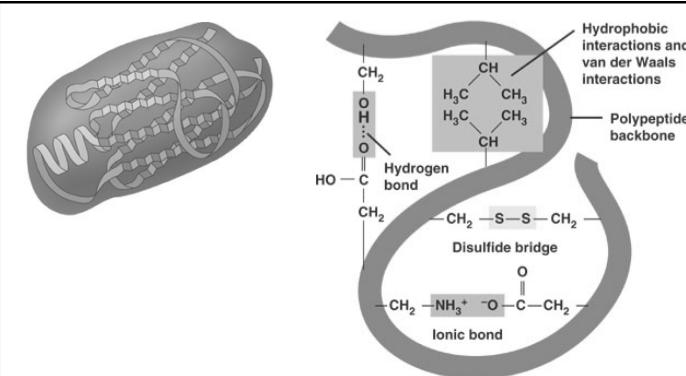
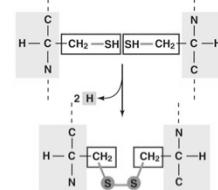


Ionic interactions occur between charged R groups.

Two nonpolar groups interact hydrophobically.

Hydrogen bonds form between two polar groups.

Cysteine R group contains  $-\text{SH}$ . Can react with second  $-\text{SH}$  to form a **disulfide bridge**

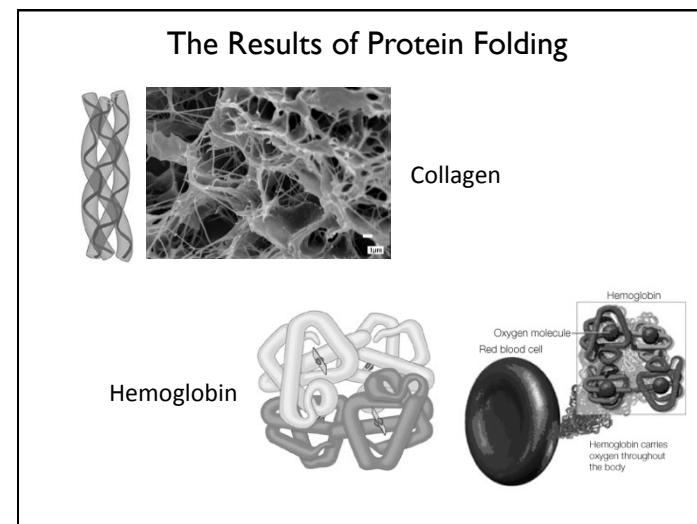
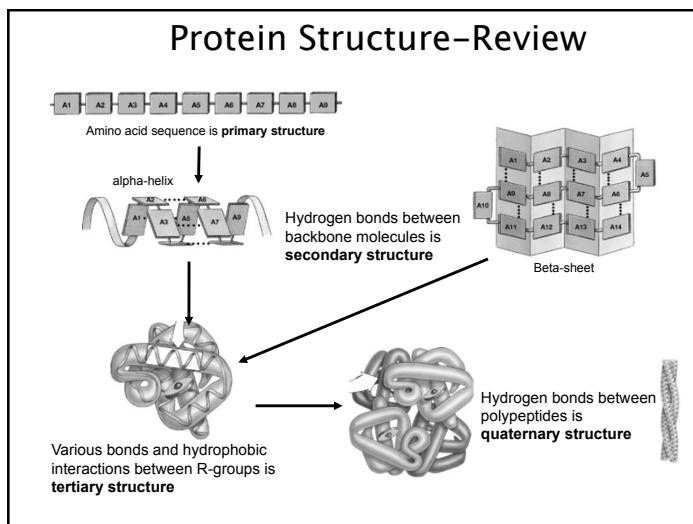
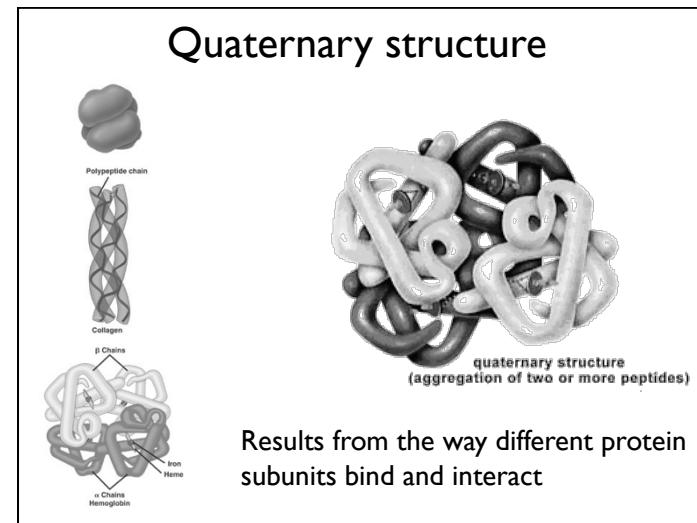
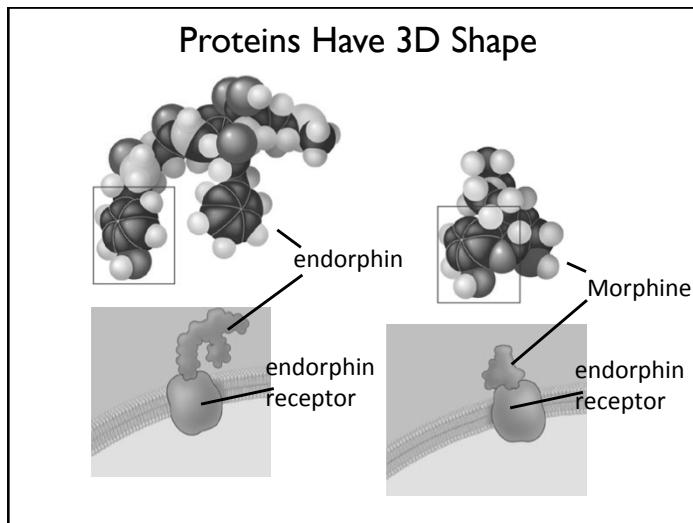


