



Protein Structure

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9/21/11





What do proteins do?

- Chemistry
 - Metabolism
 - Factories of cellular components (Fatty acid synthase)
- Transport
 - Oxygen in blood (hemoglobin)
 - Signals from outside cells (nervous system)
- Structure
 - Internal scaffolding in cells (actin)
 - Hair and nails (collagen)
- Movement
 - Myosin (animal muscles)
 - Flagella and Cilia (cellular propulsion)





How do proteins work?

- Biological Nano Machines
- Highly complex
- But... based on fundamentals of chemistry





Understanding Protein Structure

1. Primary structure
2. Secondary structure
3. Tertiary Structure

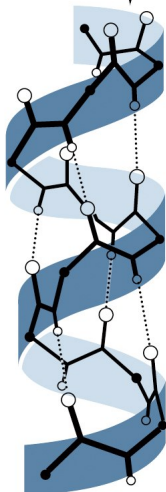


Sequence leads to structure

- Lys - Ala - His - Gly - Lys - Lys - Val - Leu - Gly - Ala -

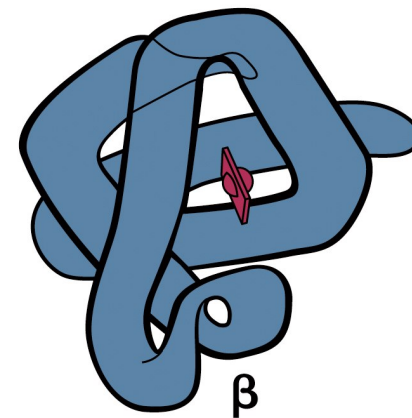
Figure 6-1a Fundamentals of Biochemistry, 2/e
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Primary structure
(amino acid sequence in a polypeptide chain)



Secondary structure
(helix)

Figure 6-1b Fundamentals of Biochemistry, 2/e



Tertiary structure:
one complete protein chain
(β chain of hemoglobin)

Figure 6-1c Fundamentals of Biochemistry, 2/e



Primary Structure - Sequence

- Amino-acid side chains along the Peptide Backbone
- Covalent bonds making a string
- Proteins have many “residues”
 - Proteins can be 100-1000s of residues
 - 20 regular amino acids



Side chains come in 4 “flavors”

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

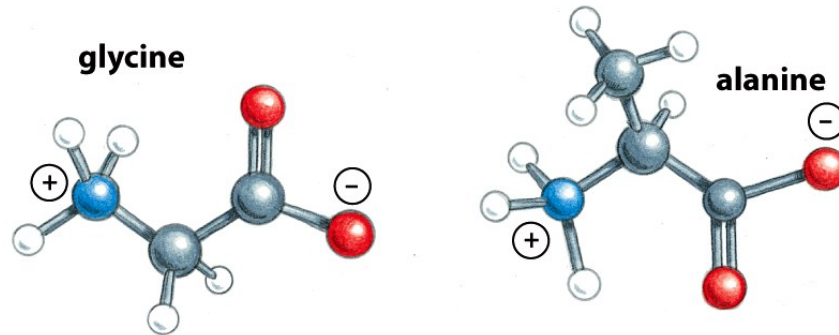
(hydrophilic)

AMINO ACID		SIDE CHAIN	
Alanine	Ala	A	nonpolar
Glycine	Gly	G	nonpolar
Valine	Val	V	nonpolar
Leucine	Leu	L	nonpolar
Isoleucine	Ile	I	nonpolar
Proline	Pro	P	nonpolar
Phenylalanine	Phe	F	nonpolar
Methionine	Met	M	nonpolar
Tryptophan	Trp	W	nonpolar
Cysteine	Cys	C	nonpolar

NONPOLAR AMINO ACIDS

(hydrophobic)

