



THE UNIVERSITY *of* TEXAS

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HEALTH SCIENCE CENTER AT HOUSTON

SCHOOL *of* HEALTH INFORMATION SCIENCES

# Introduction to Biomolecular Structure

For students of HI 6327 “Biomolecular Modeling”

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School of Health Information Sciences

<http://biomachina.org/courses/modeling/02.html>

# Early Life on Earth

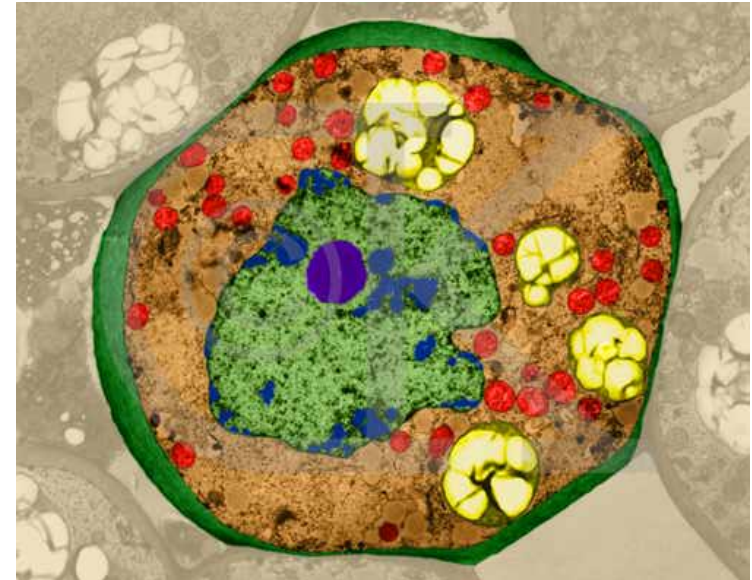
Time (Myr ago)	Event
4600	Formation of the approximately homogeneous solid Earth by planetesimal accretion
4300	Melting of the Earth due to radioactive and gravitational heating which leads to its differentiated interior structure as well as outgassing of molecules such as water, methane, ammonia, hydrogen, nitrogen, and carbon dioxide
4000	Bombardment of the Earth by planetesimals stops
3800	The Earth's crust solidifies--formation of the oldest rocks found on Earth. Condensation of atmospheric water into oceans
3500-2800	<b>Prokaryotic cell organisms develop</b>
3500-2800	<b>Beginning of photosynthesis by blue-green algae which releases oxygen molecules into the atmosphere</b>



**Stromatolites are layered mounds, columns, and sheets found in the rock. They were originally formed by the growth of layer upon layer of *cyanobacteria*, a single-celled photosynthesizing microbe growing on a sea floor.** Photo by Marjory Martin, Deakin Univ, Australia.

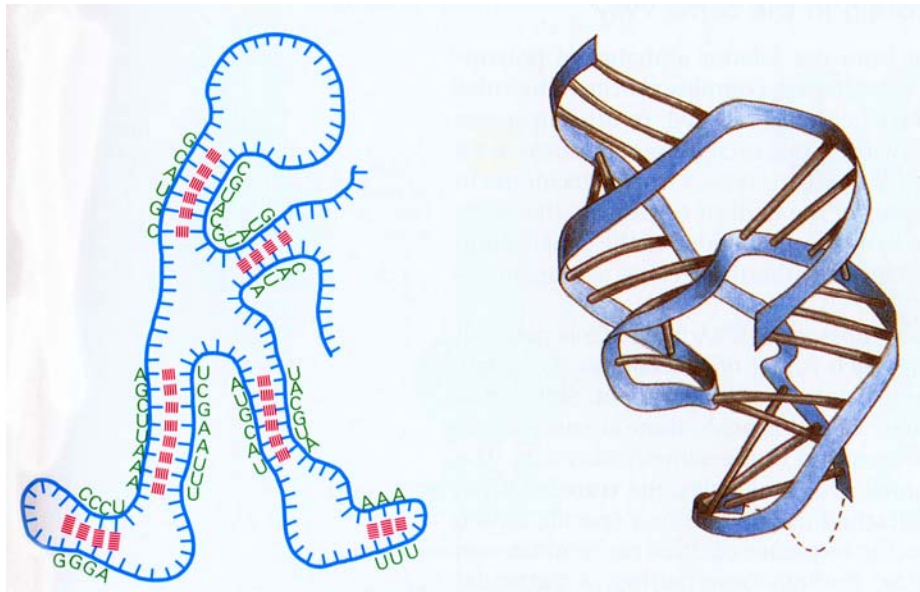
# Early Life on Earth

Time (Myr ago)	Event
1500-600	<b>Eukaryotic cell organisms develop, rise of multicellular organisms</b>
430	Waxy coated algae begin to live on land
420	Millipedes have evolved--first land animals
375	The Appalachian mountains are formed via a plate tectonic collision between North America, Africa, and Europe
200	Appearance of mammals
65	K-T (Kreide-Tertiär = Cretaceous-Tertiary) Boundary--extinction of the dinosaurs and beginning of the reign of mammals
20-12	The chimpanzee and hominid lines evolve
0.05-0	<i>Homo sapiens sapiens</i> exist



**Lily Parenchyma Cell (cross-section)** (TEM x7,210). Note the large **nucleus** and **nucleolus** in the center of the cell, **mitochondria** and **plastids** in the **cytoplasm**. Photo by Dennis Kunkel at [www.DennisKunkel.com](http://www.DennisKunkel.com)

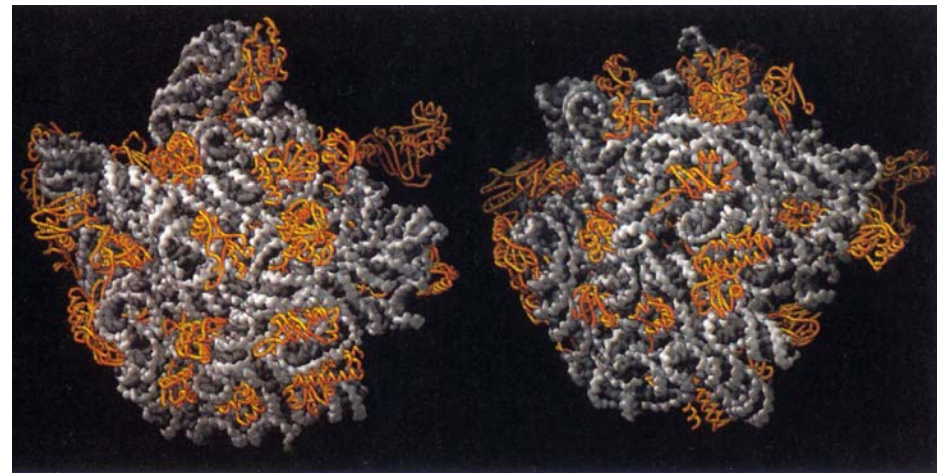
# The First Enzymes: RNA



**The conformation of an RNA molecule:  
Nucleotide pairing and 3D structure.**

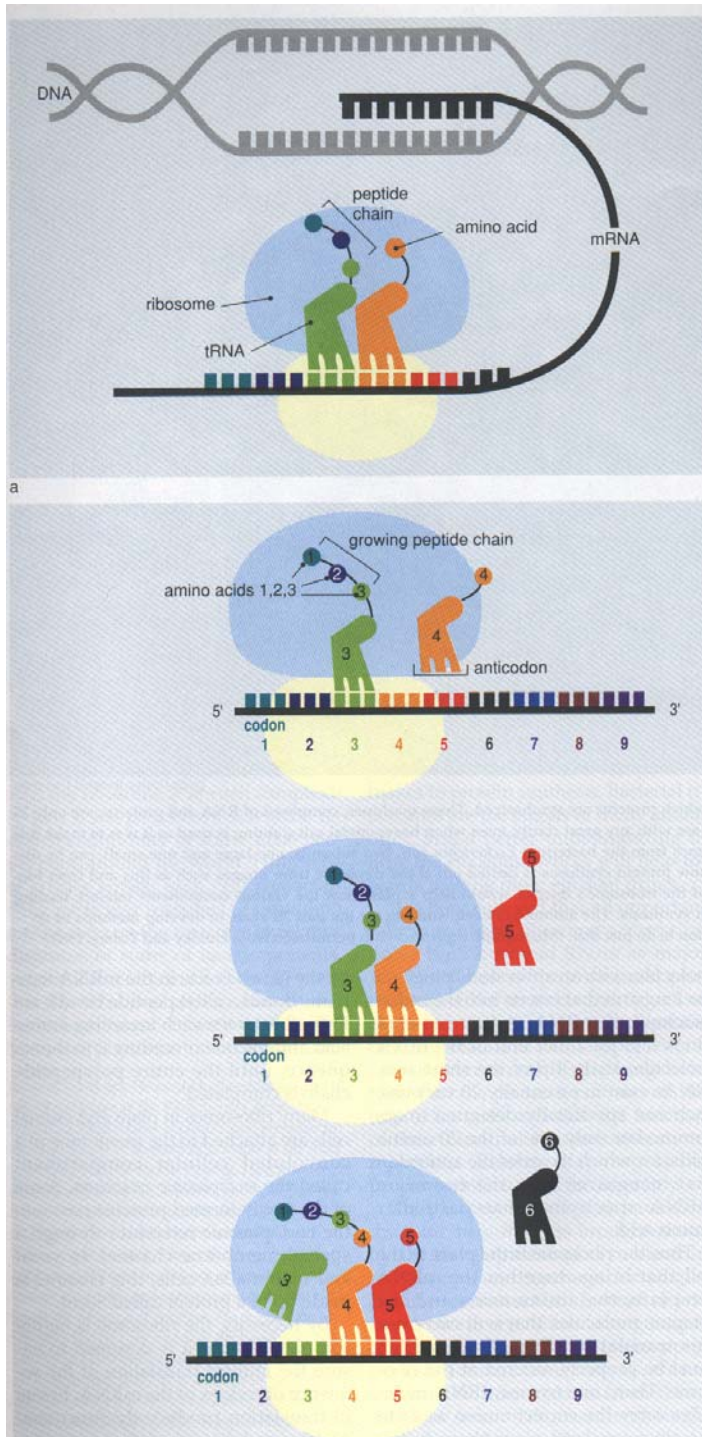
© Alberts et al. The Cell.

**Location of the protein components (gold)  
in the ribosome, that consists mainly of  
RNA (grey). © Ban et al. Science.**

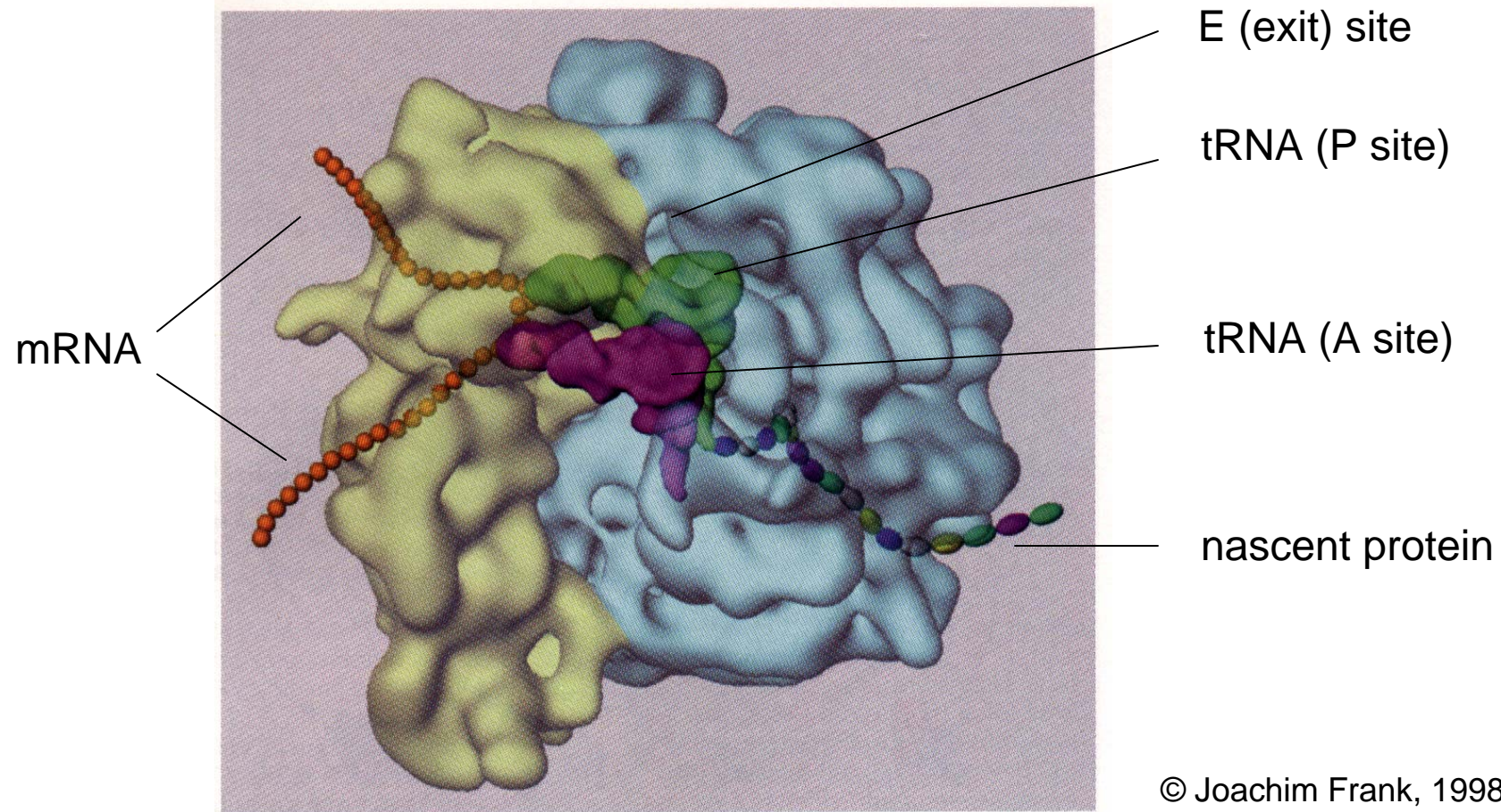


# Protein Synthesis in the Ribosomal Translation Cycle

1. mRNA synthesis with RNA polymerase
2. aa-tRNA (1 anticodon - 3b) acts as adapter
3. anticodon matches codon on mRNA
4. aa binds to polypeptide chain
5. release of tRNA
6. new tRNA binds



