

#### **Chapter 5**

# The Structure and Function of Large Biological Molecules

Lecture Presentations by Nicole Tunbridge and Kathleen Fitzpatrick

#### The Molecules of Life

- All living things are made up of four classes of large biological molecules: *carbohydrates*, *lipids*, *proteins*, and *nucleic acids*
- Macromolecules are large molecules and are complex
- Large biological molecules have unique properties that arise from the orderly arrangement of their atoms (Structure-function)

Figure 5.1a



The scientist in the foreground is using 3-D glasses to help her visualize the structure of the protein displayed on her screen.

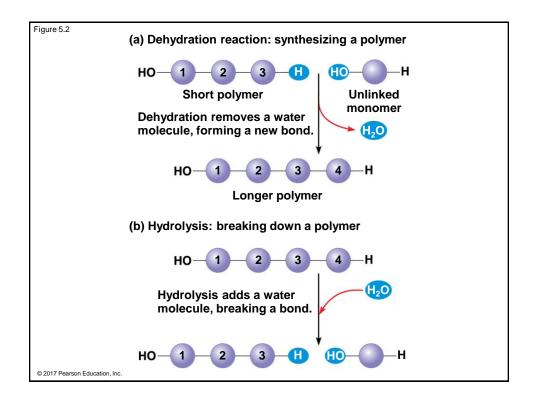
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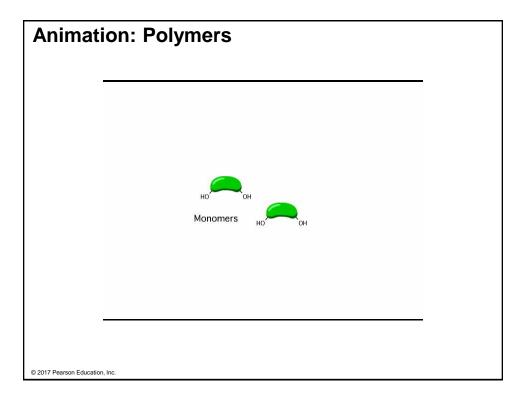
# Concept 5.1: Macromolecules are polymers, built from monomers

- A polymer is a long molecule consisting of many similar building blocks
- The repeating units that serve as building blocks are called monomers
- Carbohydrates, proteins, and nucleic acids are polymers
- Lipids are not polymers

#### The Synthesis and Breakdown of Polymers

- Enzymes are specialized macromolecules that speed up chemical reactions such as those that make or break down polymers
- A dehydration reaction occurs when two monomers bond together through the loss of a water molecule (SYNTHESIS)
- Polymers are disassembled to monomers by hydrolysis, a reaction that is essentially the reverse of the dehydration reaction (BREAKDOWN)



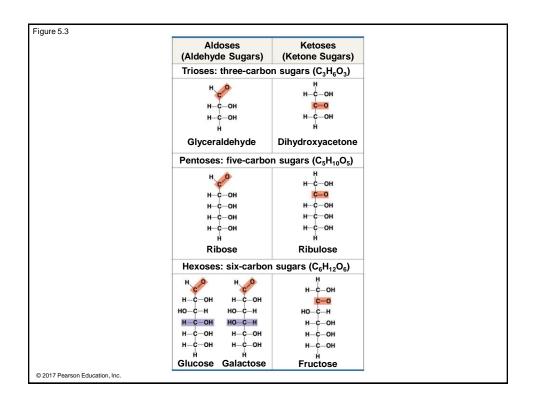


# Concept 5.2: Carbohydrates serve as fuel and building material

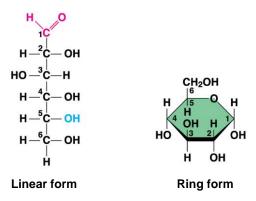
- Carbohydrates include <u>sugars</u> and the <u>polymers of</u> <u>sugars</u>
- The simplest carbohydrates are monosaccharides, or simple sugars
- Carbohydrate macromolecules are polysaccharides, polymers composed of many sugar building blocks

#### **Sugars**

- Monosaccharides serve as a major fuel for cells and as raw material for building molecules
- Monosaccharides have molecular formulas that are usually multiples of CH<sub>2</sub>O
- Glucose (C<sub>6</sub>H<sub>12</sub>O<sub>6</sub>) is the most common monosaccharide
- Monosaccharides are classified by
  - The location of the carbonyl group (as <u>aldose</u> or <u>ketose</u>)
  - The number of carbons in the carbon skeleton (<u>triose</u>, <u>tetrose</u>, <u>pentose</u>, <u>hexose</u>, etc.)