The Backbone



Proteins are long stretches of amino acids

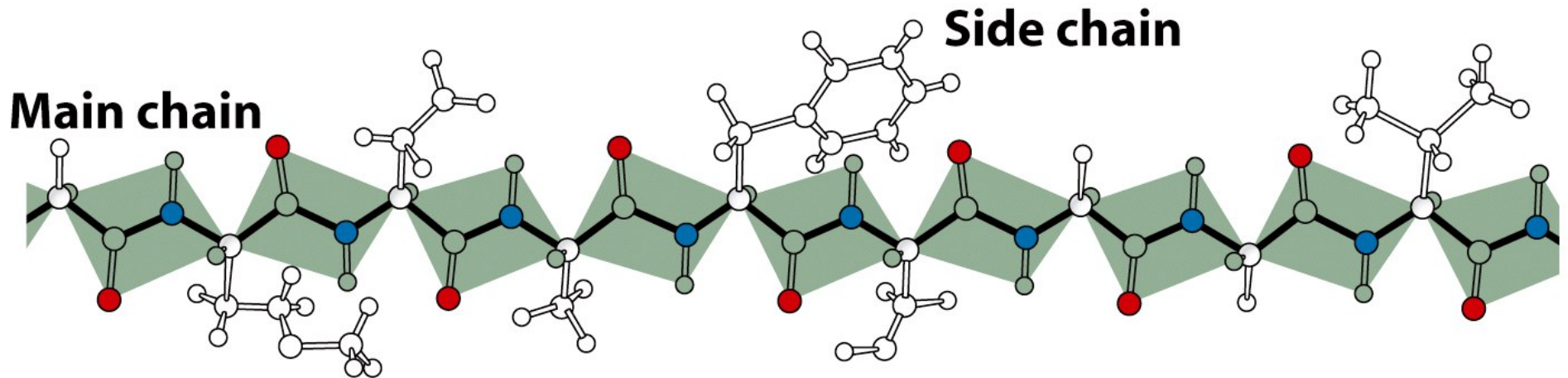


Figure 6-3 Fundamentals of Biochemistry, 2/e
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9 amino acids, 20 possible
=
How many unique sequences?



Understanding Protein Structure

1. Primary structure

2. Secondary structure

3. Tertiary Structure





Secondary Structure

- Proteins have unique structures
 - i.e. not like random mess of string
- Final structure is an assembly of secondary structure elements
- Backbone carbonyl and amide-proton hydrogen bonding are key!





