

Nucleic Acid, its properties, types and its synthesis

ABC-111

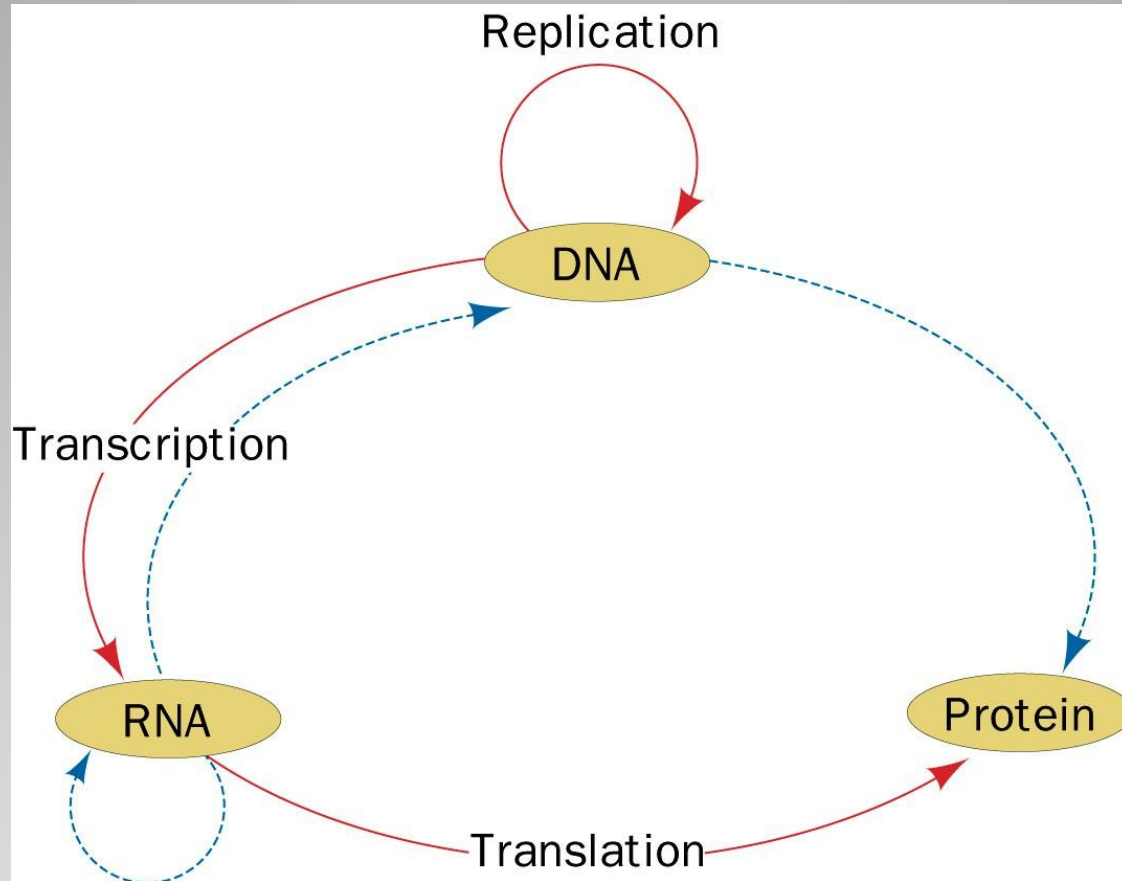
Dr. Mamta Rathore

Teaching Associates

Department of Agriculture