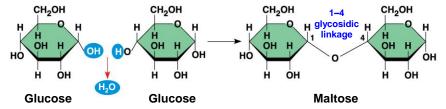


In aqueous solutions many sugars form rings

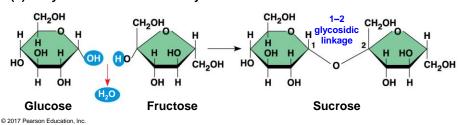


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- A disaccharide is formed when a <u>dehydration reaction</u> joins two monosaccharides
- This covalent bond is called a glycosidic linkage (bond)
  - (a) Dehydration reaction in the synthesis of maltose



(b) Dehydration reaction in the synthesis of sucrose



#### **Polysaccharides**

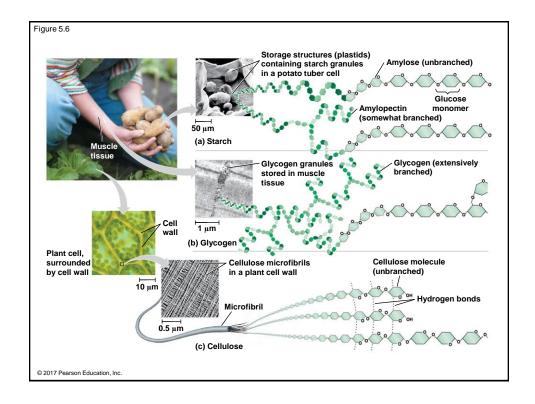
- Polysaccharides, the polymers of sugars, have storage and structural roles
- The architecture and function of a polysaccharide are determined by: <u>its sugar monomers</u> and <u>the</u> <u>positions of its glycosidic linkages</u>

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#### Storage Polysaccharides

- Starch is a storage polysaccharide of plants
- Consists of glucose monomers
- Plants store extra starch as granules within chloroplasts and other plastids
- The simplest form of starch is amylose
- Glycogen is a storage polysaccharide in animals
- Consists of glucose monomers
- Glycogen is stored as granules in liver and muscle
- Hydrolysis of glycogen in these cells releases glucose when the demand for sugar increases

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#### Structural Polysaccharides

- The polysaccharide cellulose is a major component of the tough wall of plant cells
- Like starch and glycogen, cellulose is a polymer of glucose, but the glycosidic linkages are different
- The difference is based on two ring forms for glucose: alpha (α) and beta (β)

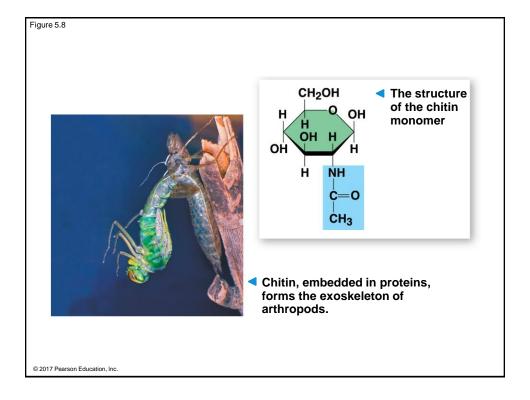
Figure 5.7b CH<sub>2</sub>OH CH<sub>2</sub>OH CH<sub>2</sub>OH CH<sub>2</sub>OH ОН ОН (b) Starch and glycogen: 1–4 linkage of a glucose monomers CH<sub>2</sub>OH CH<sub>2</sub>OH ОН ОН OH ÓН HÓ ÓН CH<sub>2</sub>OH OH CH<sub>2</sub>OH (c) Cellulose: 1-4 linkage of β glucose monomers

- Starch (α configuration) is largely helical
- Cellulose molecules (β configuration) are straight and unbranched
- Some hydroxyl groups on the monomers of cellulose can hydrogen-bond with hydroxyls of parallel cellulose molecules

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- Enzymes that digest starch by hydrolyzing α linkages can't hydrolyze β linkages in cellulose
- The cellulose in human food passes through the digestive tract as "insoluble fiber"
- Some microbes use enzymes to digest cellulose
- Many herbivores, from cows to termites, have symbiotic relationships with these microbes

- **Chitin**, another structural polysaccharide, is found in the exoskeleton of arthropods (حشرات)
- Chitin also provides structural support for the cell walls of many fungi



#### Class activity!

- How many molecules of water are needed to completely hydrolyze a polymer that is ten monomers long?
- ➤ If you eat a piece of fish, what reactions must occur for the amino acid monomers in the protein of the fish to be converted to new proteins in your body?

#### Class activity!

➤ A dehydration reaction joins 2 molecules of glucose to form maltose. The formula for glucose is C6H12O6. What is the formula for maltose?