Basic forms of Secondary Structure

- Random Coil
- Alpha-Helix
 - α -helix
- Beta-sheet
 - β -sheet
- Loops



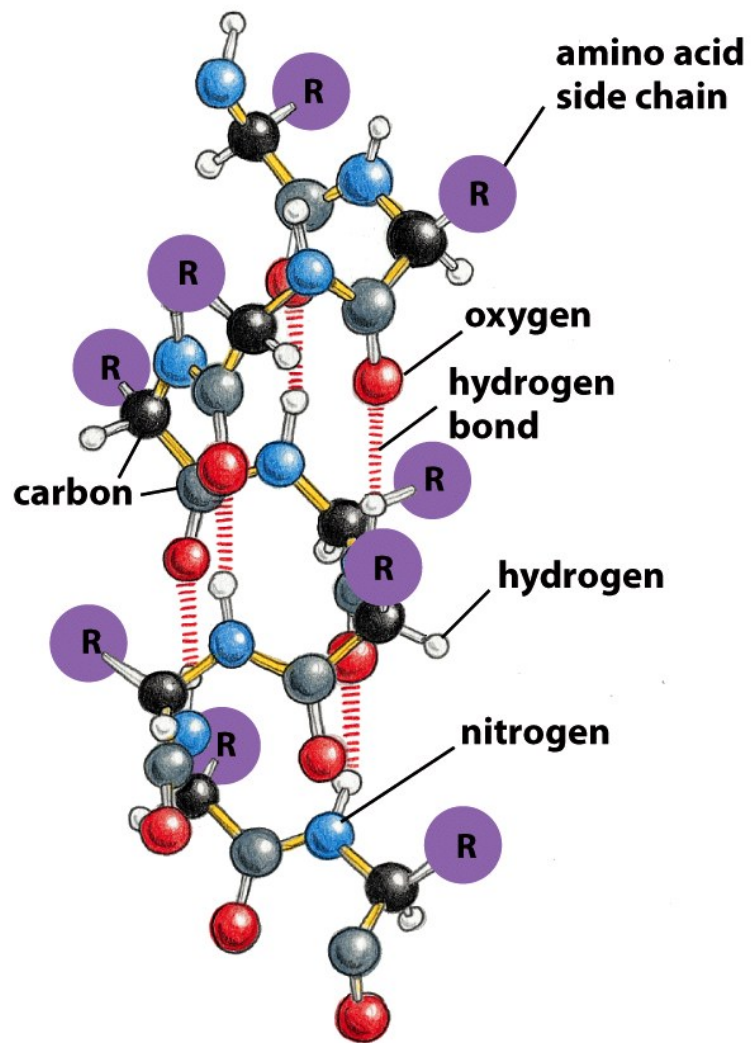


Basic forms of Secondary Structure

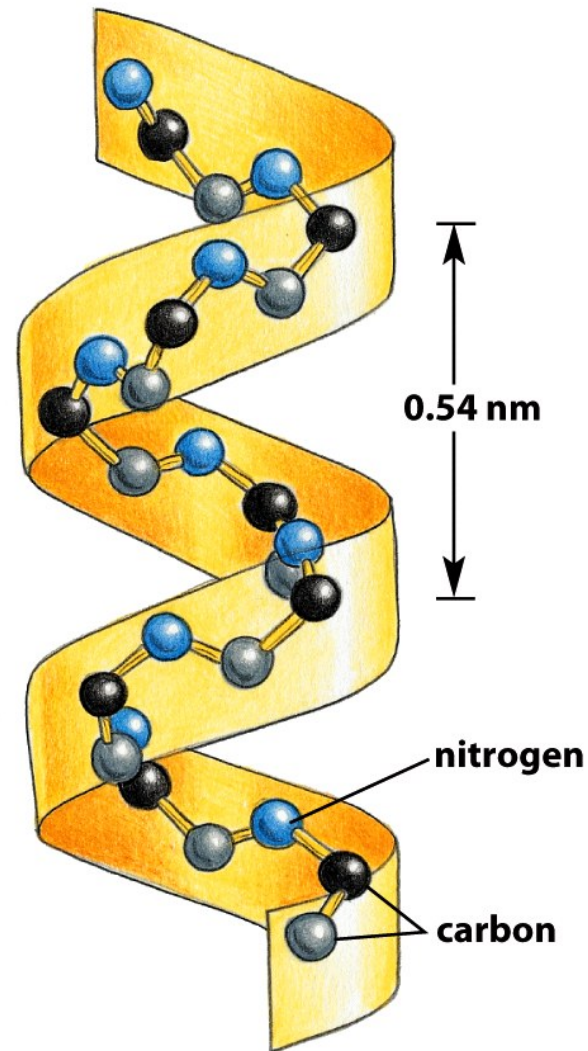
- **Random Coil**
- Alpha-Helix
 - α -helix
- Beta-sheet
 - β -sheet
- Loops



Alpha Helices

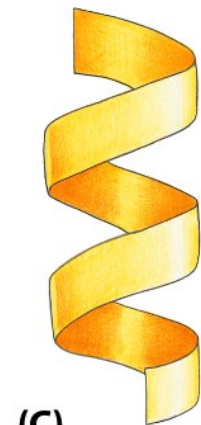


(A)



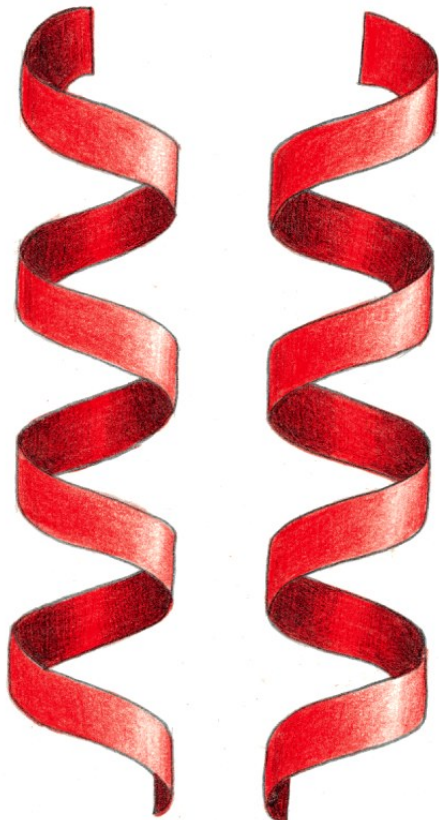
(B)

α helix



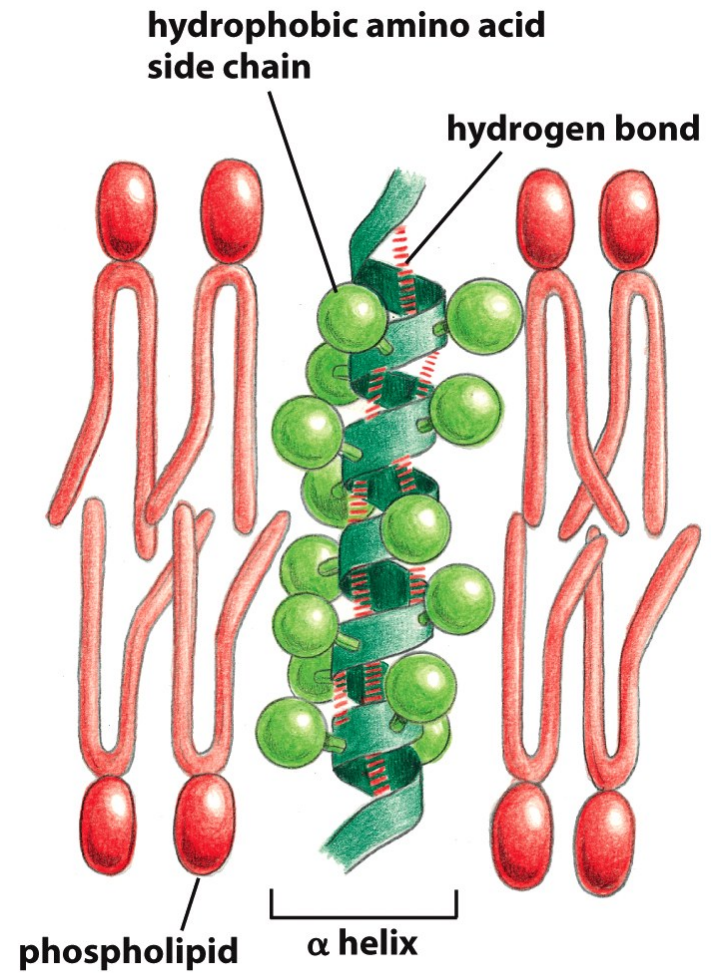
(C)