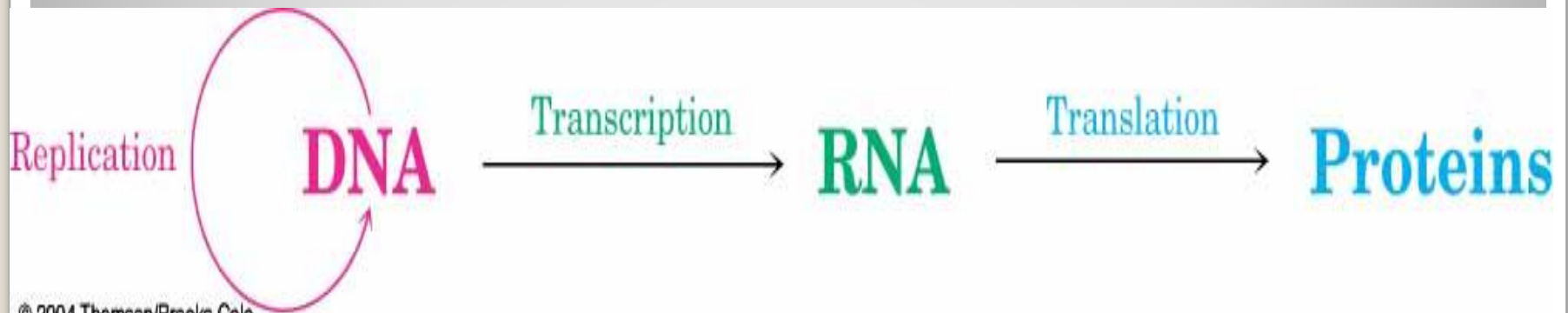
Biochemistry

C.S.A.U.A & T. Kanpur

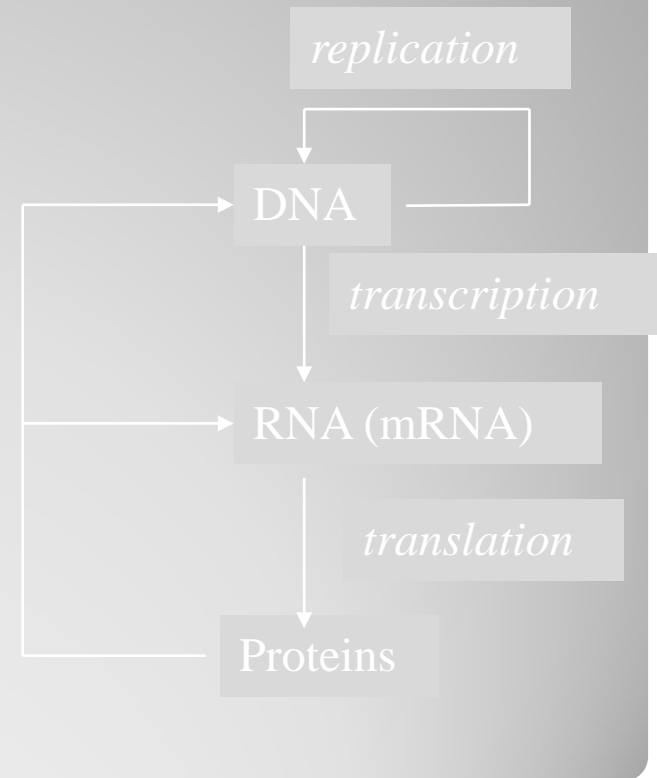
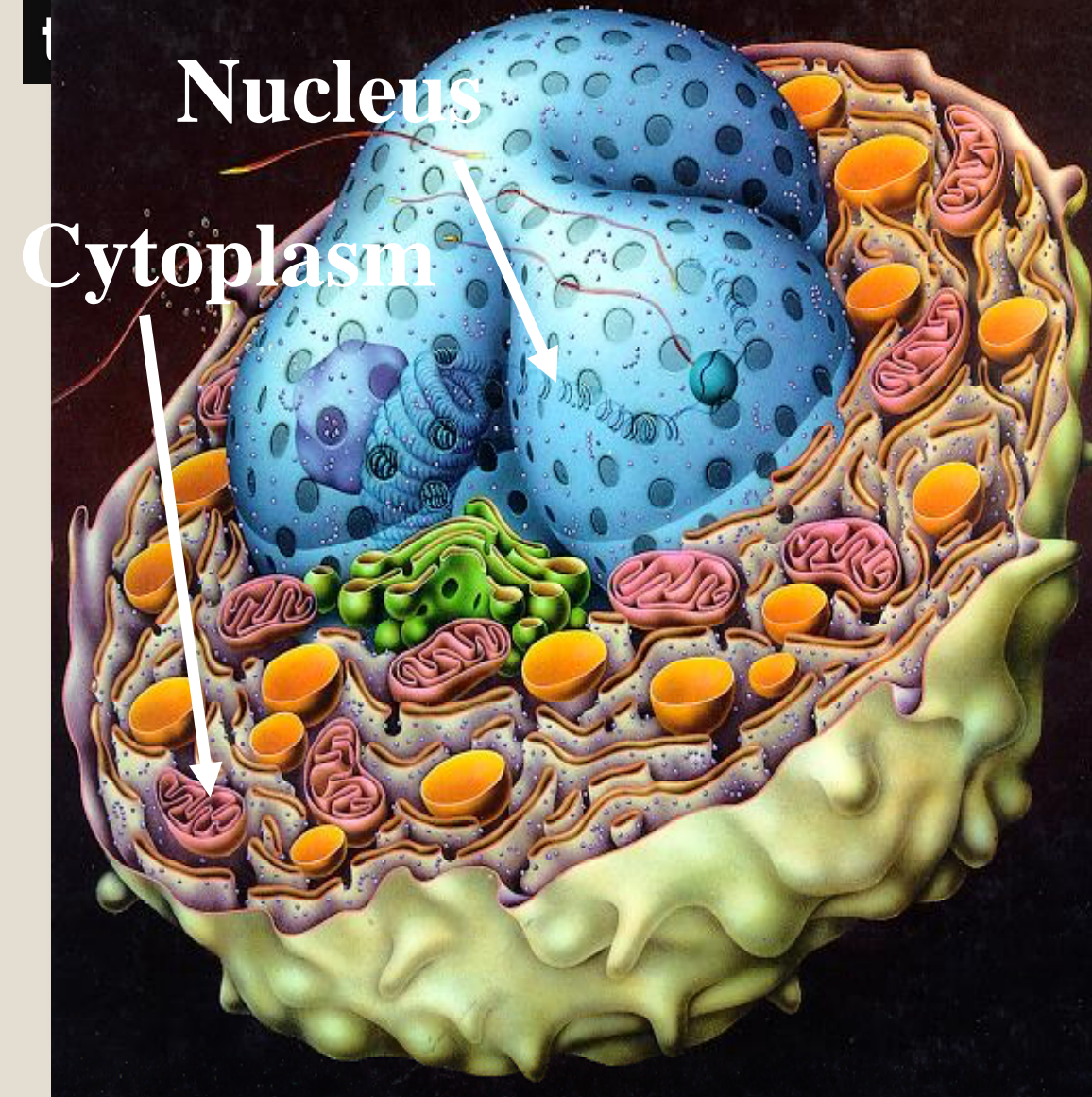