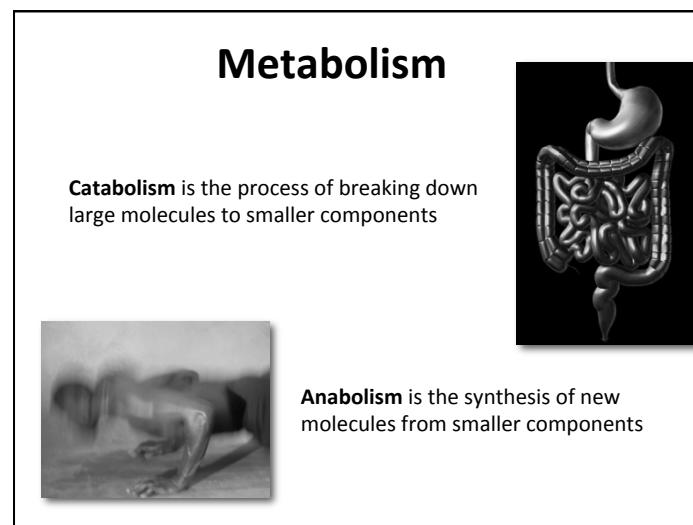
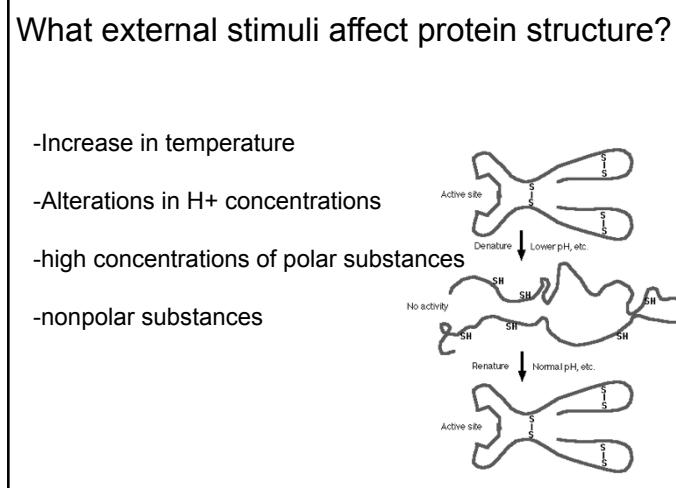
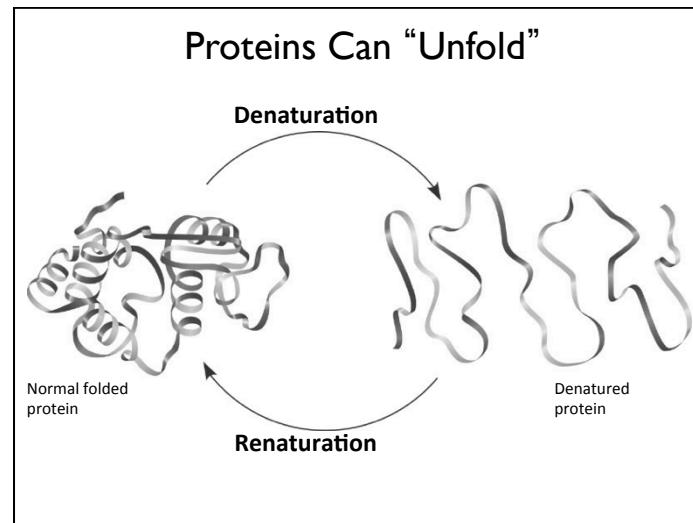
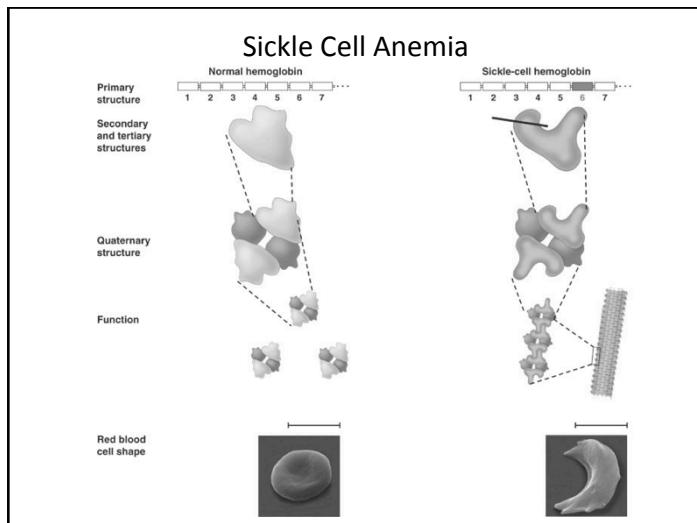
**H-bonds** between side chains stabilize folds