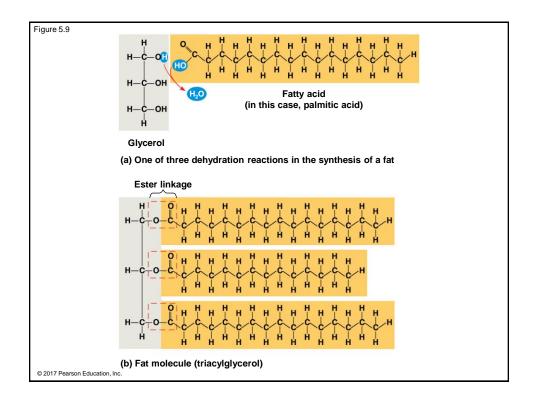
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# Concept 5.3: Lipids are a diverse group of hydrophobic molecules

- Lipids are the only class of large biological molecules that does not include true polymers
- The unifying feature of lipids is that they do not like water and don't mix well with it (HYDROPHOBIC)
- Lipids consist mostly of hydrocarbon regions
- The most biologically important lipids are fats, phospholipids, and steroids

#### **Fats**

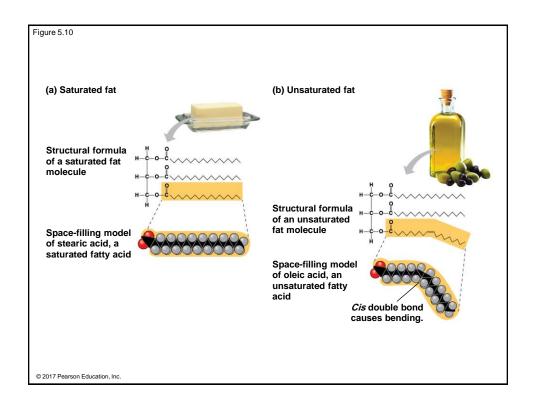
- Fats are made from two types of smaller molecules: <u>glycerol</u> and <u>fatty acids</u>
- Glycerol is a three-carbon alcohol with a hydroxyl group attached to each carbon
- A fatty acid consists of a carboxyl group attached to a long carbon skeleton



- Fats separate from water because water molecules hydrogen-bond to each other and exclude the fats
- In a fat, three fatty acids are joined to glycerol by an ester linkage, creating a triacylglycerol, or triglyceride
- The fatty acids in a fat can be all the same or different

#### **Animation: Fats**

- Fatty acids vary in length (number of carbons) and in the number and locations of double bonds
- Saturated fatty acids have the maximum number of hydrogen atoms possible and no double bonds
- Unsaturated fatty acids have one or more double bonds

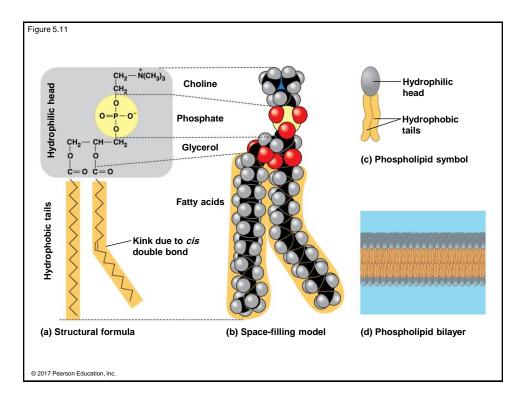


- Fats made from saturated fatty acids are called saturated fats and are solid at room temperature
- Most animal fats are saturated
- Fats made from unsaturated fatty acids are called unsaturated fats or oils and are *liquid* at room temperature
- Plant fats and fish fats are usually unsaturated

- The major function of fats is energy storage
- Humans and other mammals store their long-term food reserves in adipose cells
- Adipose tissue also cushions vital organs and insulates the body

#### **Phospholipids**

- In a phospholipid, two fatty acids and a phosphate group are attached to glycerol
- The two fatty acid tails are hydrophobic, but the phosphate group and its attachments form a hydrophilic head
- The molecule is therefore AMPHIPATHIC (both hydrophobic and hydrophilic)



- When phospholipids are added to water, they self-assemble into double-layered sheets called bilayers
- At the surface of a cell, phospholipids are also arranged in a bilayer, with the hydrophobic tails pointing toward the interior
- The phospholipid bilayer forms a boundary between the cell and its external environment

#### **Steroids**

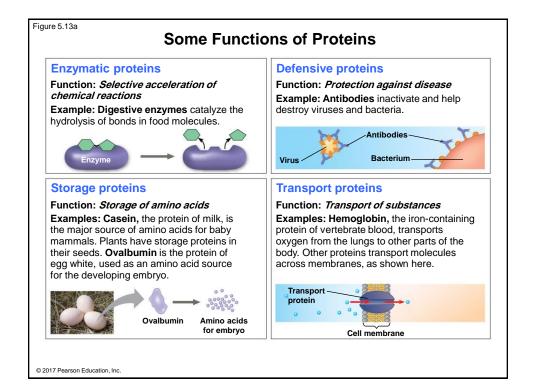
- Steroids are lipids characterized by a carbon skeleton consisting of four fused rings
- Cholesterol, a type of steroid, is a component in animal cell membranes and a precursor from which other steroids are synthesized
- A high level of cholesterol in the blood may contribute to cardiovascular disease
- Some hormones, such as sex hormones, are made from cholesterol