Two important helical notes

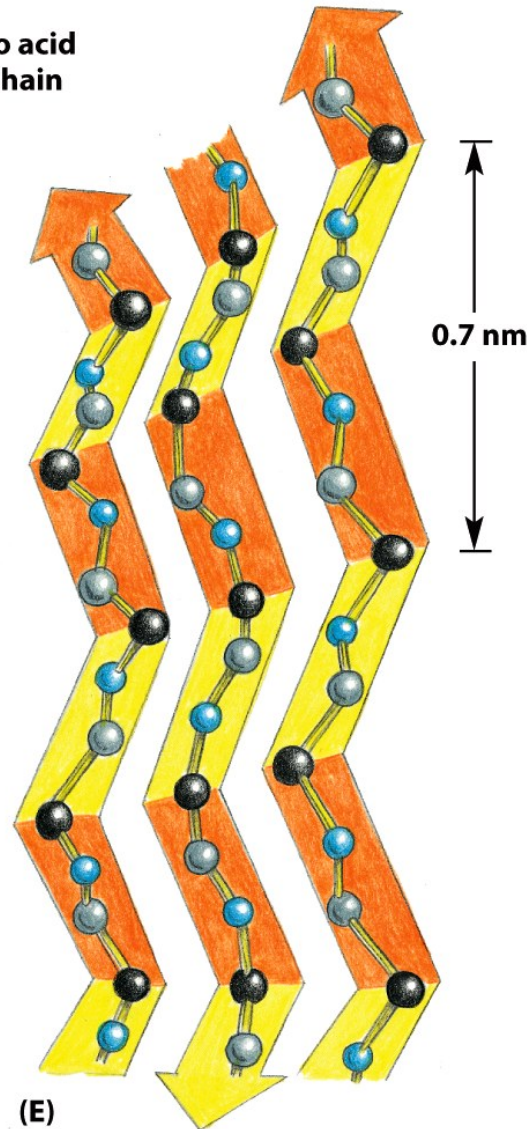
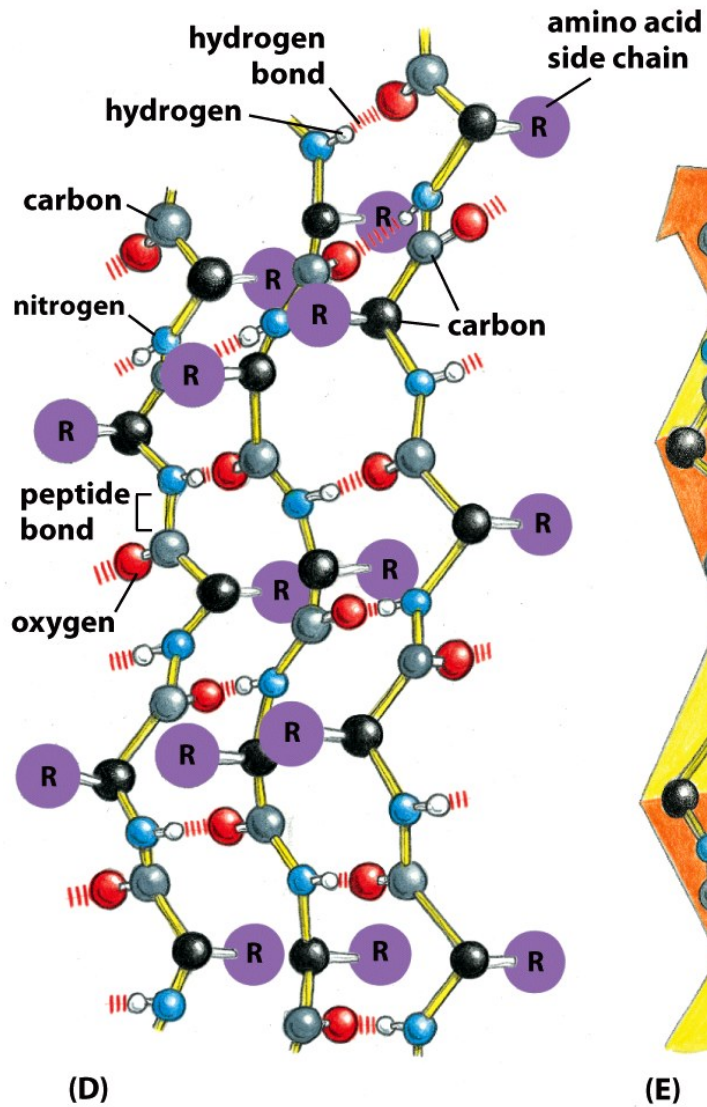


left-handed
(E)