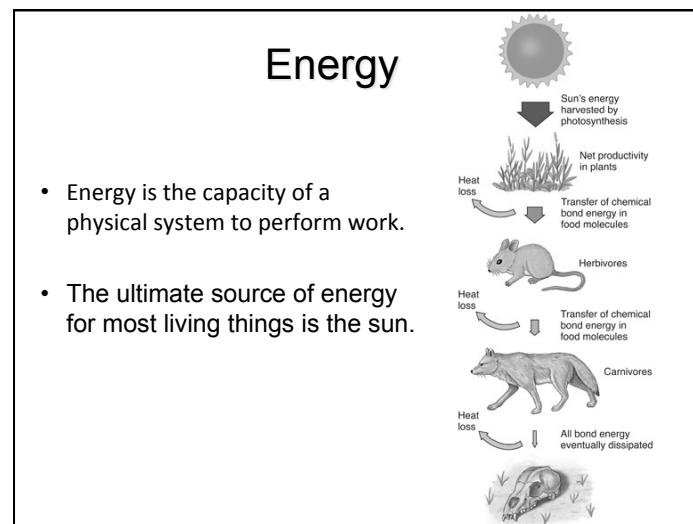
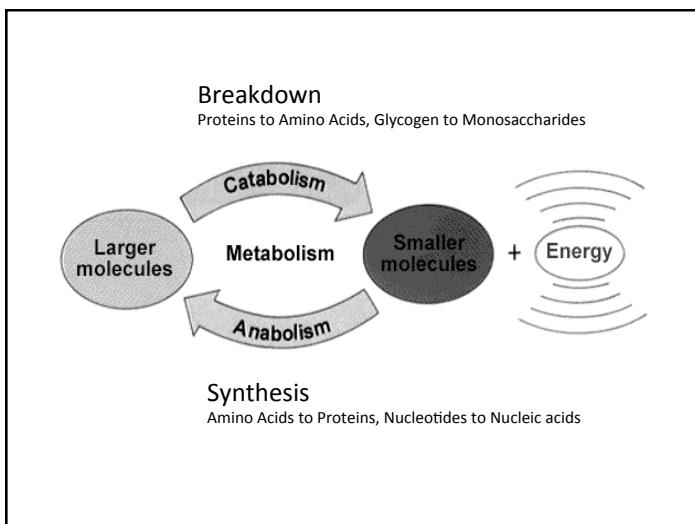
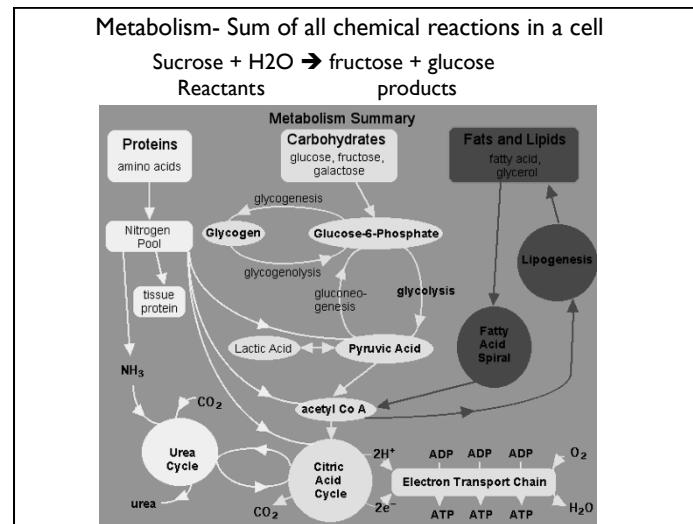
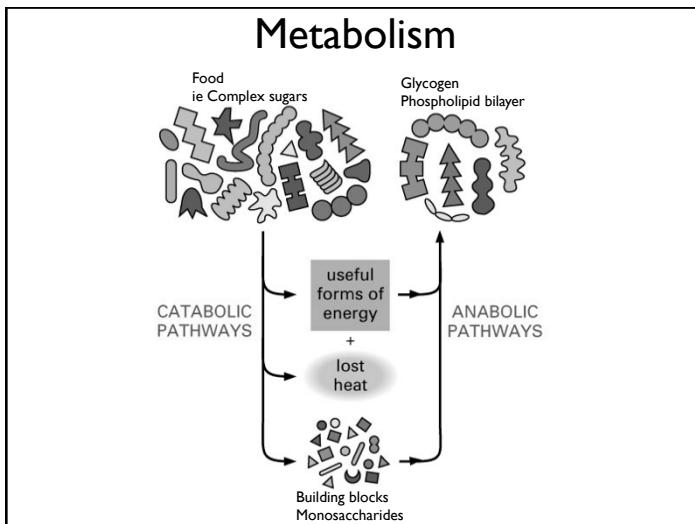
**Ionic** interactions form salt bridges