# EM Map of the Ribosome at 15Å Resolution

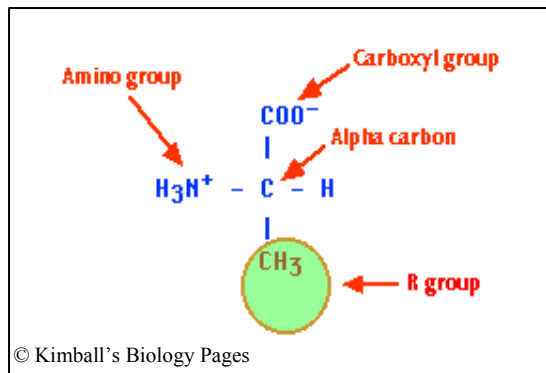


# The Amino Acids

- **Proteins are polymers of the 20 naturally occurring amino acids**
- **A.a. are abbreviated by 3 and 1 letter codes (learn these!)**
- **A.a. can be grouped based on electrostatic and size of side chain R**

## Aliphatic:

Alanine	Ala A
Valine	Val V
Leucine	Leu L
Isoleucine	Ile I



## Polar:

Asparagine	Asn N
Glutamine	Gln Q
Serine	Ser S
Threonine	Thr T

## Non-polar:

Glycine	Gly G
Proline	Pro P
Cysteine	Cys C
Methionine	Met M

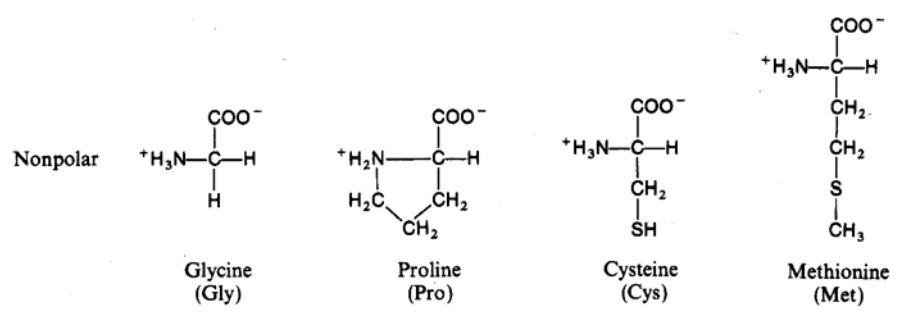
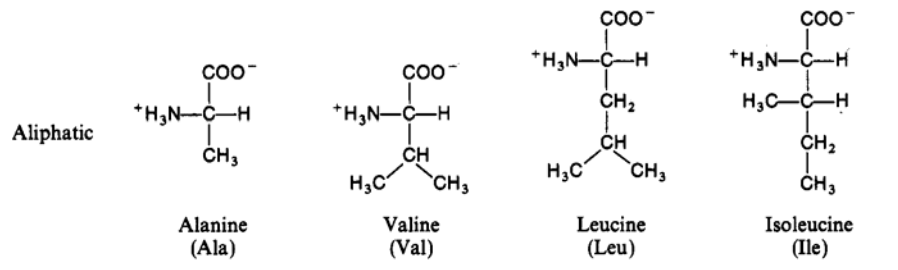
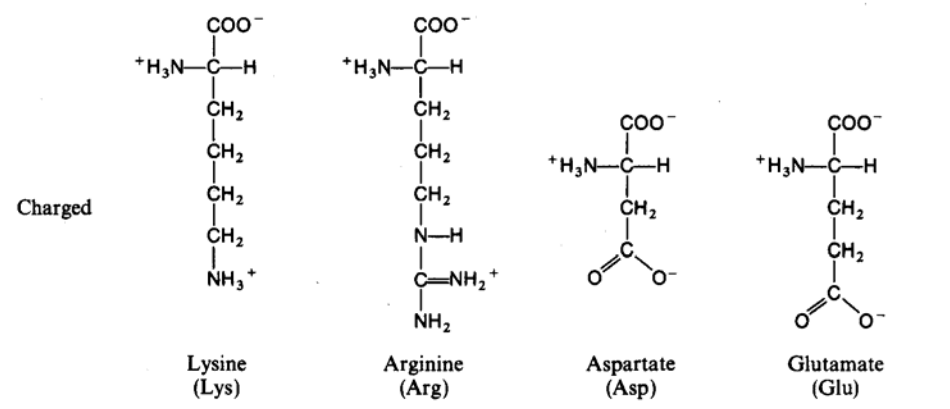
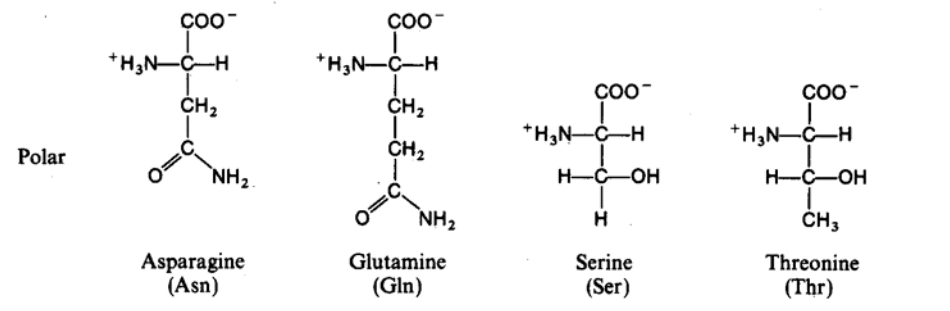
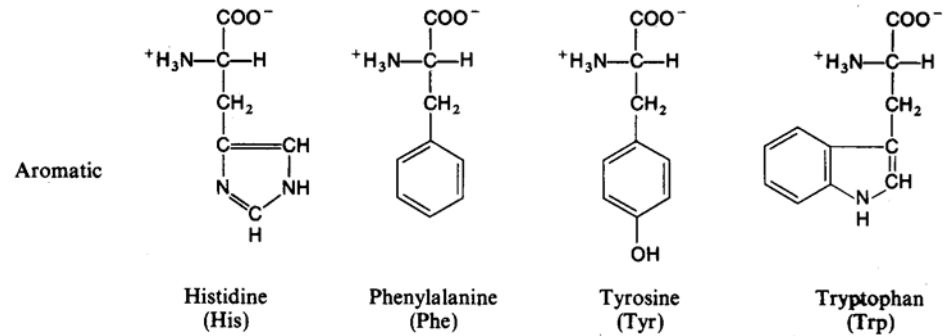
## Aromatic:

Histidine	His H
Phenylalanine	Phe F
Tyrosine	Tyr Y
Tryptophan	Trp W

## Charged:

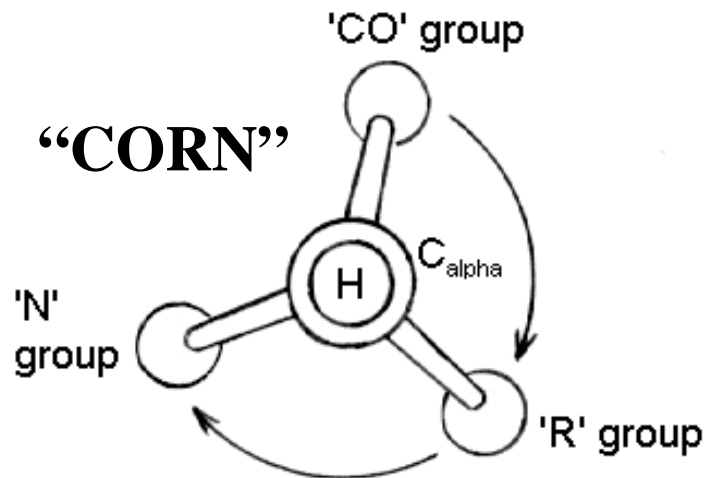
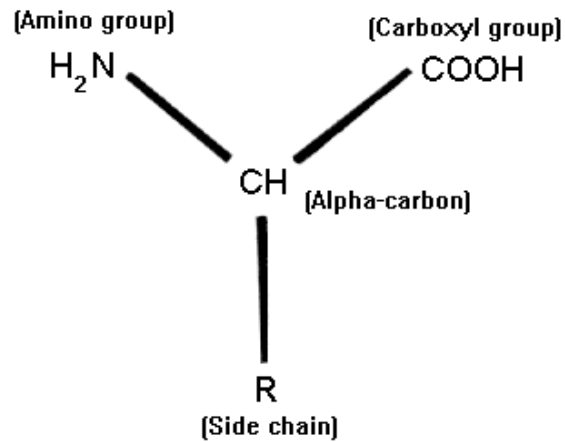
Lysine	Lys K
Arginine	Arg R
Aspartic Acid	Asp D
Glutamic Acid	Glu E

# 2D Structures of Amino Acids





# 3D Structure



An example of the atom colour conventions used with amino acids. Can you identify this residue?

Red is used to represent oxygen atoms, in this case the side chain hydroxyl.

Carbon atoms in methyl or methylene groups are usually represented in white.

Blue shows nitrogen atoms. This one is the main chain amino group.

Sulphur atoms are coloured yellow. Which amino acids contain this element?

This red atom is the main chain carbonyl oxygen.

# Side Chain Protonation and pH

- **pH** measures the concentration of  $H^+$  ions in solution.
- $H^+$  from dissociation of an acid when this is dissolved in water.

The pH value is the negative logarithm of the  $H^+$  concentration in mol/L:

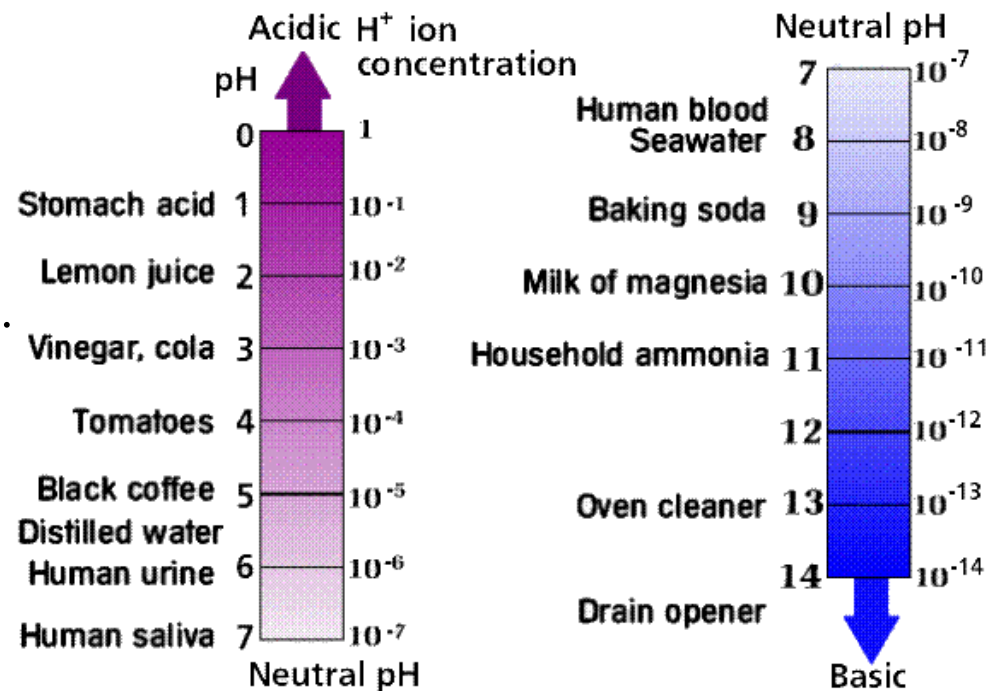
$$\text{pH} = -\log_{10}[\text{H}^+]$$

The  $[\text{H}^+]$  in pure water is  $10^{-7}$ ; therefore the **neutral** pH of pure water is:

$$\text{pH} = 7$$

- Below pH 7: higher  $[\text{H}^+]$  acidic.
- Above pH 7: lower  $[\text{H}^+]$  basic
- Cellular pH is approximately 7.2-7.4.