The central dogma of molecular biology.

- Processes in the transfer of genetic information:
- **Replication**: identical copies of DNA are made
- **Transcription**: genetic messages are read and carried out of the cell nucleus to the ribosomes, where protein synthesis occurs.
- **Translation**: genetic messages are decoded to make proteins.



**The nucleus contains the cell's DNA (genome)
RNA is synthesized in the nucleus and exported**



Two types of Nucleotides

(depending on the sugar they contain)

1- Ribonucleic acids (RNA)

The pentose sugar is **Ribose**
(has a hydroxyl group in the 3rd
carbon---**OH**)

2- Deoxyribonucleic acids (DNA)

The pentose sugar is
Deoxyribose (has just an
hydrogen in the same place--- **H**)
Deoxy = "minus oxygen"

Definitions

Nucleic acids are polymers of nucleotides

Nucleotides are carbon ring structures containing nitrogen linked to a 5-carbon sugar (a ribose)

5-carbon sugar is either a ribose or a deoxy-ribose making the nucleotide either a ribonucleotide or a deoxyribonucleotide

In eukaryotic cells nucleic acids are either:

Deoxyribose nucleic acids (DNA)

Ribose nucleic acids (RNA)

Messenger RNA (mRNA)

Transfer RNA (tRNA)

Ribosomal RNA (rRNA)

Nucleic Acid Function

DNA

Genetic material - sequence of nucleotides encodes different amino acids

RNA

Involved in the transcription/translation of genetic material (DNA)

Genetic material of some viruses

Nucleotide Structure

Despite the complexity and diversity of life the structure of DNA is dependent on only 4 different nucleotides

Diversity is dependent on the nucleotide sequence

All nucleotides are 2 ring structures composed of:

5-carbon sugar :	β -D-ribose (RNA) β -D-deoxyribose (DNA)
-------------------------	---------------------------------------------------------