**Hydrophobic** side chains aggregate in interior

**van der Waals** stabilize hydrophobic side chains







## Energy

**Energy** - The capacity to do work (to move matter against an opposing force).

**Potential Energy** - The energy stored by matter as a result of its location or spatial arrangement.

*Potential Energy (gravitational)*



*Kinetic Energy*



**Kinetic Energy** - The energy of motion, which is directly related to the speed of that motion. Moving matter does work by imparting motion to other matter.

## Thermodynamics I

A. Energy cannot be created or destroyed

B. Energy can only be transformed into another form



## Thermodynamics II

A *closed* system moves toward entropy, increasing disorder.

Living systems are open systems that maintain organization and increase it during development



**ENTROPY**

It's not so much 'letting yourself go' as it is 'succumbing to the inevitable'

Life increases **order**...  
but increases overall **disorder** (entropy) in the process

## Chaos will reign

Entropy is a measure of disorder in the system



**ENTROPY**

Eg. Anabolic reactions required to make 1kg of your body requires 10kg of food. Rest is "waste".

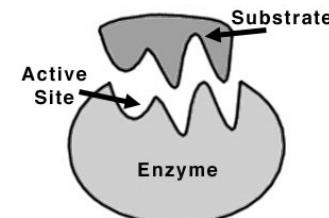
## 5 principles governing metabolism

- Complex chemical transformations occur through a series of separate intermediate reactions.
- These reactions are catalyzed by specific enzymes.
- These enzymes are activated/inactivated to control reaction rate.
- Most metabolic pathways are similar between organisms.
- In eukaryotes these metabolic pathways are compartmentalized in organelles.

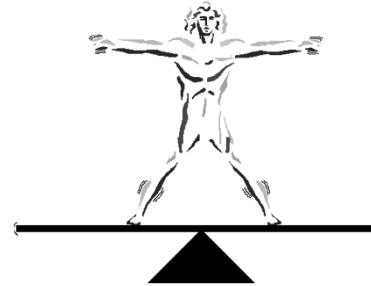
## Biological catalysts: Enzymes....

**Catalysts:** Proteins that speed up biochemical reactions  
Increase reaction rates without themselves being altered  
these reactions may occur without catalysts, but too slowly

**Enzyme:** most common type of catalyst



Enzymes catalyze metabolic reactions  
and maintain **homeostasis**



How do cells create this balance?  
 ▪ regulation of enzyme expression levels  
 ▪ regulation of enzyme activity

## Free Energy ( $\Delta G$ )

- **Free energy** – the energy available for doing work.
  - Most chemical reactions release free energy
  - they are **exergonic**.
    - Downhill
  - Some reactions require the input of free energy
  - they are **endergonic**.
    - Uphill