**pH of some common items.**  
Image from Purves et al.,  
Life: The Science of Biology



# pKa

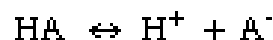
“dissociation point”

**pH < pKa:**  
**H<sup>+</sup> on**

**pH > pKa:**  
**H<sup>+</sup> off**

## acid-base equilibrium

The ionization equilibrium of a weak acid is given by



The equilibrium constant  $K$  for this weak acid is

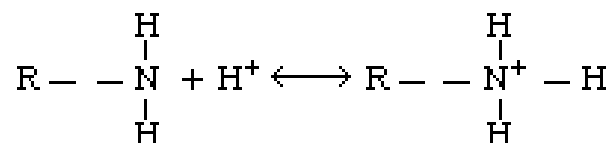
$$K = \frac{[\text{H}^+][\text{A}^-]}{[\text{HA}]}$$

The  $\text{p}K_a$  of an acid is defined as

$$\text{p}K_a = -\log K = \log \frac{1}{K}$$

By looking again at the 2nd equation, it can easily be shown that  $\text{p}K_a = \text{pH}$  when the acid is half dissociated,  $[\text{A}^-] = [\text{HA}]$ .

The relationship between pH and pKa is very important to understand, because this relationship describes how chemicals change states in biological systems as the pH varies. For example, an amino group has two possible states representable by  $\text{A}^-$  and  $\text{HA}$ :



It is important to know which state of the equilibrium is favored at different pH's.

This relationship between pH and pKa is described by the Henderson-Hasselbalch Equation:

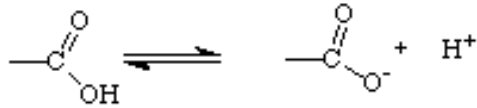
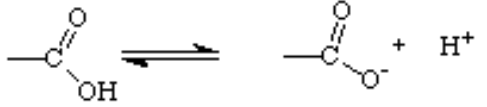
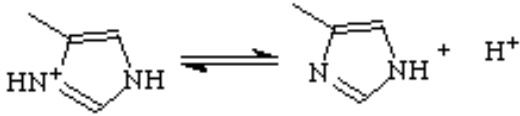
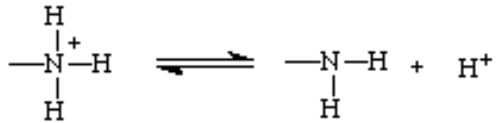

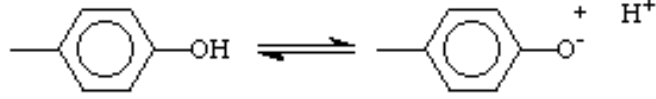
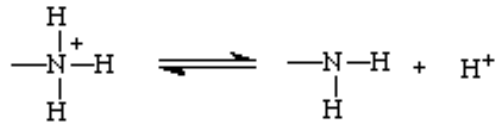
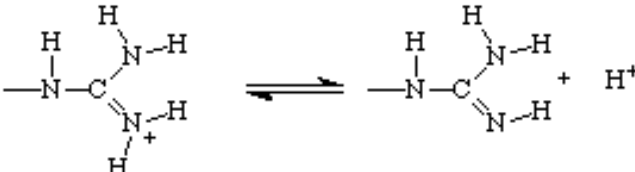
$$\text{pH} = \text{pKa} + \log \frac{[\text{A}^-]}{[\text{HA}]}$$

This equation is quite useful. With it you can now predict what state a molecule will be in at a given pH, among other things.

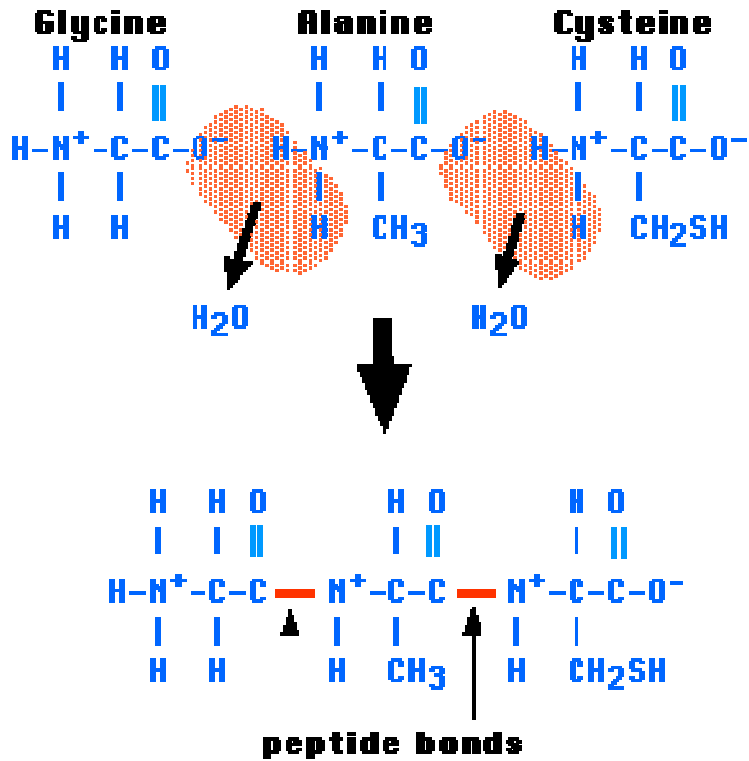
# pKa Chart

## for amino acids

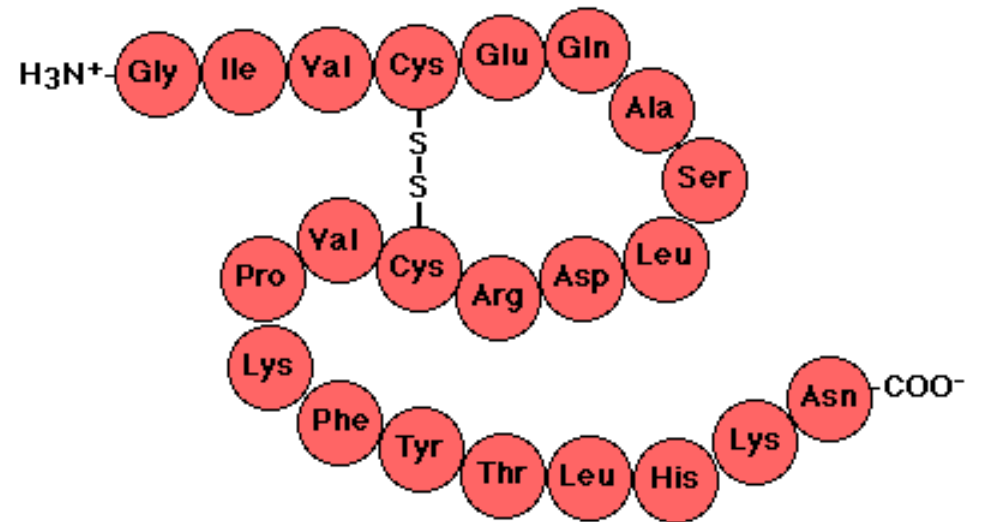
**Note:** Histidine protonation depends on protein environment (pKa shift) at neutral pH. Can have either N, or both, protonated.

<u>Group</u>	<u>Acid</u> $\rightleftharpoons$ <u>Base</u> + $H^+$	<u>pKa</u>
Terminal carboxyl		3.1
aspartic acid or glutamic acid		4.4
Histidine		6.5
Terminal Amino		8.0
Cysteine		8.5
Tyrosine		10.0
Lysine		10.0
Arginine		12.0

# Polypeptides



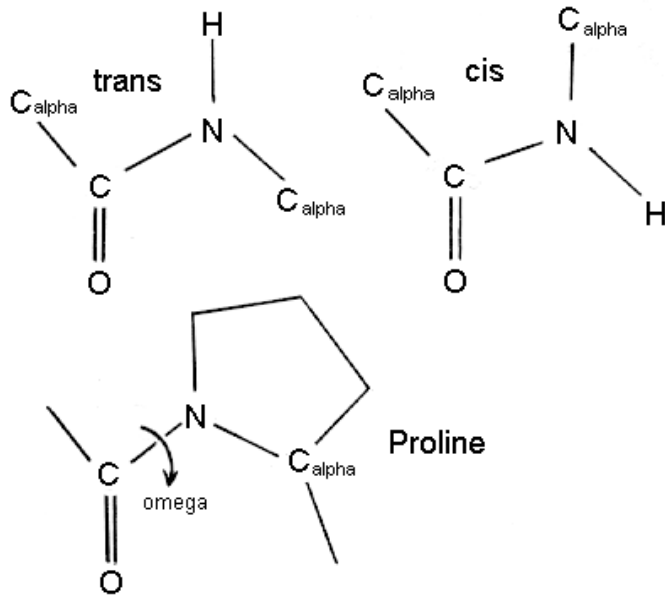
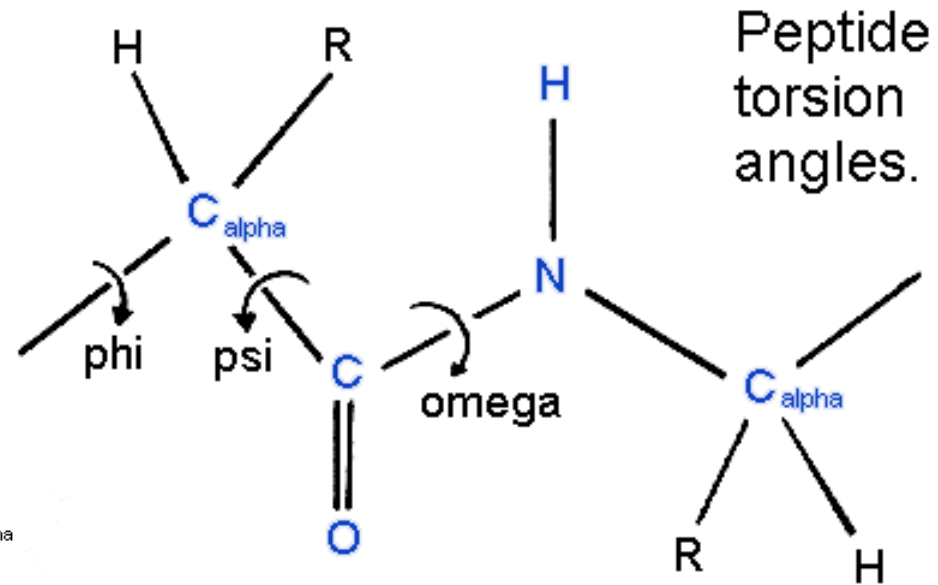
“N-terminus”



“C-terminus”

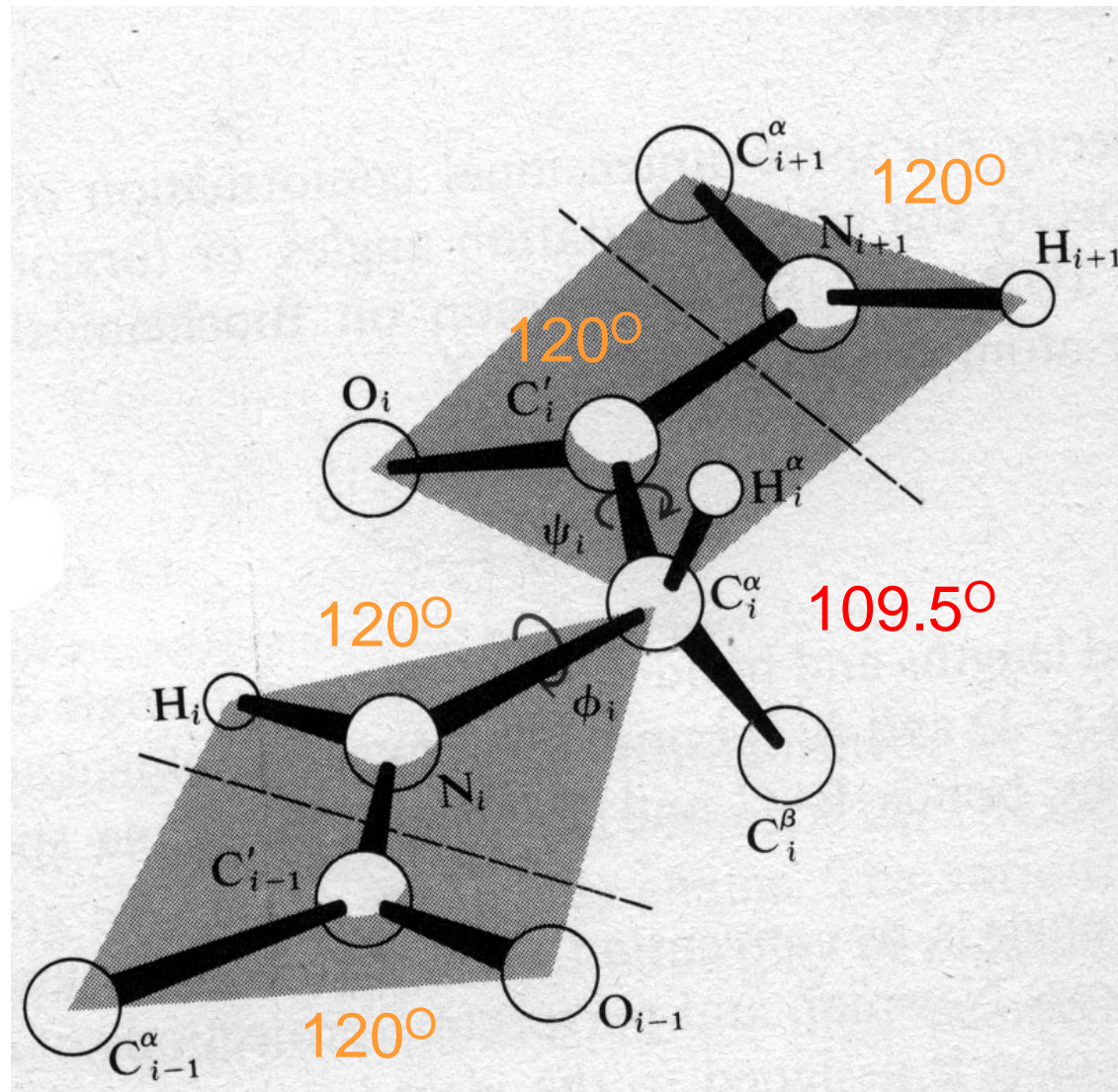
# Planarity of the Peptide Bond

Calpha(i), C(i), O(i),  
N(i+1) H(i+1) and  
Calpha(i+1) are co-planar



In all a.a. (except proline) steric hindrance favors the trans configuration ( $\omega = 180^\circ$ ), in proline sometimes  $\omega = 0$ .