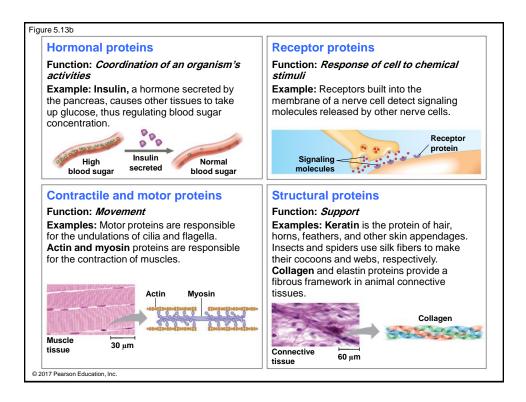
#### Class activity!

- Why are human sex hormones are considered lipids?
- Suppose a membrane surrounded an oil droplet, as it does in the cells of plant seeds and in some animal cells. Describe and explain the form it might take?

# Concept 5.4: Proteins include a diversity of structures, resulting in a wide range of functions

- Proteins make more than 50% of the dry mass of most cells
- Functions:
- Some proteins <u>speed up</u> chemical reactions
   (<u>Enzymes</u> act as <u>catalysts</u> to accelerate reactions)
- Other protein functions include defense, storage, transport, cellular communication, movement, and structural support



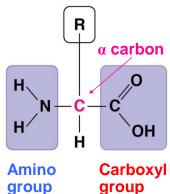


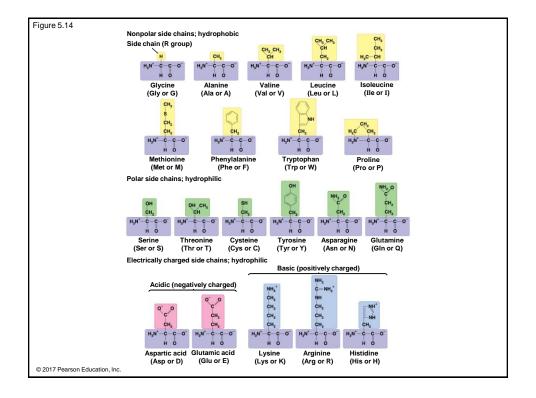
- Proteins are all made from the same set of 20 amino acids
- Polypeptides are <u>unbranched</u> polymers built from these amino acids
- A protein is a biologically functional molecule that consists of one or more polypeptides

#### **Amino Acid Monomers**

- Amino acids are organic molecules with <u>amino</u> and <u>carboxylic acid</u> groups
- Amino acids differ in their properties due to differing side chains, called <u>R groups</u>

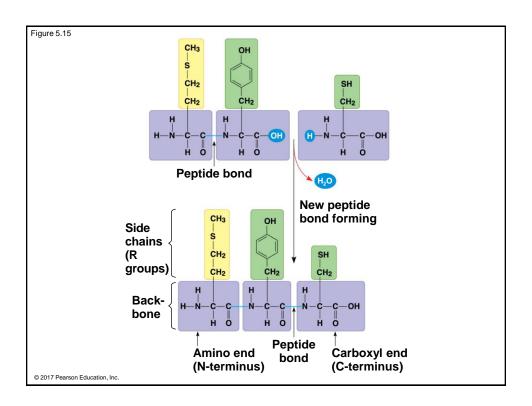
#### Side chain (R group)





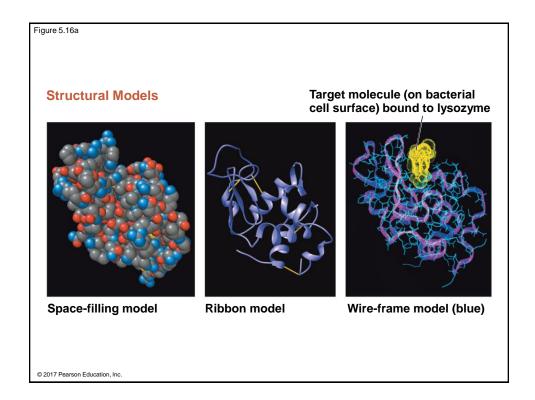
#### **Polypeptides (Amino Acid Polymers)**

- Amino acids are joined by <u>covalent bonds</u> called peptide bonds
- A polypeptide is a polymer of amino acids
- Polypeptides range in length from a few to more than 1,000 monomers
- Each polypeptide has a unique linear sequence of amino acids, with a carboxyl end (C-terminus) and an amino end (N-terminus)