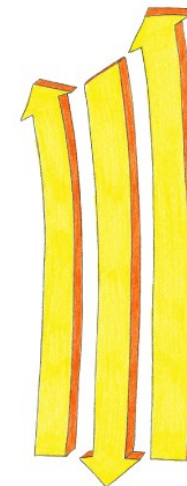
right-handed



Beta Sheets

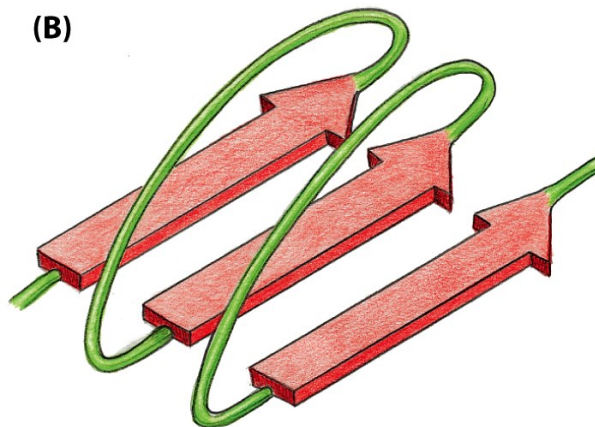
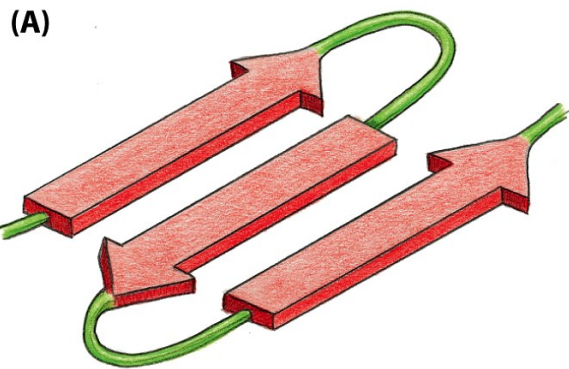


β sheet

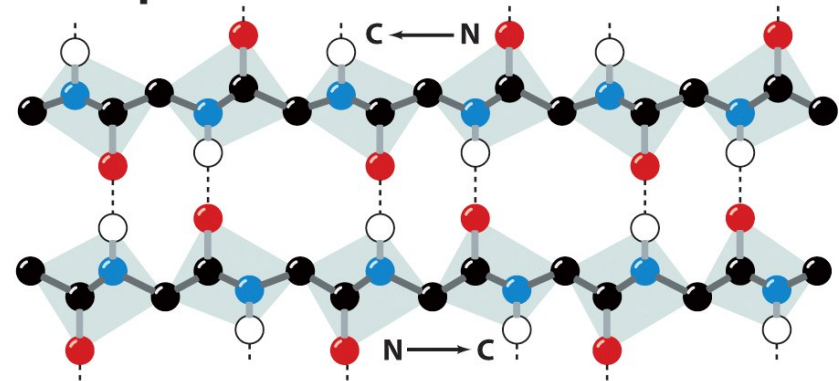


(F)

Beta sheets can assemble different ways



(a) Antiparallel



(b) Parallel

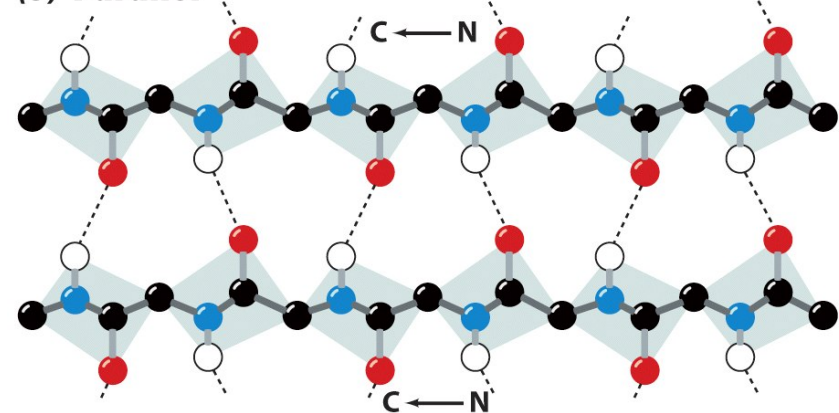



Figure 6-9 Fundamentals of Biochemistry, 2/e



Loops

- Connect helices and sheets
 - Are often structured, but can be very flexible.
 - Glycine often present in loops
 - Mathematica Again
- 



Tertiary Structure

- The assembly of secondary structure
- Side chains and backbone interactions determine how the secondary structure elements align
- Non-covalent bonding primary stabilization
- Only mildly stable, can be broken up by heating or using chemical denaturants



Side Chain Residues Come in 4 “flavors”