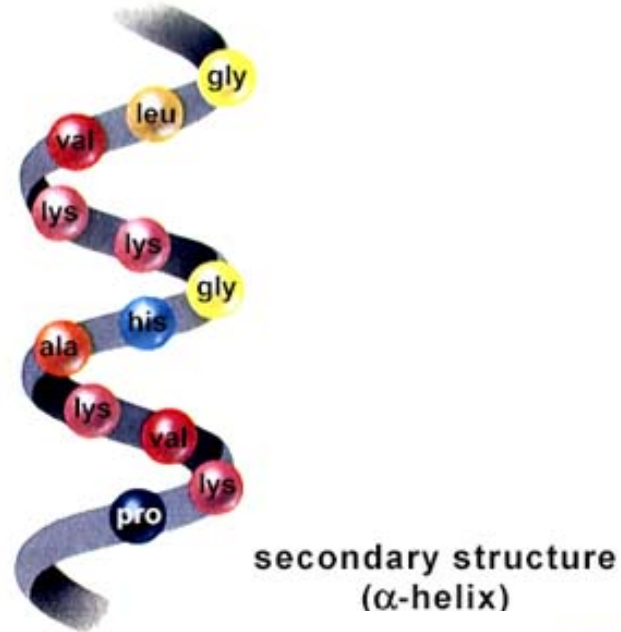
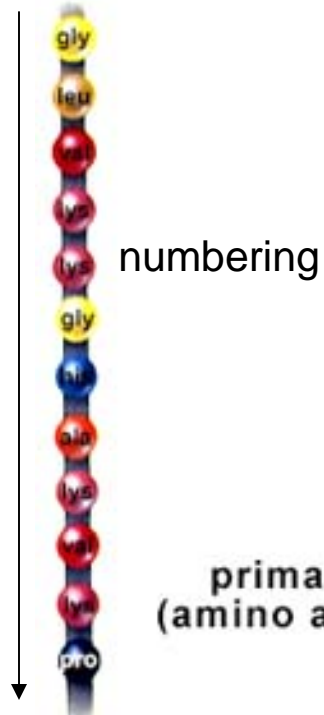
# Peptide Bond in 3D



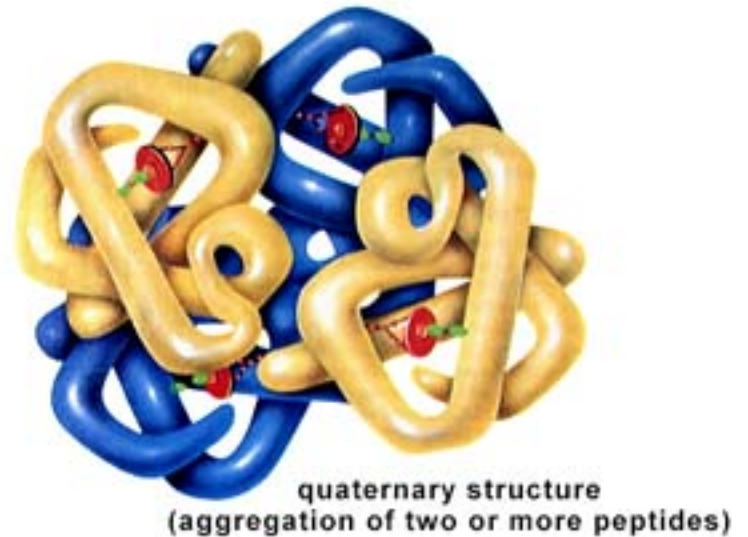
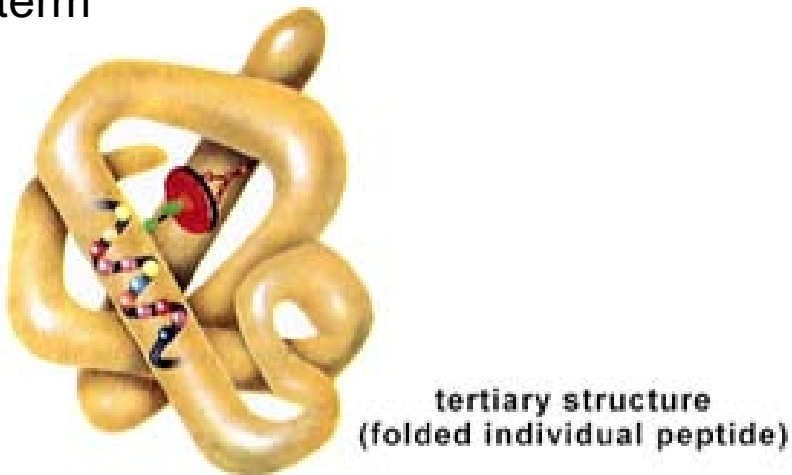
These angles are approximate and should be used in HW 1

# Protein Structural Hierarchy

N-term

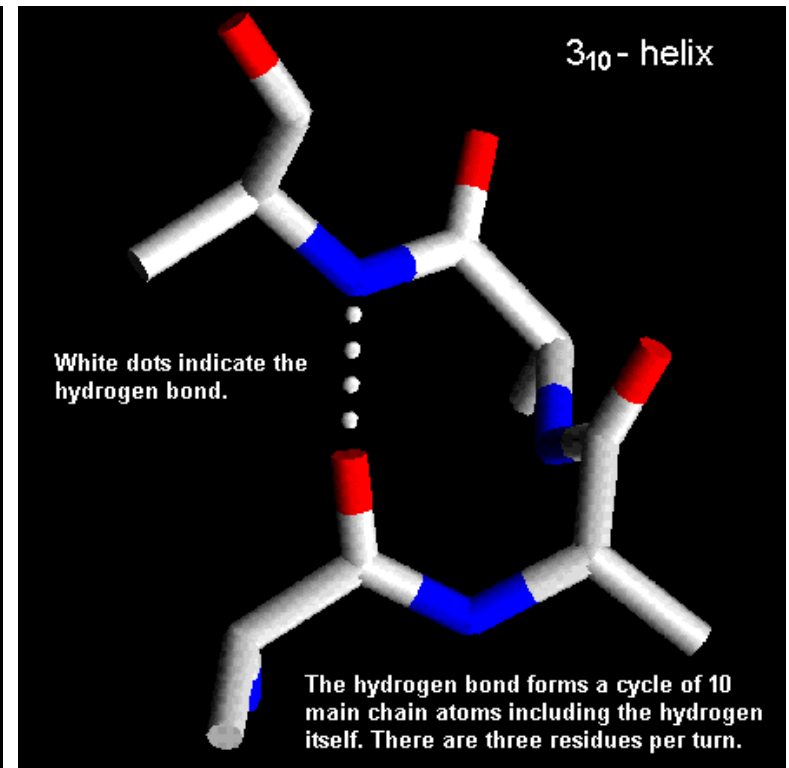
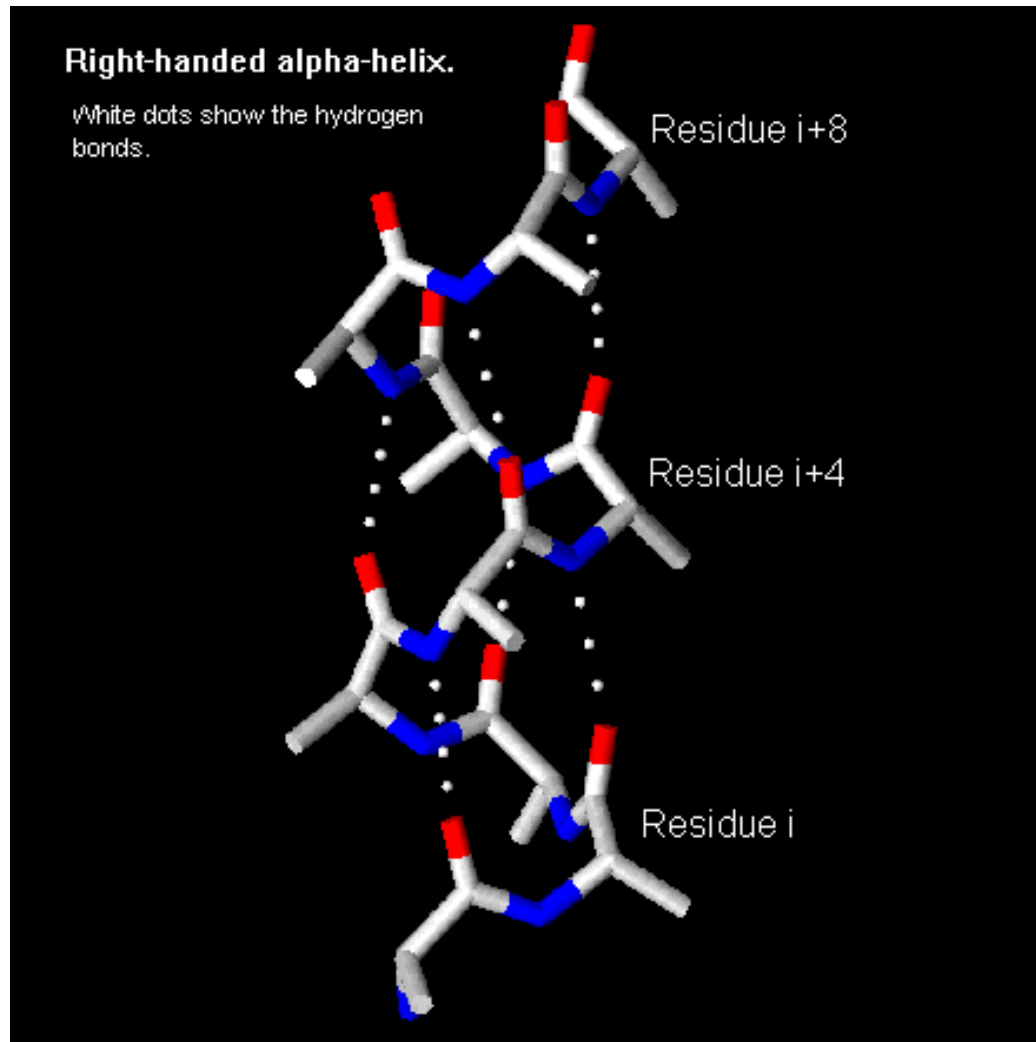