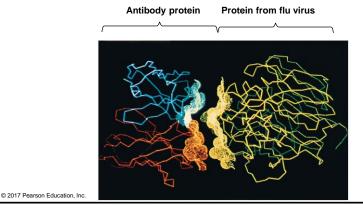


#### **Protein Structure and Function**

- The specific activities of proteins are due to their detailed three-dimensional architecture
- A functional protein consists of one or more polypeptides precisely twisted, folded, and coiled into a unique shape

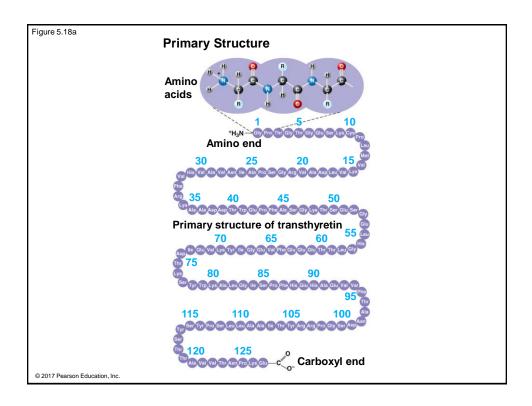


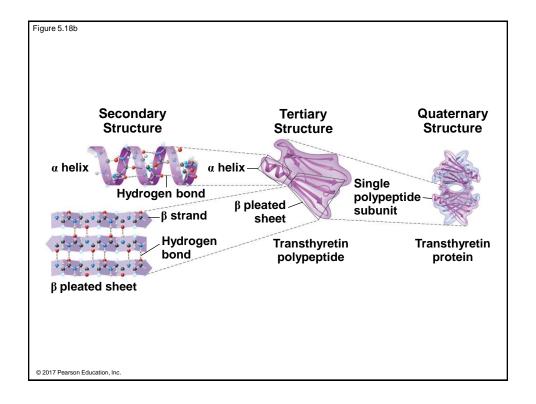
- The <u>amino acids sequence</u> determines a <u>protein's</u> three-dimensional structure
- A <u>protein's structure</u> determines how it works (function)
- The function of a protein usually depends on its ability to recognize and bind to other molecules



#### Four Levels of Protein Structure

- The primary structure of a protein is its unique sequence of amino acids
- Secondary structure, found in most proteins, consists of coils and folds in the polypeptide chain
- Tertiary structure is determined by interactions among various side chains (R groups)
- Quaternary structure results when a protein consists of multiple polypeptide chains





- The primary structure of a protein is its <u>sequence</u> of amino acids
- Primary structure is like the <u>order of letters in a long</u> word
- Primary structure is determined <u>by inherited genetic</u> <u>information</u>
- E.g.
   H<sub>3</sub>N<sup>+</sup>-Met-Tyr-Trp-Ala-Pro-Lys-Ala-Gly-Asp-COO<sup>-</sup>

H<sub>3</sub>N<sup>+</sup>–Asp-Gly-Ala-Lys-Pro-Ala-Trp-Tyr-Met–COO<sup>-</sup>

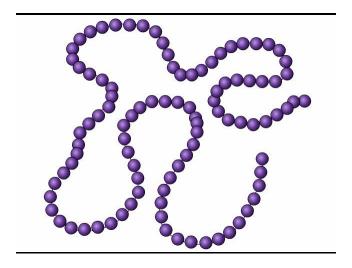
(Remember: RAT and TAR are different)

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Is different from:

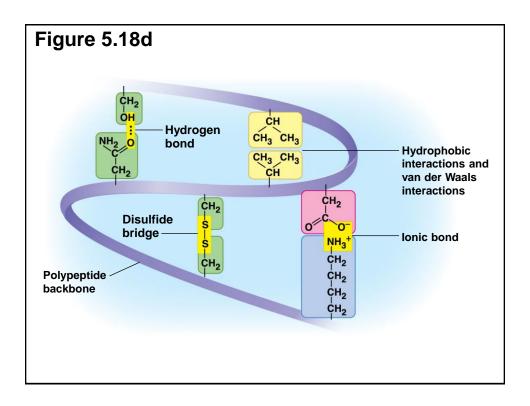
- The coils and folds of secondary structure result from hydrogen bonds between <u>repeating units of the</u> <u>polypeptide backbone</u>
- Typical secondary structures are a coil called an α helix and a folded structure called a β pleated sheet