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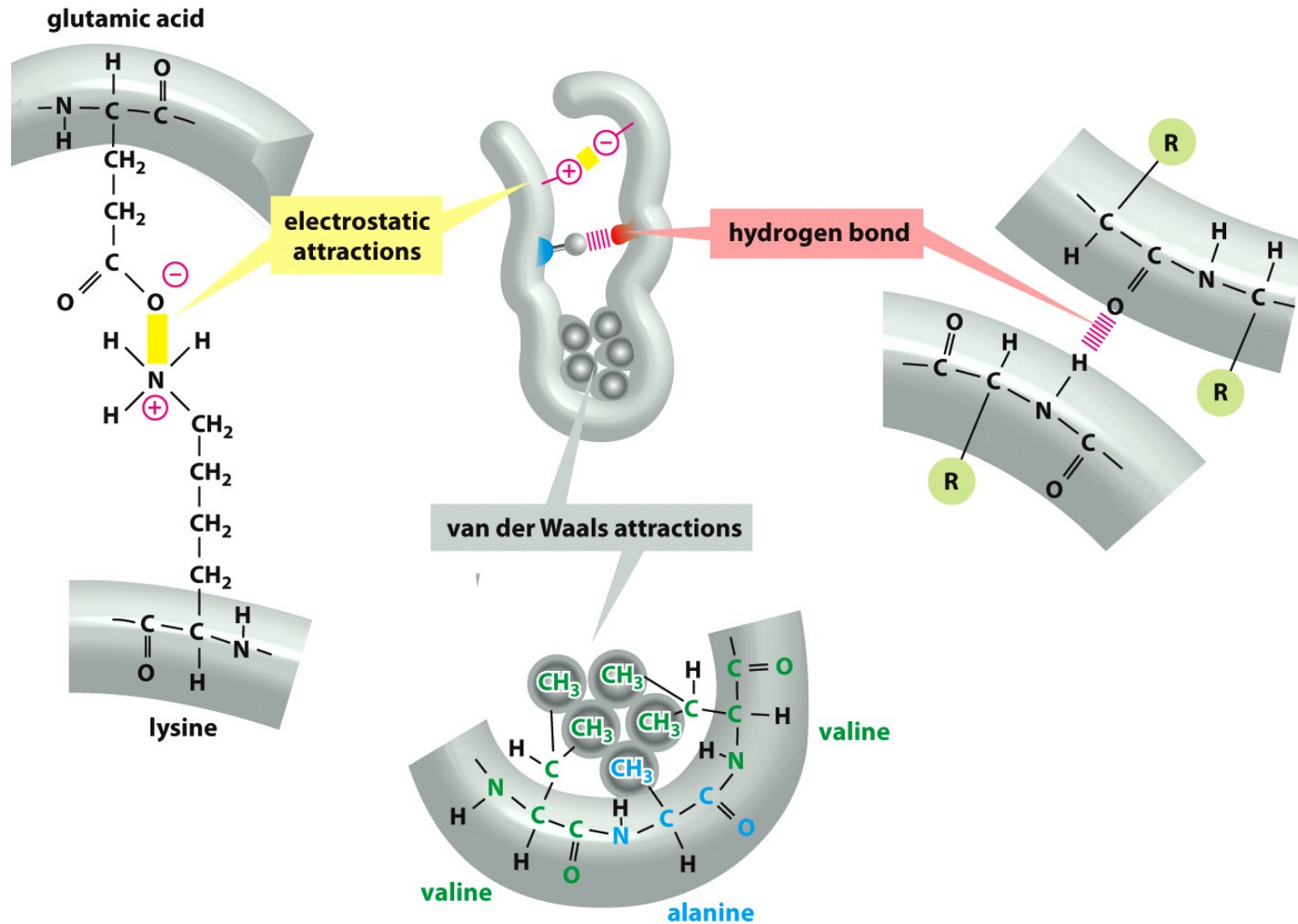
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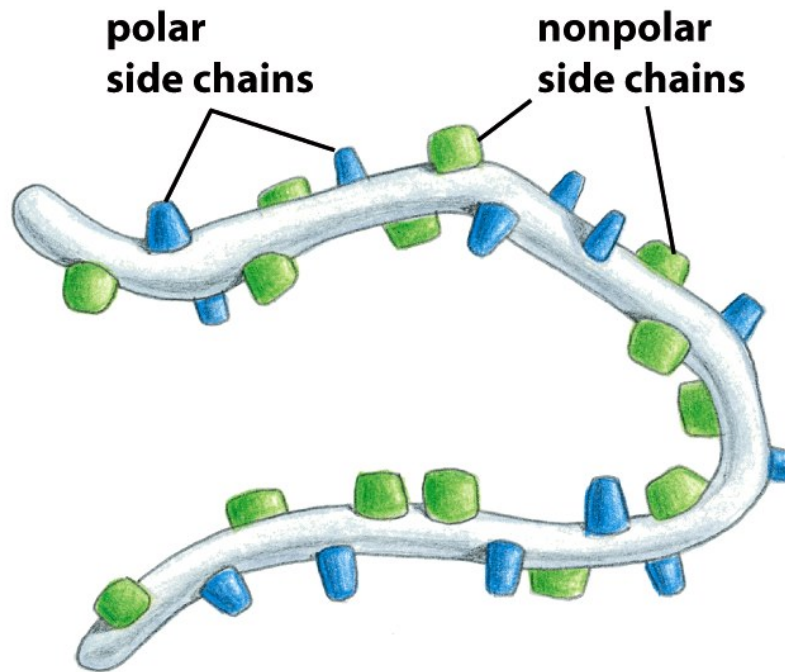
NONPOLAR AMINO ACIDS