C-term



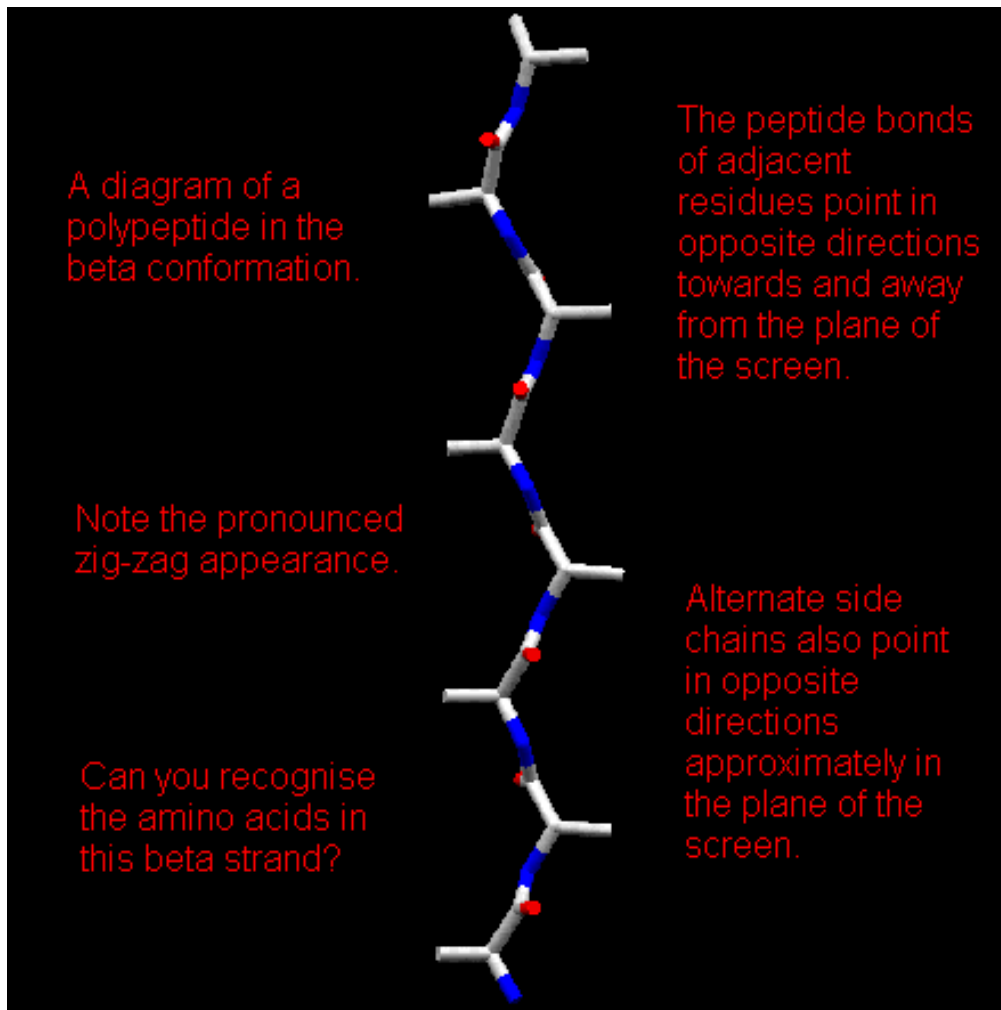


# Secondary Structure: $\alpha$ -Helix

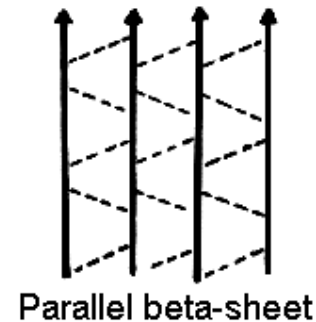
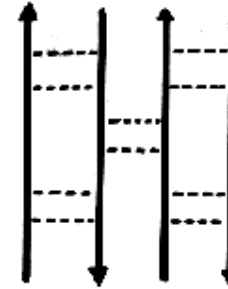


- $\alpha$ -helix (R):
- repeats every  $5.4\text{\AA}$ ,
  - 3.6 a.a. per turn
  - frequently terminated by  $3(10)$  helix

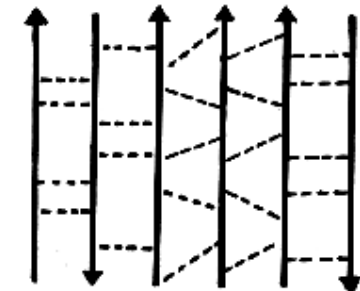
# Secondary Structure: $\beta$ -Sheet



Antiparallel beta-sheet

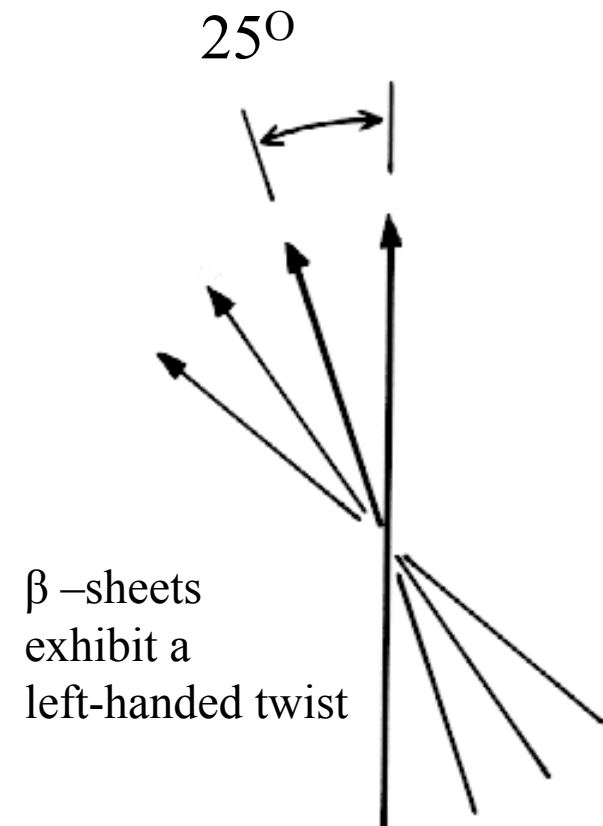
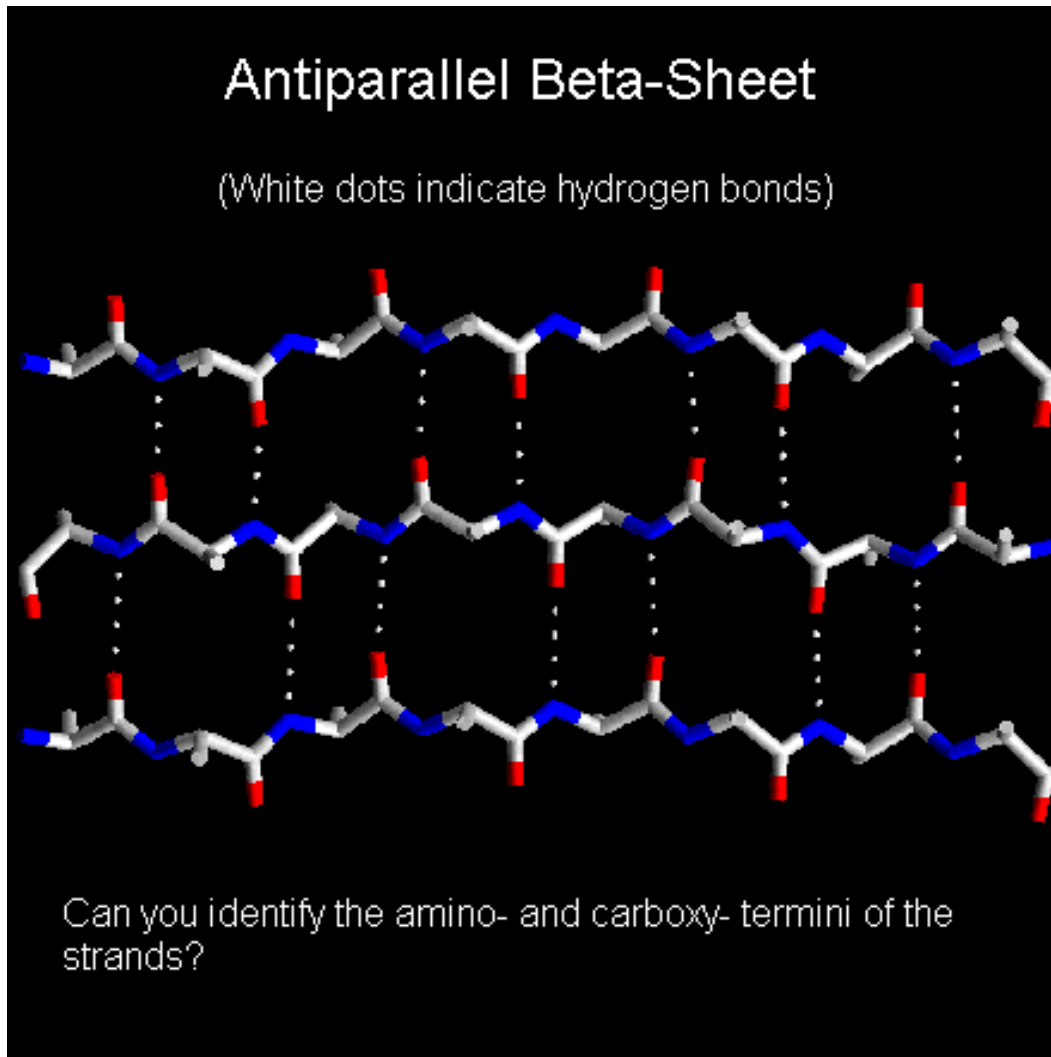


The different types of beta-sheet. Dashed lines indicate main chain hydrogen bonds.