#### **Animation: Secondary Protein Structure**

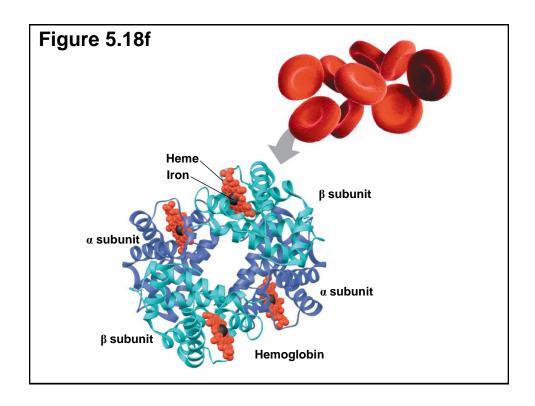


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- Tertiary structure, the <u>overall shape</u> of a polypeptide, results from <u>interactions between</u> <u>R groups</u>, (not interactions between backbone units)
- These interactions include hydrogen bonds, ionic bonds, hydrophobic interactions, and van der Waals interactions
- <u>Strong covalent bonds</u> called disulfide bridges may strengthen the protein's structure

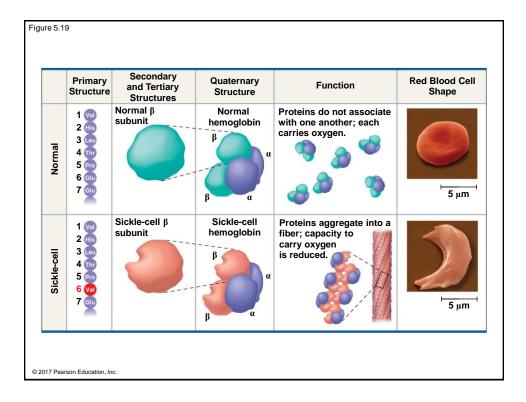


- Quaternary structure results when <u>two or more</u> <u>polypeptide chains form one macromolecule</u>
- E.g.
- *Collagen* is a fibrous protein consisting of three polypeptides coiled like a rope
- *Hemoglobin* is a globular protein consisting of four polypeptides: two  $\alpha$  and two  $\beta$  subunits



## Sickle-Cell Disease: A Change in Primary Structure

- A slight change in primary structure can affect a protein's structure and ability to function
- Sickle-cell disease, an inherited blood disorder, results from <u>a single amino acid substitution</u> in the protein hemoglobin
- The abnormal hemoglobin molecules cause the red blood cells to aggregate into chains and to deform into a sickle shape



#### Class activity!

- ➤ In a DNA double helix, a region along one strand has this sequence 5'-TAGGCCT-3'.
- Copy this sequence and write down the complementary strand, clearly indicating the orientation of the 2 strands.

#### Homework!

➤ Comparisons of amino acid sequences can shed light of the evolutionary divergence of related species. If you were comparing two living species, would you expect all proteins to show the same degree of divergence? Why or Why no?

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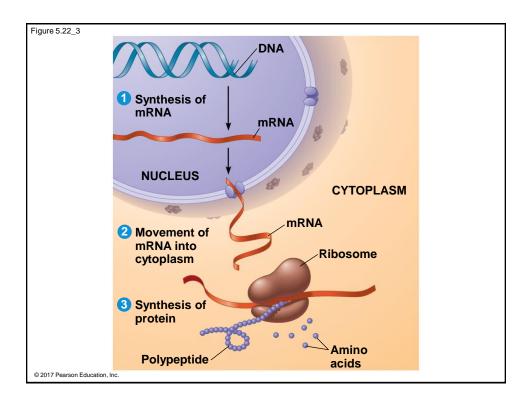
# Concept 5.5: Nucleic acids store, transmit, and help express hereditary information

- The amino acid sequence of a polypeptide is programmed by a unit of inheritance called a gene
- Genes consist of DNA, a nucleic acid made of monomers called nucleotides

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#### The Roles of Nucleic Acids

- There are two types of nucleic acids
  - Deoxyribonucleic acid (DNA)
  - Ribonucleic acid (RNA)
- DNA provides directions for its own replication
- DNA directs synthesis of messenger RNA (mRNA) and, through mRNA, controls protein synthesis
- This process is called gene expression



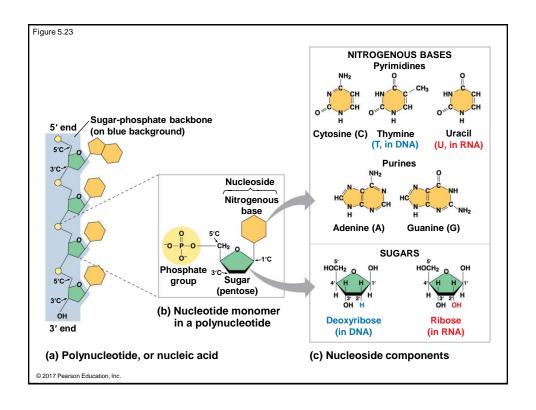
- Each gene along a DNA molecule directs synthesis of a messenger RNA (mRNA)
- The mRNA molecule interacts with the cell's proteinsynthesizing machinery to direct production of a polypeptide
- The flow of genetic information can be summarized as DNA → RNA → protein