(hydrophobic)

Non-covalent interactions hold it together



Protein's "hide" their hydrophobic core



unfolded polypeptide

Disulphide Bonds

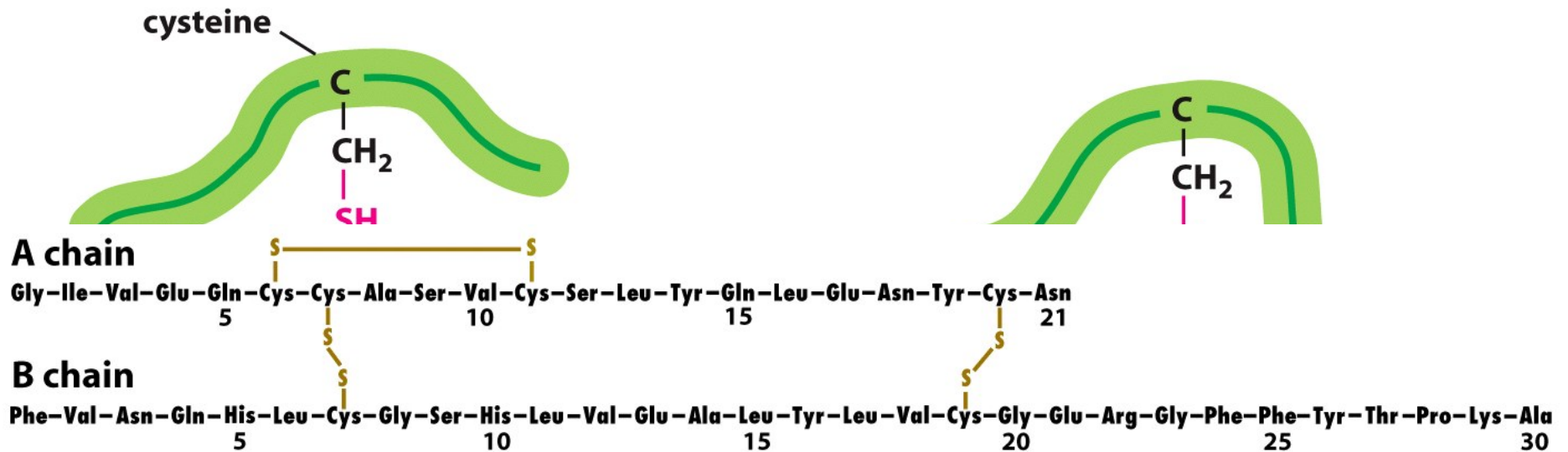
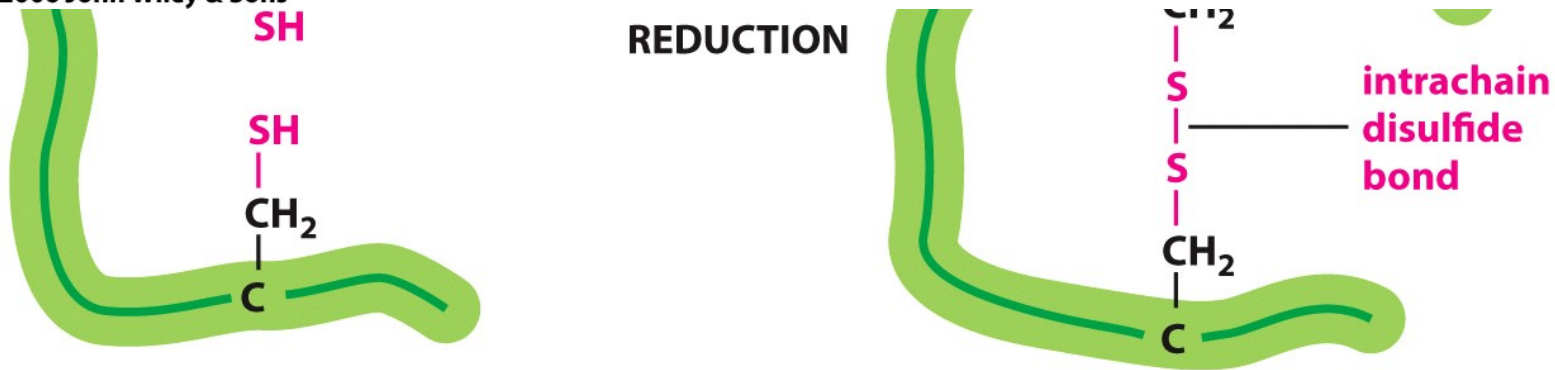


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Relative strength of Bonds

BOND TYPE	LENGTH (nm)	STRENGTH IN kcal/mole	
		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.**