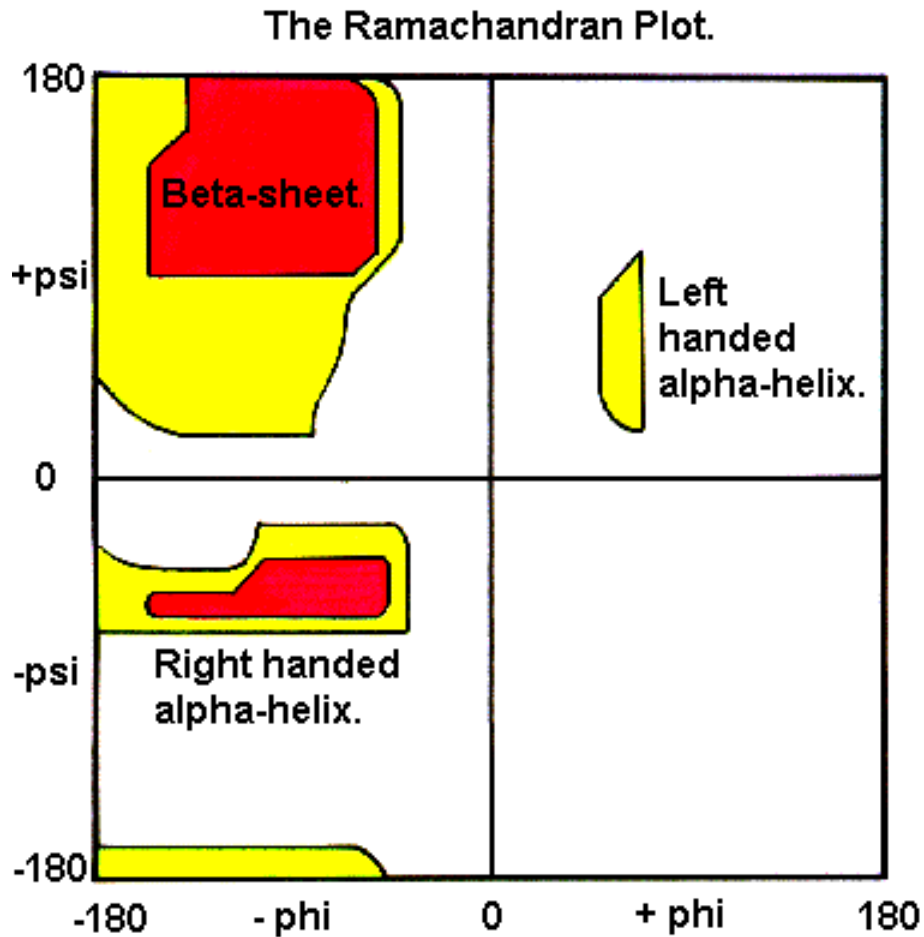


Mixed beta-sheet

# Secondary Structure: $\beta$ -Sheet

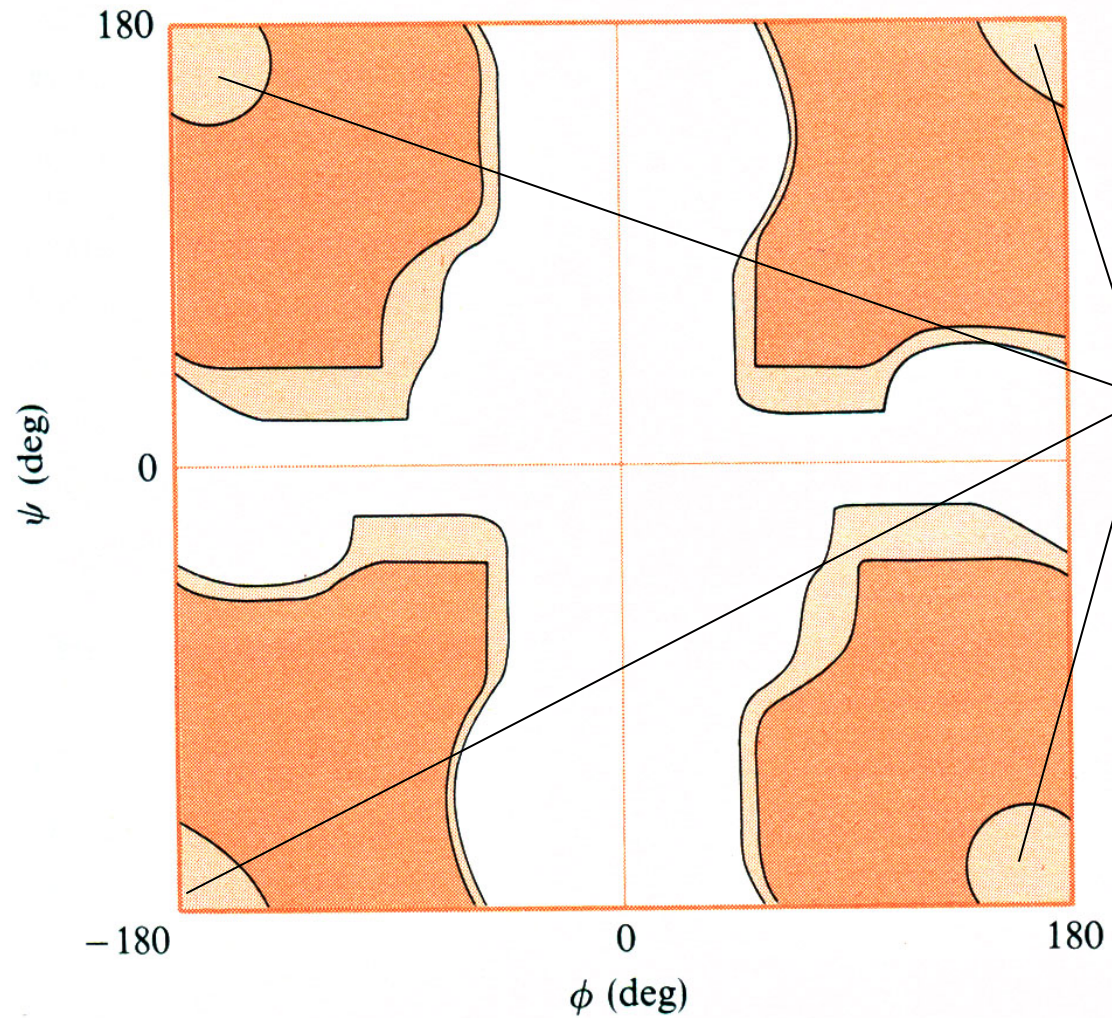


# The Ramachandran Plot



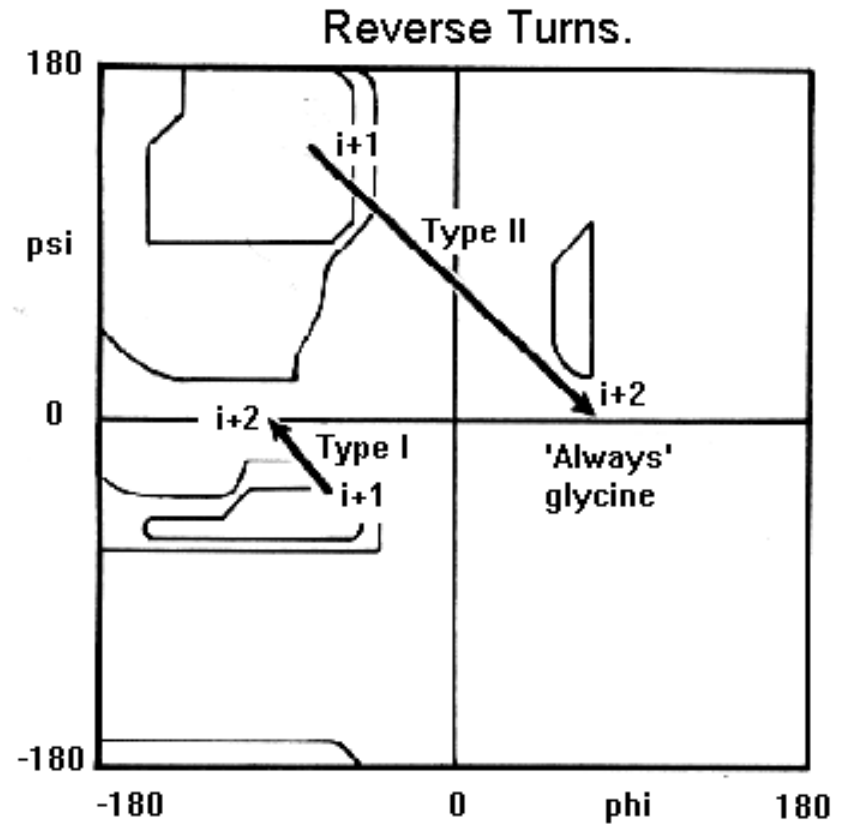
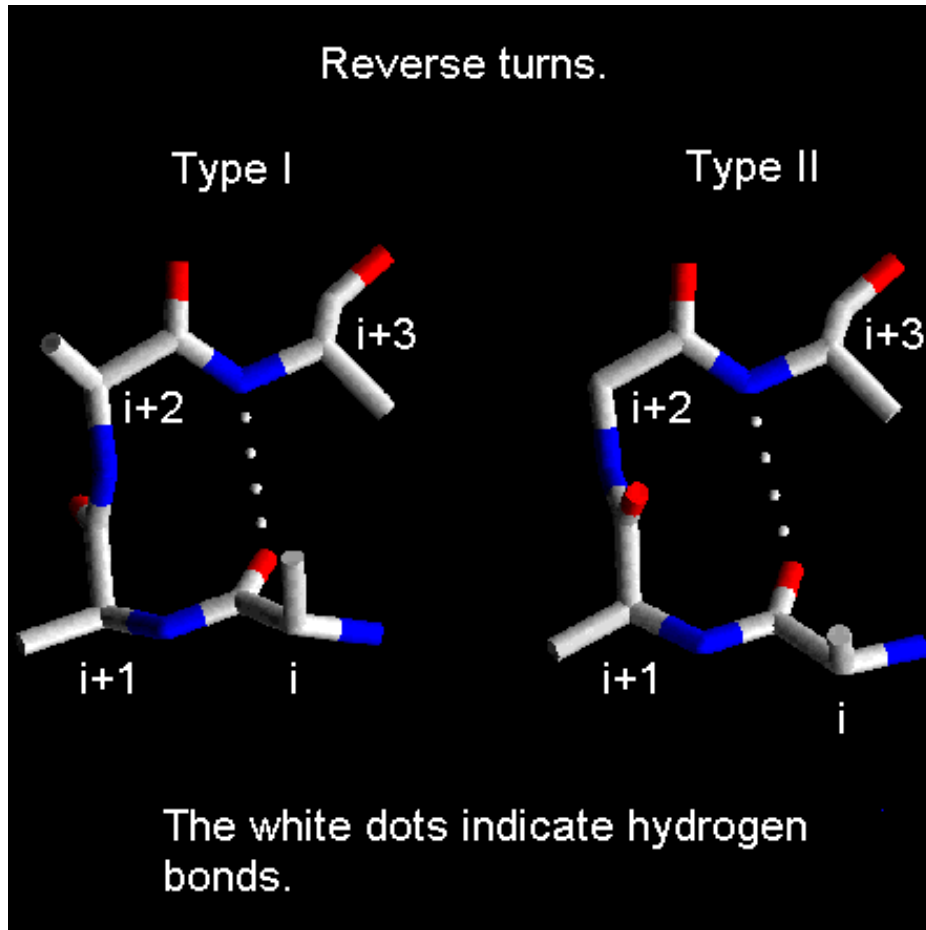
- hard-sphere model
- sterically disallowed regions
- valid for all a.a. except Gly
- Gly has no side chain

# Glycine Ramachandran Plot



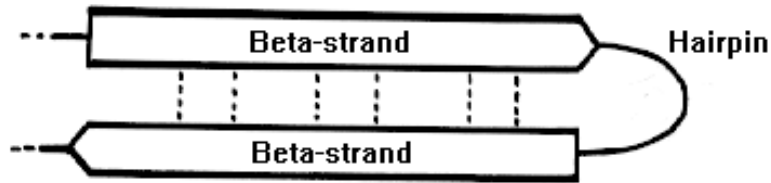
(hydrogen,  
ignore in  
HW 1)

# Reverse Turns

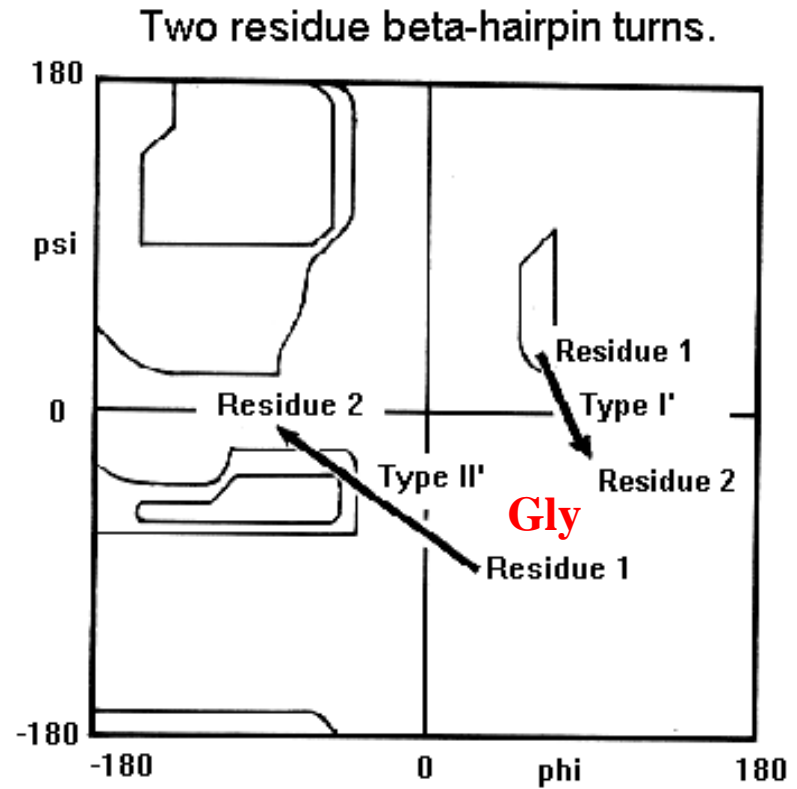
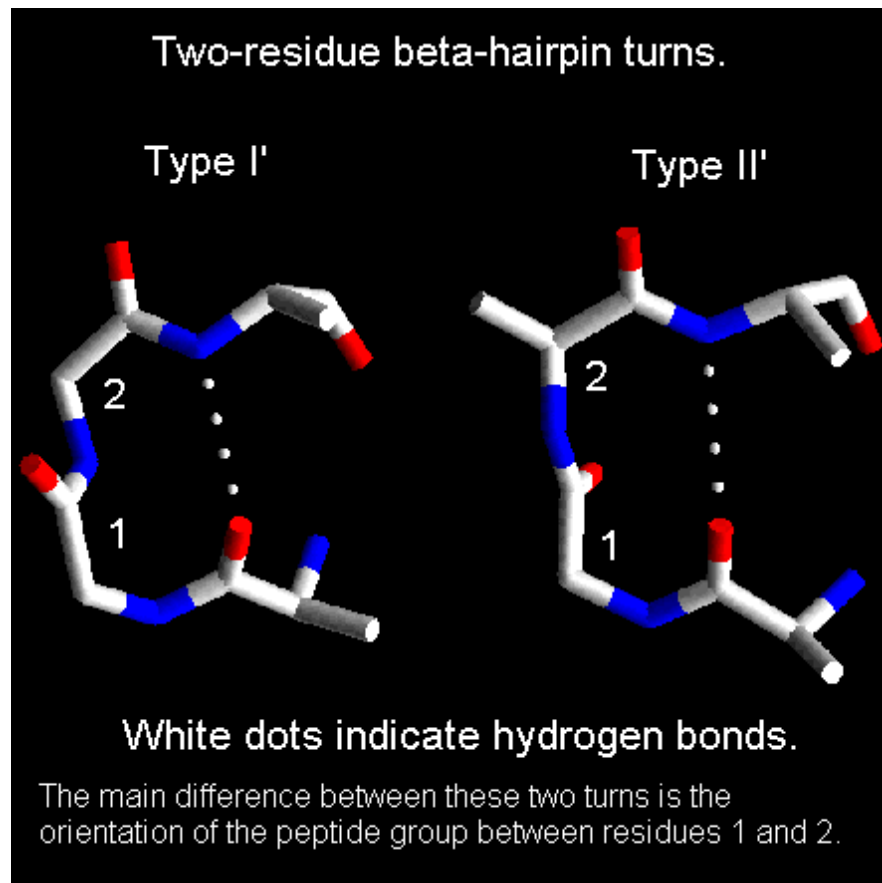


- abundant in globular proteins
- occur on surface of molecule
- possibly nucleation center for folding

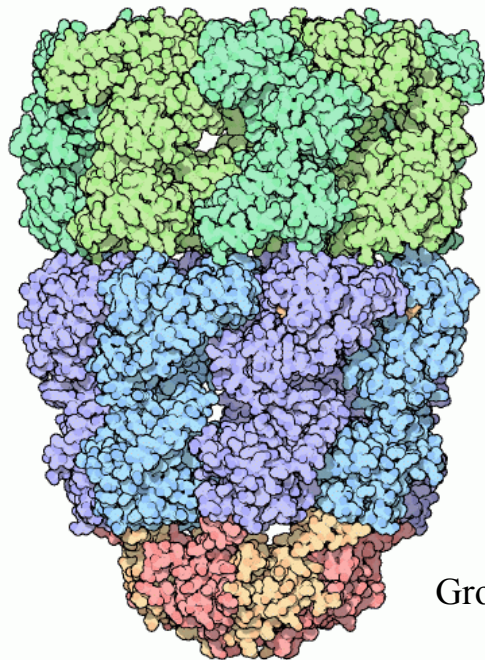
# $\beta$ –Hairpins



The dashed lines indicate main chain hydrogen bonds.