Base	Purine Pyrimidine
-------------	------------------------------------

Phosphate group	A nucleotide WITHOUT a phosphate group is a <u>NUCLEOSIDE</u>
------------------------	-----------------------------------------------------------------------------

NUCLEIC ACIDS (DNA and RNA) Notes

DNA – Deoxyribonucleic Acid

- DNA controls all living processes including production of new cells – cell division
- DNA carries the genetic code – stores and transmits genetic information from one generation to the next
- Chromosomes are made of DNA
- DNA is located in the nucleus of the cell

Nucleotides and Nucleosides

- Nucleotide =
 - Nitrogenous base
 - Pentose
 - Phosphate
- Nucleoside =
 - Nitrogenous base
 - Pentose
- Nucleobase =
 - Nitrogenous base

- Simple units called **nucleotides**, connected in long chains
- **Nucleotides have 3 parts:**
 - 1- 5-Carbon sugar (pentose)
 - 2- Nitrogen containing base
(made of C, H and N)
 - 3- A phosphate group (P)
- The P groups make the links that unite the sugars (hence a "**sugar-phosphate backbone**")

What are they made of ?

TABLE 8–1**Nucleotide and Nucleic Acid Nomenclature**

Base	Nucleoside	Nucleotide	Nucleic acid
Purines			
Adenine	Adenosine	Adenylate	RNA
	Deoxyadenosine	Deoxyadenylate	DNA
Guanine	Guanosine	Guanylate	RNA
	Deoxyguanosine	Deoxyguanylate	DNA
Pyrimidines			
Cytosine	Cytidine	Cytidylate	RNA
	Deoxycytidine	Deoxycytidylate	DNA
Thymine	Thymidine or deoxythymidine	Thymidylate or deoxythymidylate	DNA
Uracil	Uridine	Uridylate	RNA

Note: “Nucleoside” and “nucleotide” are generic terms that include both ribo- and deoxyribo- forms. Also, ribonucleosides and ribonucleotides are here designated simply as nucleosides and nucleotides (e.g., riboadenosine as adenosine), and deoxyribonucleosides and deoxyribonucleotides as deoxynucleosides and deoxynucleotides (e.g., deoxyriboadenosine as deoxyadenosine). Both forms of naming are acceptable, but the shortened names are more commonly used. Thymine is an exception; “ribothymidine” is used to describe its unusual occurrence in RNA.

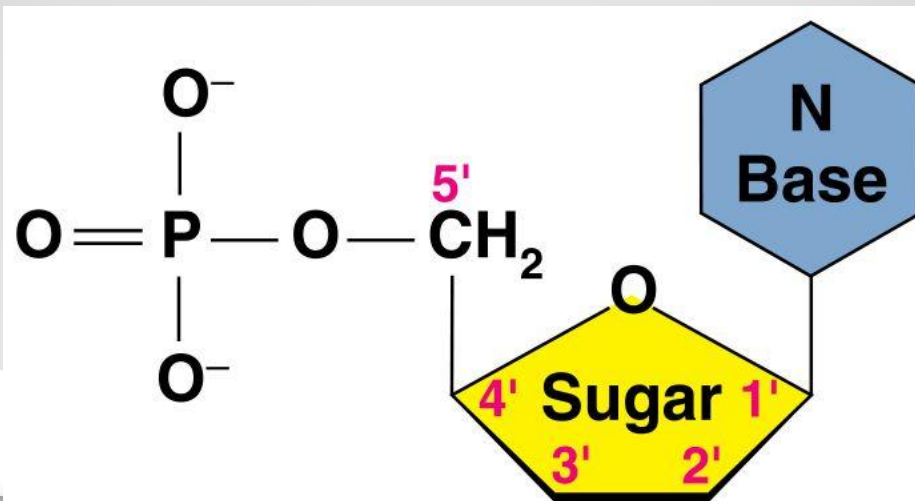
Table 8-1

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Nucleic Acids

- **Nucleic acids** are molecules that store information for cellular growth and reproduction
- There are two types of nucleic acids:
 - **deoxyribonucleic acid (DNA)** and **ribonucleic acid (RNA)**
- These are polymers consisting of long chains of monomers called nucleotides
- A **nucleotide** consists of a nitrogenous base, a pentose sugar and a phosphate group:

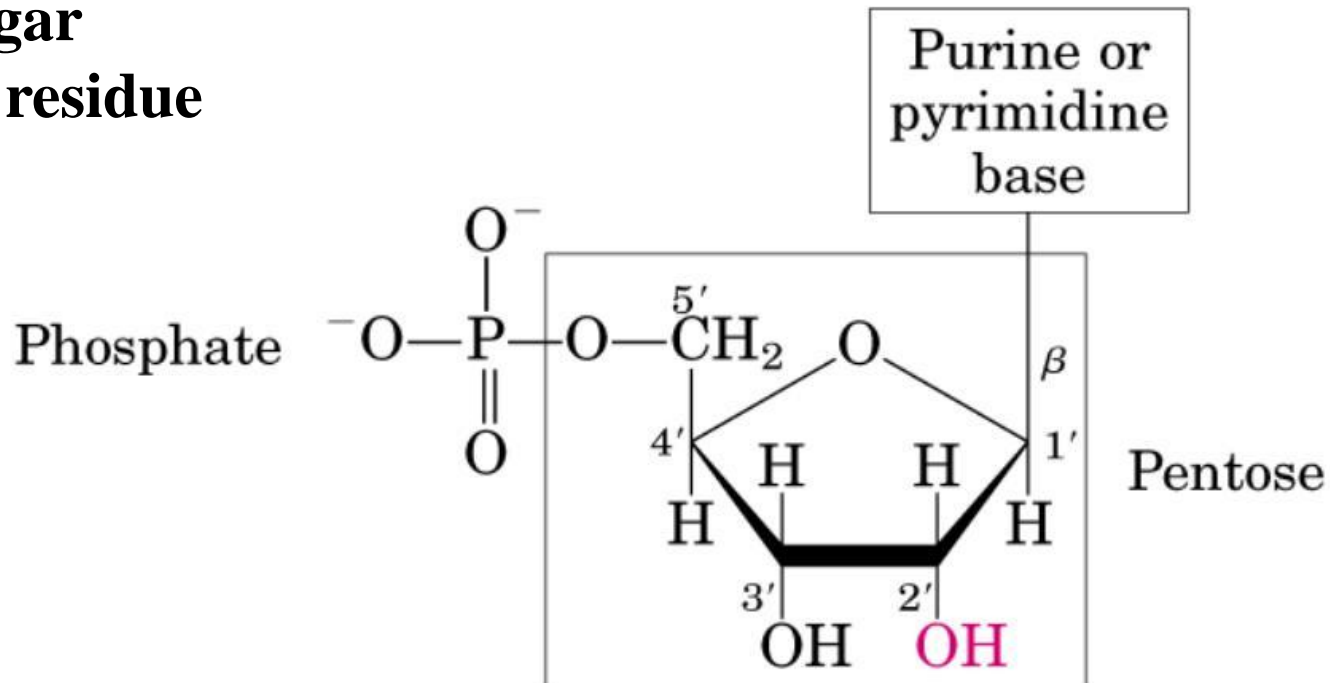


Nucleic Acids

DNA and RNA are nucleic acids, long, thread-like polymers made up of a linear array of monomers called nucleotides

All nucleotides contain three components:

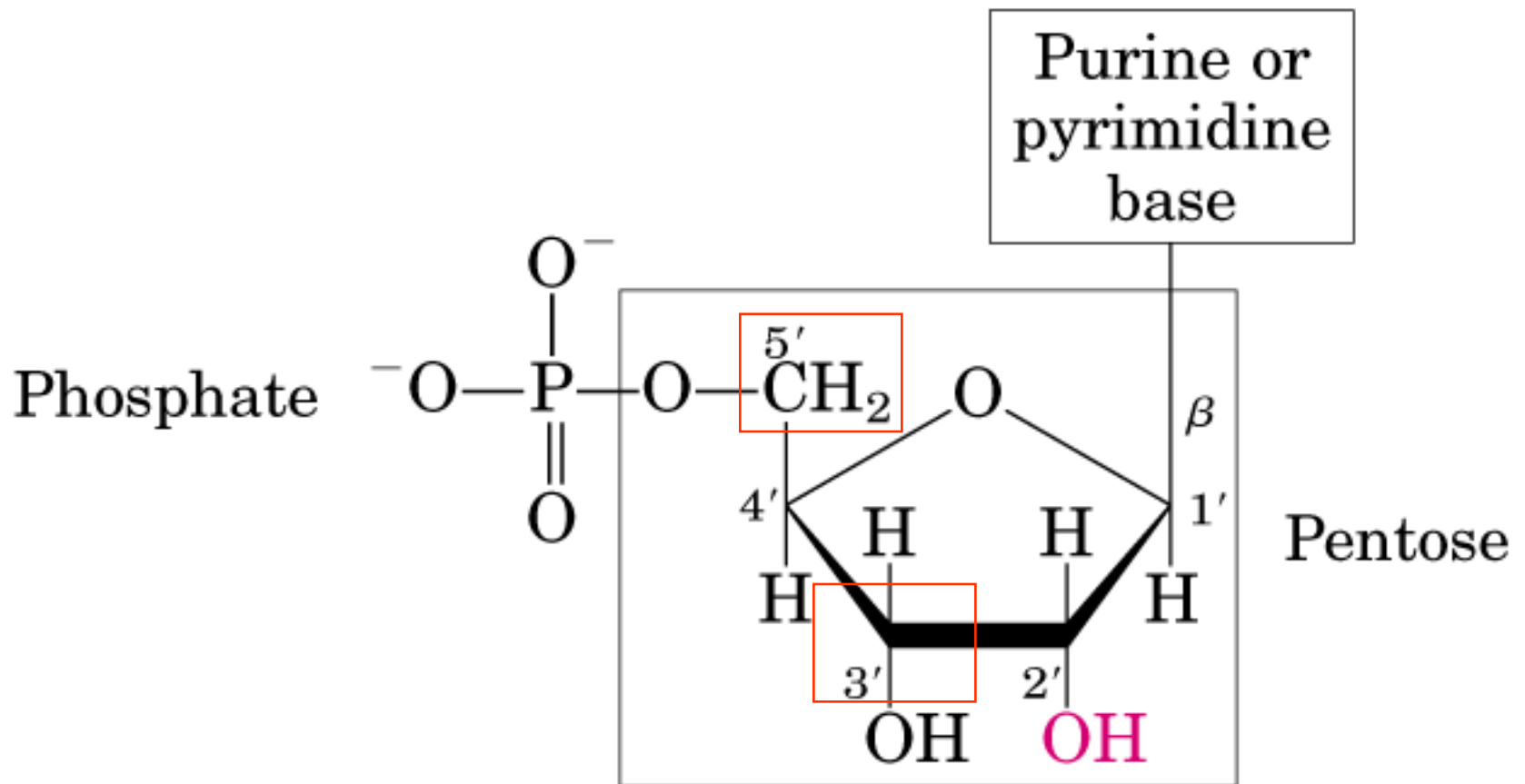
1. A nitrogen heterocyclic base
2. A pentose sugar
3. A phosphate residue