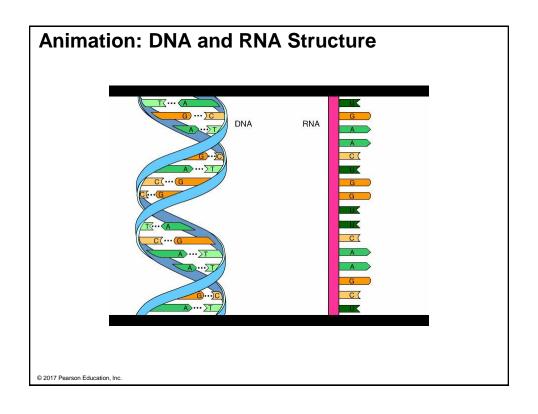
#### The Components of Nucleic Acids

- Nucleic acids are polymers called polynucleotides
- Each polynucleotide is made of monomers called nucleotides
- Each nucleotide consists of a nitrogenous base, a pentose sugar, and one or more phosphate groups
- The portion of a nucleotide without the phosphate group is called a nucleoside (nitrogenous base + pentose sugar only)
- Nucleoside = nitrogenous base + sugar
- Nucleotide = nucleoside + phosphate group

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- There are two families of nitrogenous bases
  - Pyrimidines (cytosine (C), thymine (T), and uracil (U))
     have a single six-membered ring
  - Purines (adenine (A) and guanine (G)) have a sixmembered ring fused to a five-membered ring
- In DNA, the sugar is deoxyribose; in RNA, the sugar is ribose





#### **Nucleotide Polymers**

- Nucleotides are linked together by phosphodiester linkages to build a polynucleotide
- A phosphodiester linkage consists of a <u>phosphate</u> group that links the sugars of two nucleotides
- These links create a backbone of sugarphosphate units with nitrogenous bases as appendages
- The <u>sequence of bases</u> along a DNA or mRNA polymer is unique for each gene

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#### The Structures of DNA and RNA Molecules

- DNA molecules have two polynucleotides spiraling around an imaginary axis, forming a double helix
- The backbones <u>run in opposite 5' → 3' directions</u> from each other, an arrangement referred to as antiparallel
- One DNA molecule includes many genes

- Only certain bases in DNA pair up and form hydrogen bonds: adenine (A) always with thymine (T), and guanine (G) always with cytosine (C)
- This is called complementary base pairing
- This feature of DNA structure makes it possible to generate two identical copies of each DNA molecule in a cell preparing to divide

- RNA, in contrast to DNA, is <u>single-stranded</u>
- Complementary pairing can also occur between two RNA molecules or between parts of the same molecule
- In RNA, thymine is replaced by uracil (U), so
   A can pair with U
- RNA can also have other base pairing
- While DNA always exists as a double helix, RNA molecules are more variable in form

#### **DNA and RNA Base Pairing Rules**

#### **DNA to DNA**

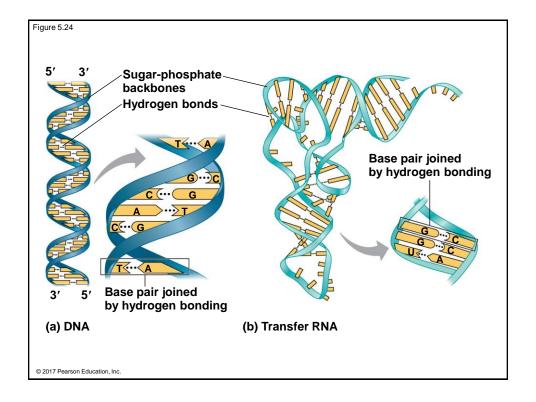
- Possible Bases: Adenine, Thymine, Cytosine, Guanine
- G↔C, A↔T
- A and G are purines (double-ring), C and T are pyrimidines (single-ring)

#### DNA to mRNA

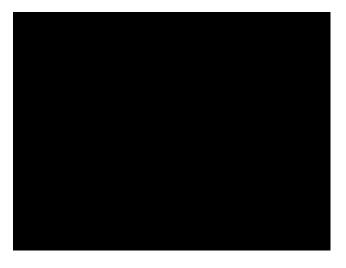
- Possible Bases: Adenine, Thymine, Cytosine, Guanine, Uracil (RNA only)
- G↔C, A→U, T→A
- A and G are purines (double-ring), C, T, and U are pyrimidines (single-ring)