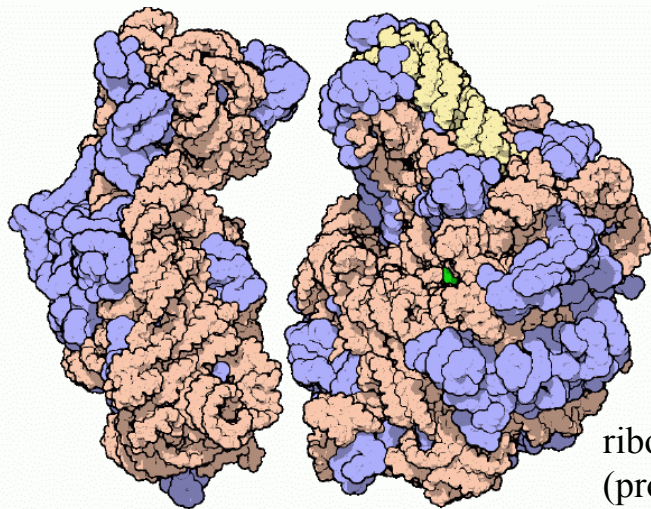
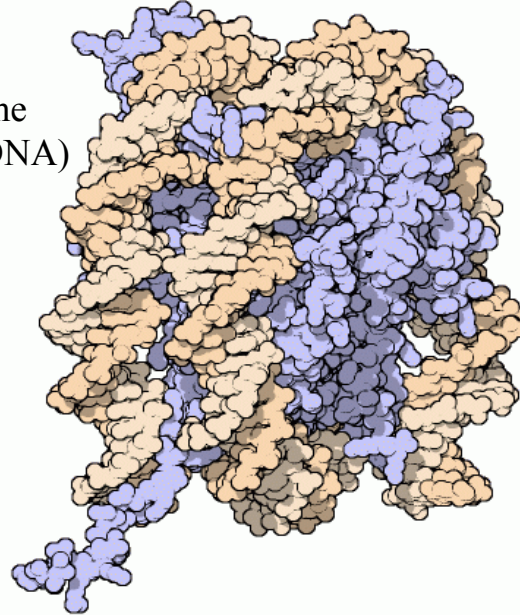


# Tertiary and Quaternary Structure

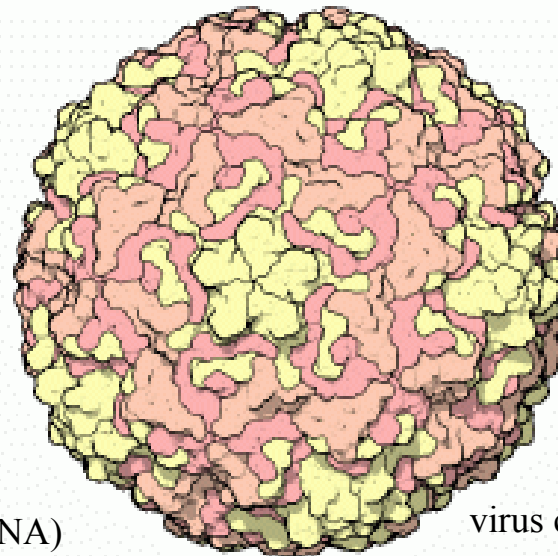


GroEL chaperonin

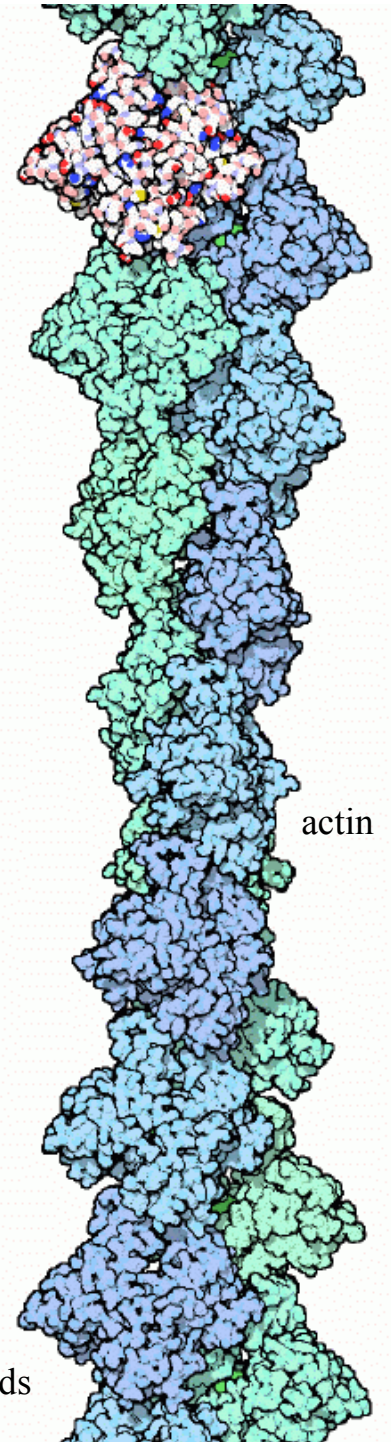
nucleosome  
(protein/DNA)



ribosome  
(protein/RNA)



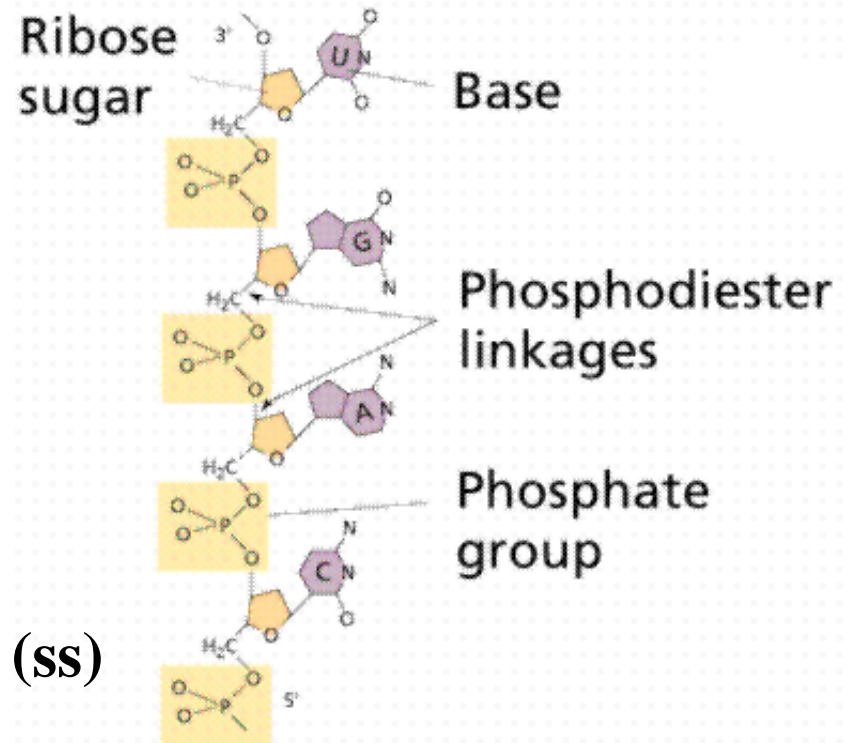
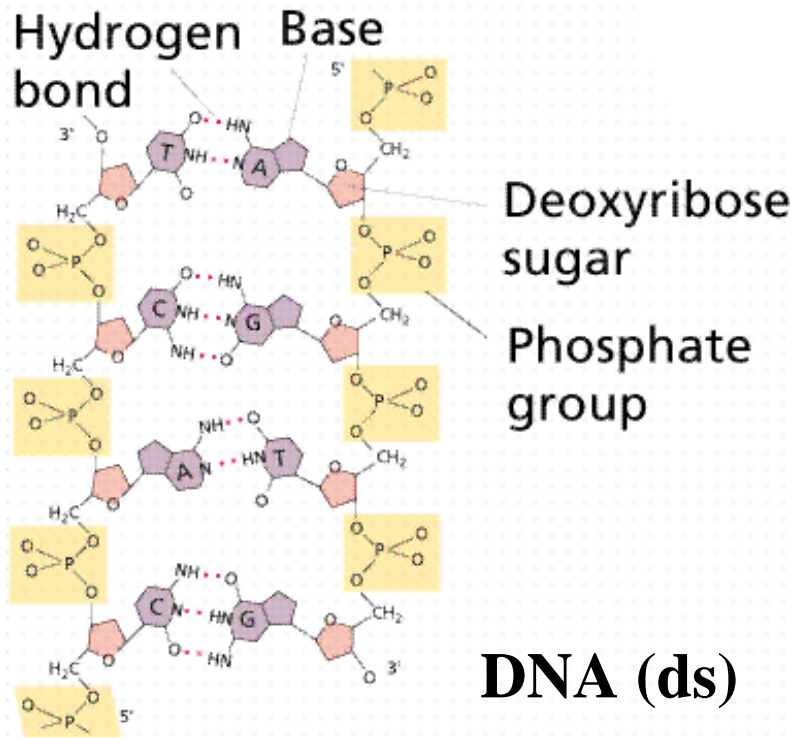
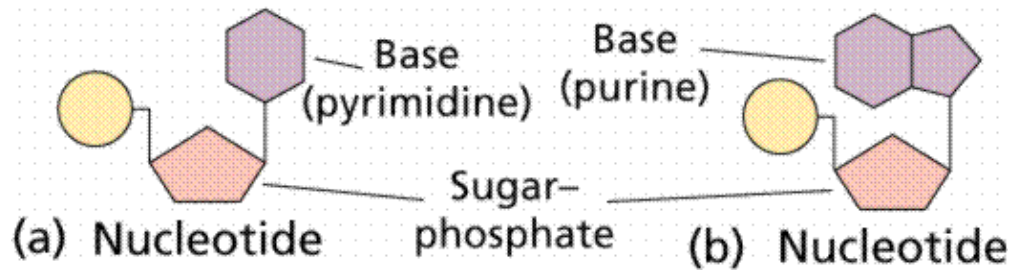
virus capsids



actin



# Polynucleotides



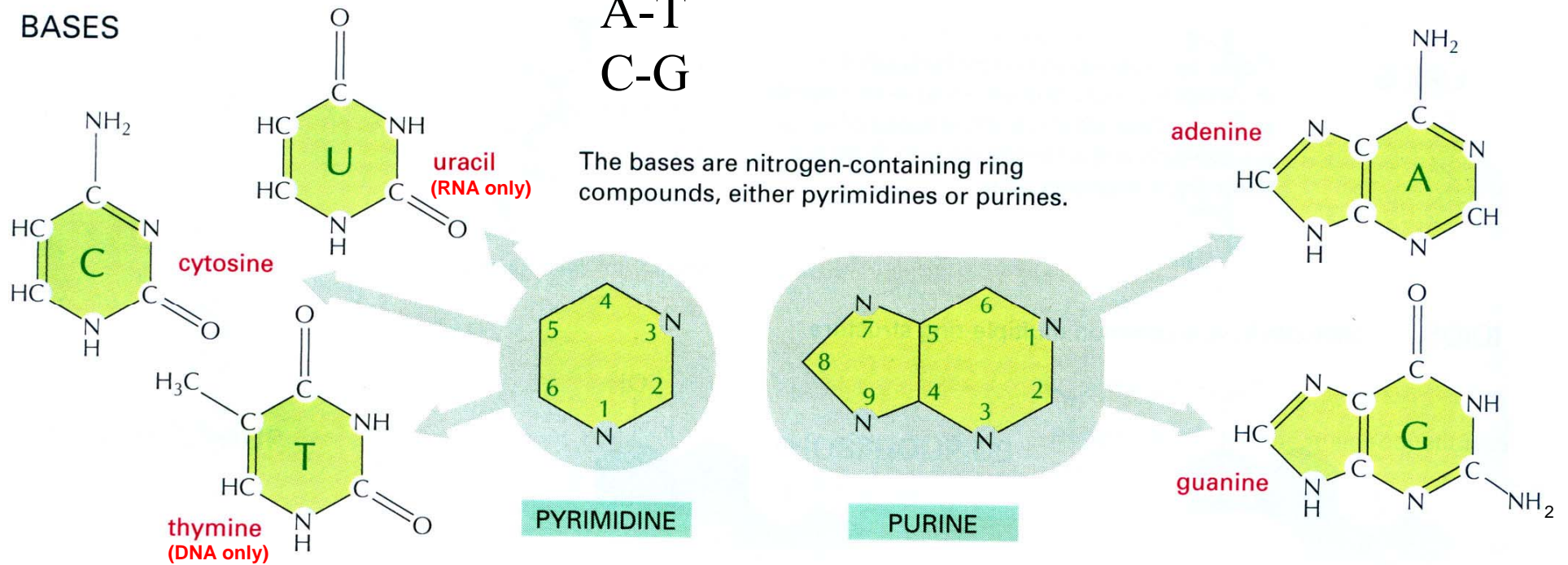
# Bases

Pairings (in DNA):