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



The folded state is the
“lowest energy” state

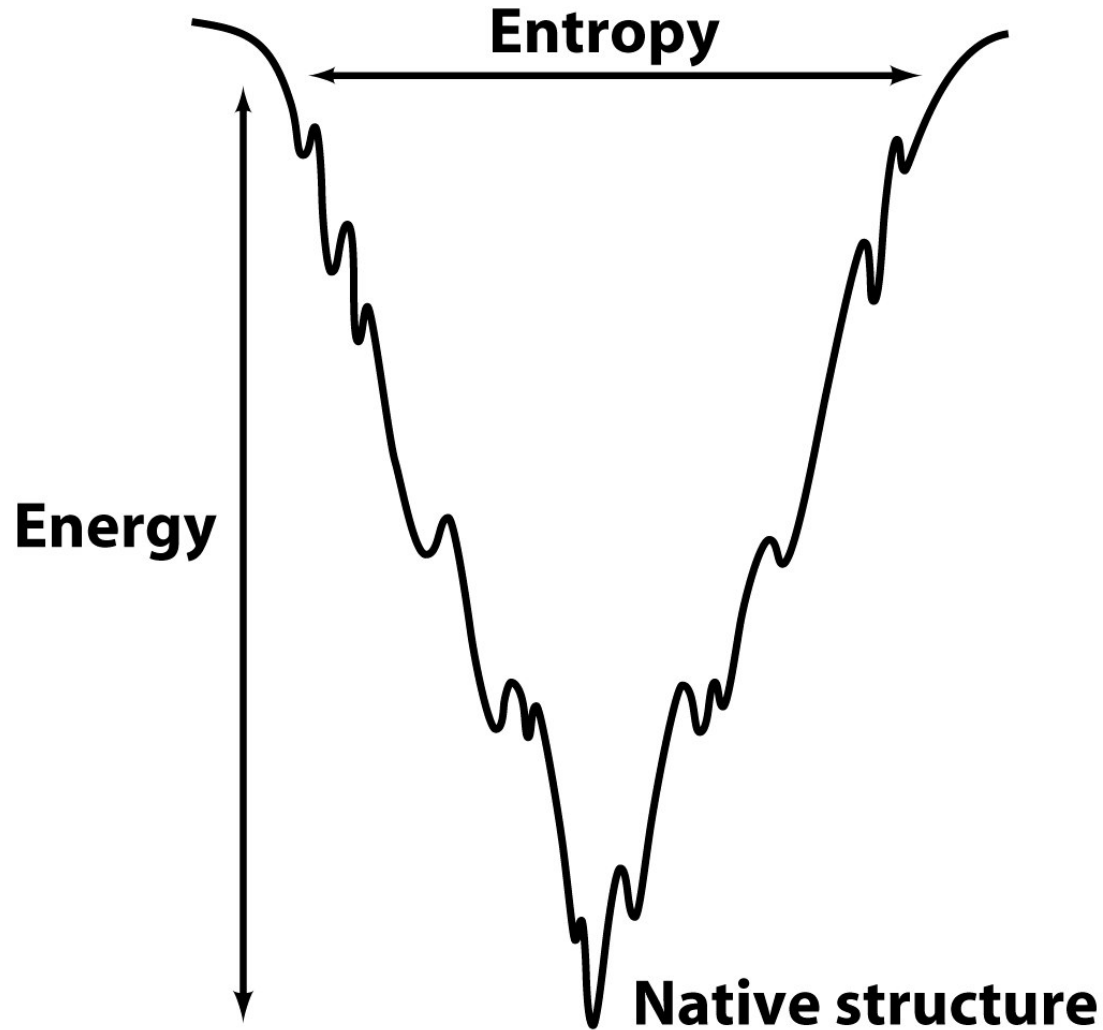


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Tertiary Structure

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└── POLAR AMINO ACIDS ─┘
(hydrophilic)

└── NONPOLAR AMINO ACIDS ─┘
(hydrophobic)

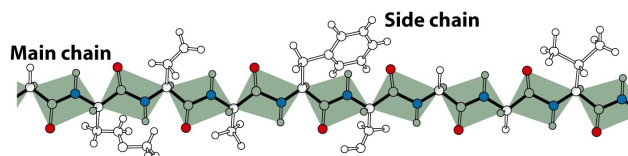
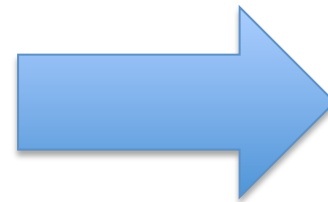
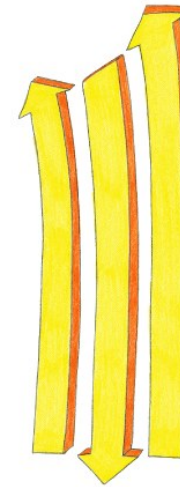


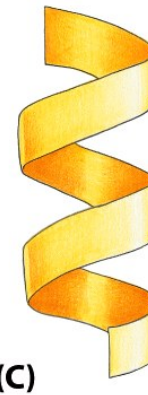
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β sheet



α helix



(c)

(a)



(b)



Figure 6-13 Fundamentals of Biochemistry, 2/e
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From Noodles to Nanomachines