#### mRNA to tRNA

- · Possible Bases: Adenine, Cytosine, Guanine, Uracil
- G↔C, A↔U
- A and G are purines (double-ring), C, T, and U are pyrimidines (single-ring)







#### **Concept 5.6: Genomics and proteomics have** transformed biological inquiry and applications

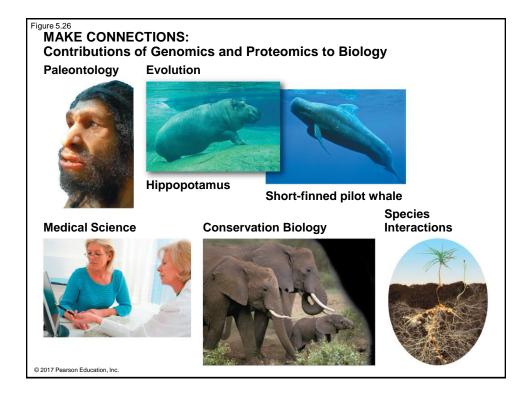
- Once the structure of DNA and its relationship to amino acid sequence was understood, biologists wanted to "decode" genes by learning their base sequences
- The first chemical techniques for DNA sequencing were developed in the 1970s and made better over the next 20 years
- The rapid development of faster and less expensive methods of sequencing was a side effect of the **Human Genome Project**
- Many genomes (including human) have been sequenced, generating large sets of data

Figure 5.25



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- Bioinformatics uses <u>computer software</u> and other <u>computational tools</u> to deal with the data resulting from sequencing many genomes
- Analyzing large sets of genes or even comparing whole genomes of different species is called genomics
- A similar analysis of large sets of proteins including their sequences is called **proteomics**



### **DNA** and Proteins as Tape Measures of Evolution

- Sequences of genes and their protein products document the *hereditary background* of an organism
- Linear sequences of DNA molecules are passed from parents to offspring
- We can extend the concept of "molecular genealogy" to relationships between species
- Molecular biology has added a new measure to the toolkit of evolutionary biology

Figure 5.UN02a



> Rhesus monkey



> Gibbon



Species	Alignment of Amino Acid Sequences of β-globin						
Human	1	VHLTPEEKSA	VTALWGKVNV	DEVGGEALGR	LLVVYPWTQR	FFESFGDLST	
Monkey	1	VHLTPEEKNA	VTTLWGKVNV	DEVGGEALGR	LLLVYPWTQR	FFESFGDLSS	
Gibbon	1	VHLTPEEKSA	VTALWGKVNV	DEVGGEALGR	LLVVYPWTQR	FFESFGDLST	
Human	51	PDAVMGNPKV	KAHGKKVLGA	FSDGLAHLDN	LKGTFATLSE	LHCDKLHVDP	
Monkey	51	PDAVMGNPKV	KAHGKKVLGA	FSDGLNHLDN	LKGTFAQLSE	LHCDKLHVDP	
Gibbon	51	PDAVMGNPKV	KAHGKKVLGA	FSDGLAHLDN	LKGTFAQLSE	LHCDKLHVDP	
Human	101	ENFRLLGNVL	VCVLAHHFGK	EFTPPVQAAY	QKVVAGVANA	LAHKYH	
Monkey	101	ENFKLLGNVL	VCVLAHHFGK	EFTPQVQAAY	QKVVAGVANA	LAHKYH	
Gibbon	101	ENFRLLGNVL	VCVLAHHFGK	EFTPQVQAAY	QKVVAGVANA	LAHKYH	

Data from Human: http://www.ncbi.nlm.nih.gov/protein/AAA21113.1; rhesus monkey: http://www.ncbi.nlm.nih.gov/protein/122634; gibbon: http://www.ncbi.nlm.nih.gov/protein/122616

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Figure 5.UN04

#### **Summary**

Components	Examples	Functions	
СН₂ОН Н С Н	Monosaccharides: glucose, fructose	Fuel; carbon sources that can be converted to other molecules or combined into polymers	
но он	Disaccharides: lactose, sucrose	combined into polymers	
H OH Monosaccharide monomer	Polysaccharides:  Cellulose (plants) Starch (plants) Glycogen (animals) Chitin (animals and fungi)	Strengthens plant cell walls     Stores glucose for energy     Stores glucose for energy     Strengthens exoskeletons and fungal cell walls	

Components	Examples	Functions
Glycerol 3 fatty acids	Triacylglycerols (fats or oils): glycerol + three fatty acids	Important energy source
Head with P 2 fatty acids	Phospholipids: glycerol + phosphate group + two fatty acids	Lipid bilayers of membranes Hydrophobio tails Hydrophilic heads
Steroid backbone	Steroids: four fused rings with attached chemical groups	Component of cell membranes (cholesterol) Signaling molecules that travel through the body (hormones)