A-T

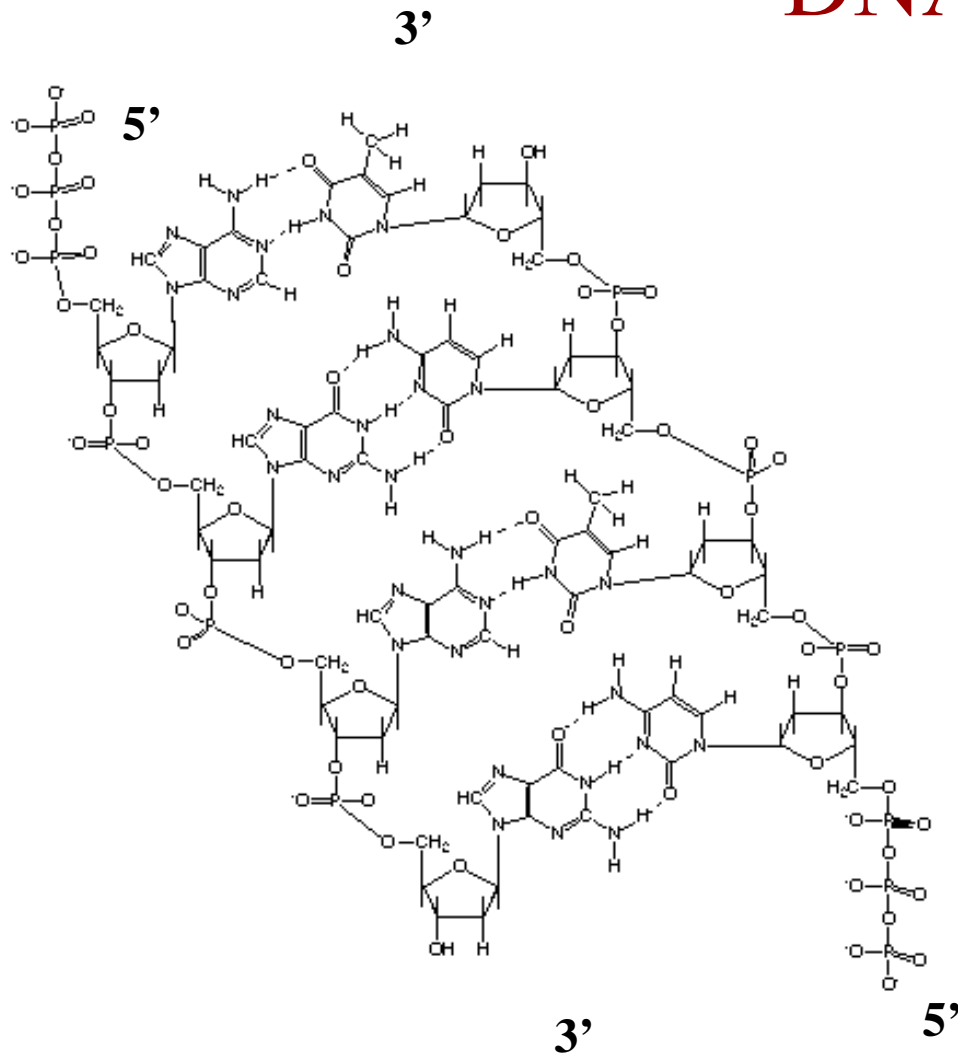
C-G

The bases are nitrogen-containing ring compounds, either pyrimidines or purines.



Convention: sequences written 5' → 3'

# DNA

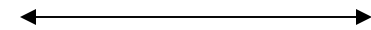
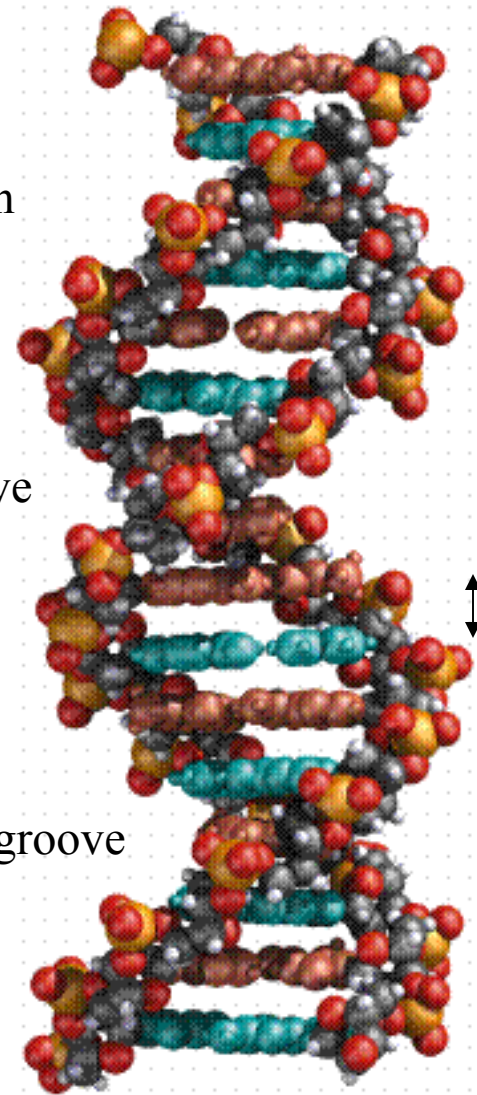


35.7Å pitch

minor groove

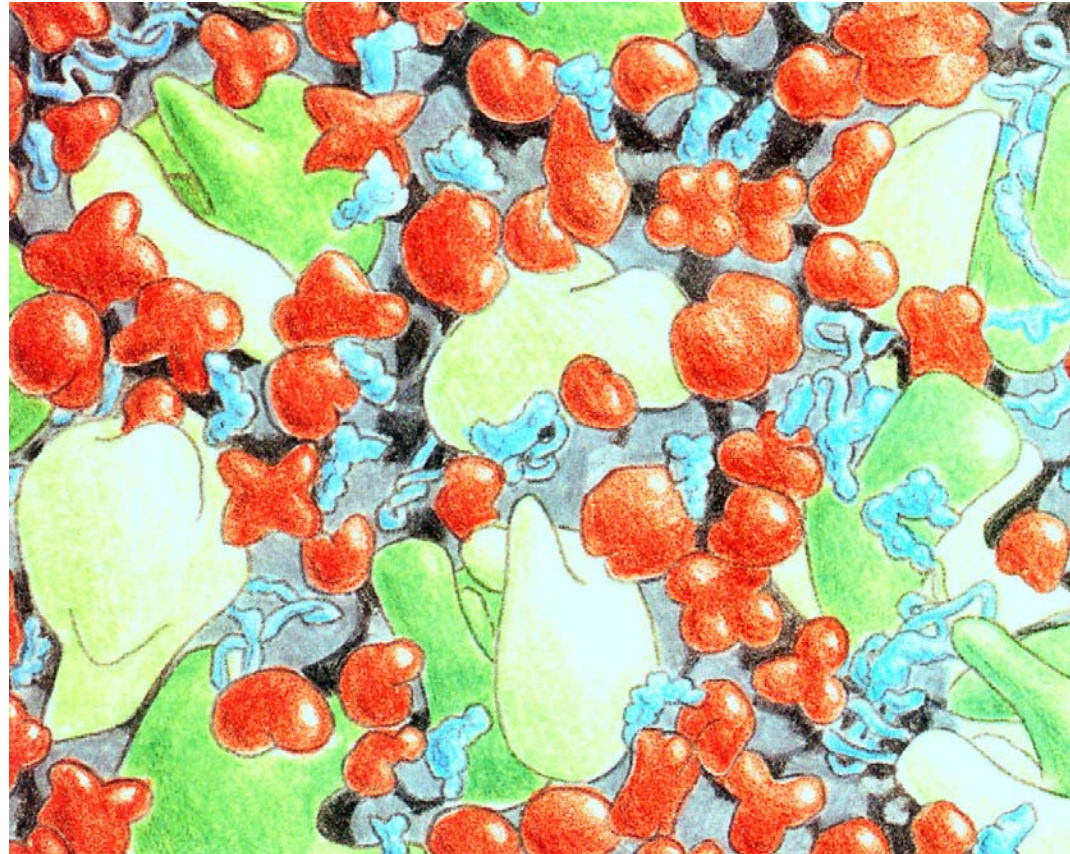
3.4Å

major groove



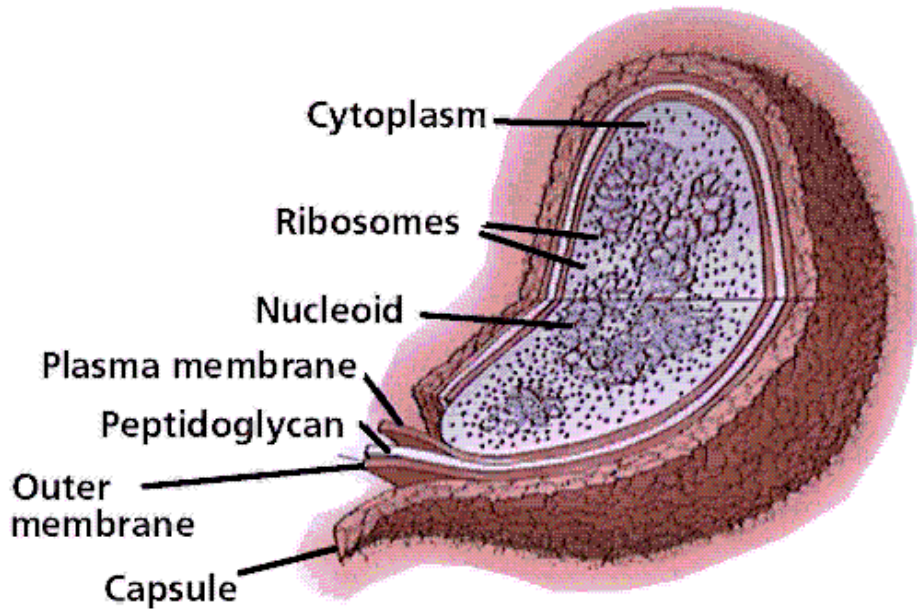
20Å

# Biomolecules in the Cell Cytoplasm

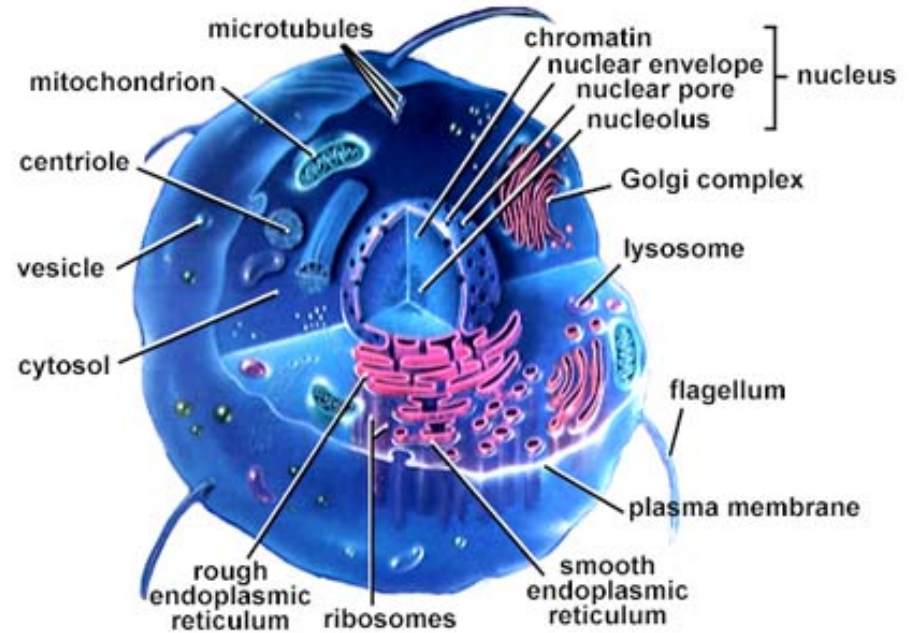


RNAs, ribosomes, and proteins

# Cell Structure



Prokaryotic cell



Eukaryotic cell

# Resources and Reading Assignment

WWW:

<http://www.emc.maricopa.edu/faculty/farabee/BIOBK/BioBookTOC.html>

<http://web.mit.edu/esgbio/www>

<http://users.rcn.com/jkimball.ma.ultranet/BiologyPages>

Textbooks:

Schlick, Chapters 1, 3, 4, and 5

Bourne & Weissig, Chapters 2 and 3

Handouts:

P.G. Debrunner, 1993, Proteins and Nucleic Acids