Chemical Structure of DNA vs RNA

Ribonucleotides have a 2'-OH

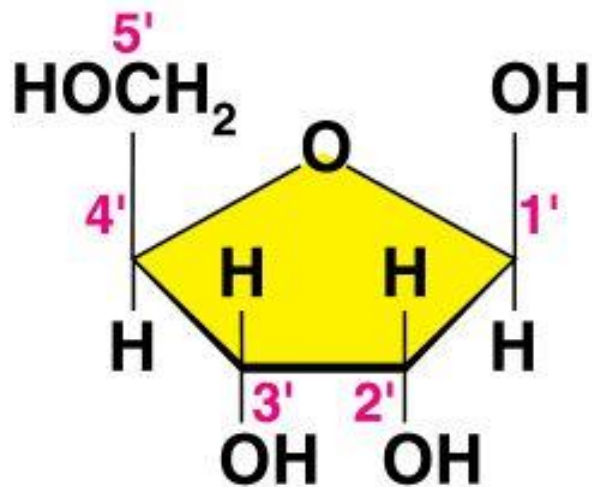
Deoxyribonucleotides have a 2'-H



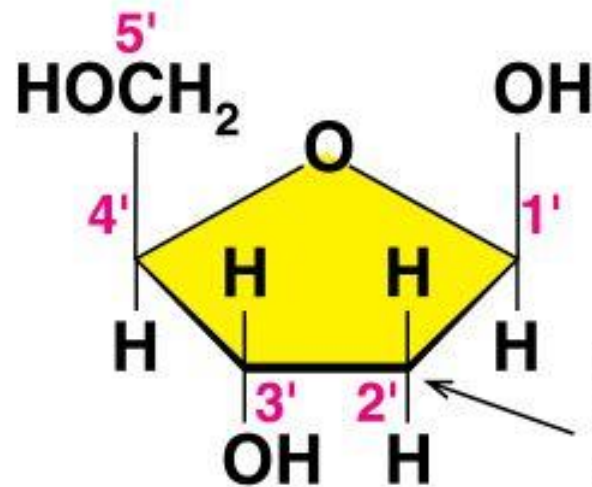
Pentose Sugars

- There are two related **pentose sugars**:
 - RNA contains **ribose**
 - DNA contains **deoxyribose**
- The sugars have their carbon atoms numbered with primes to distinguish them from the nitrogen bases

Pentose sugars in RNA and DNA




Ribose in RNA



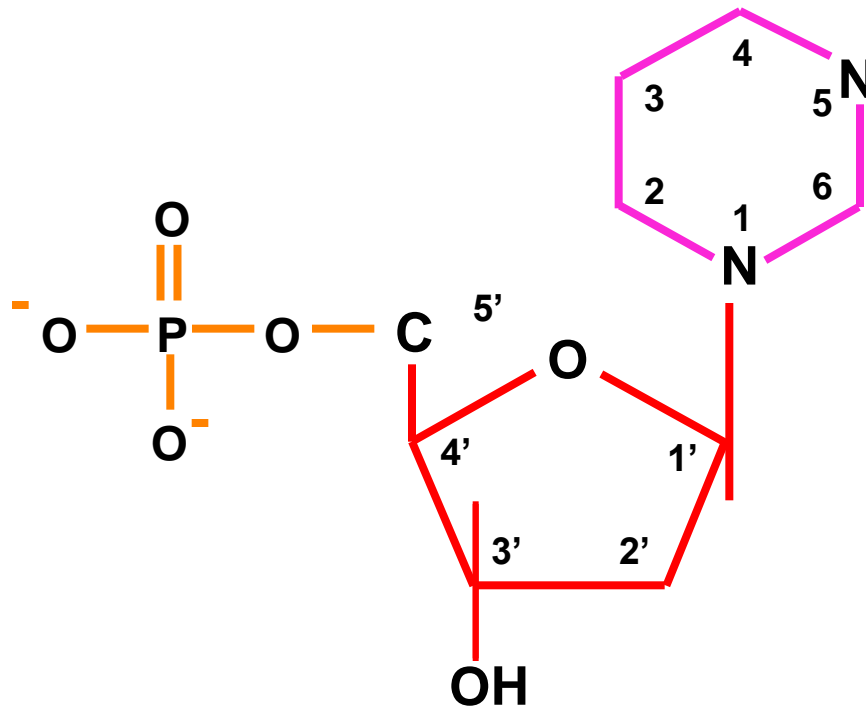
Deoxyribose in DNA

No oxygen
is bonded
to this carbon



Nucleotide Structure - 4

Base-Sugar-PO₄²⁻



Monophosphate

Nucleotide Function

Building blocks for DNA and RNA

Intracellular source of energy - Adenosine triphosphate (ATP)

Second messengers - Involved in intracellular signaling (e.g. cyclic adenosine monophosphate [cAMP])

Intracellular signaling switches (e.g. G-proteins)

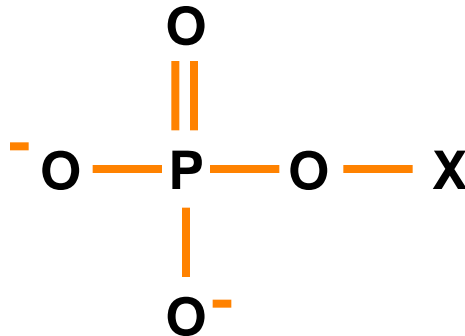
Nucleotide Structure - 4

Phosphate Groups

Phosphate groups are what makes a nucleoside a nucleotide

Phosphate groups are essential for nucleotide polymerization

Basic structure:



Nucleotide Structure - 4

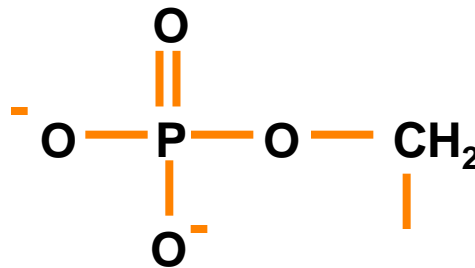
Phosphate Groups

Number of phosphate groups determines nomenclature

Monophosphate

e.g. AMP

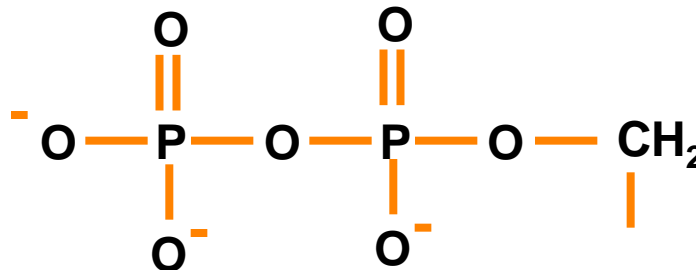
Free = inorganic
phosphate (Pi)



Diphosphate

e.g. ADP

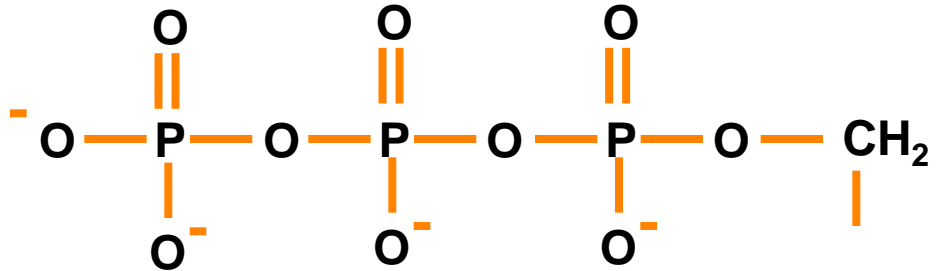
Free = Pyro-
phosphate (PPi)



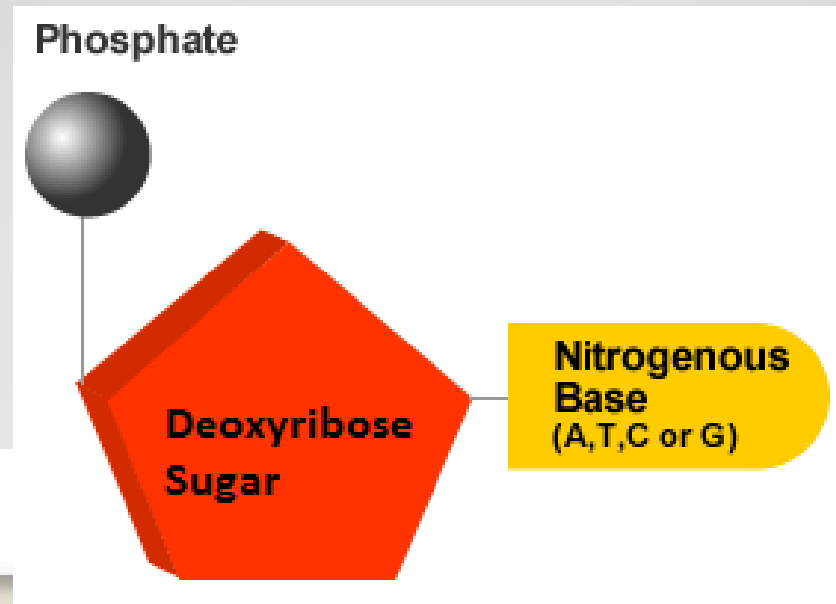
Nucleotide Structure - 4 Phosphate Groups

Triphosphate
e.g. ATP

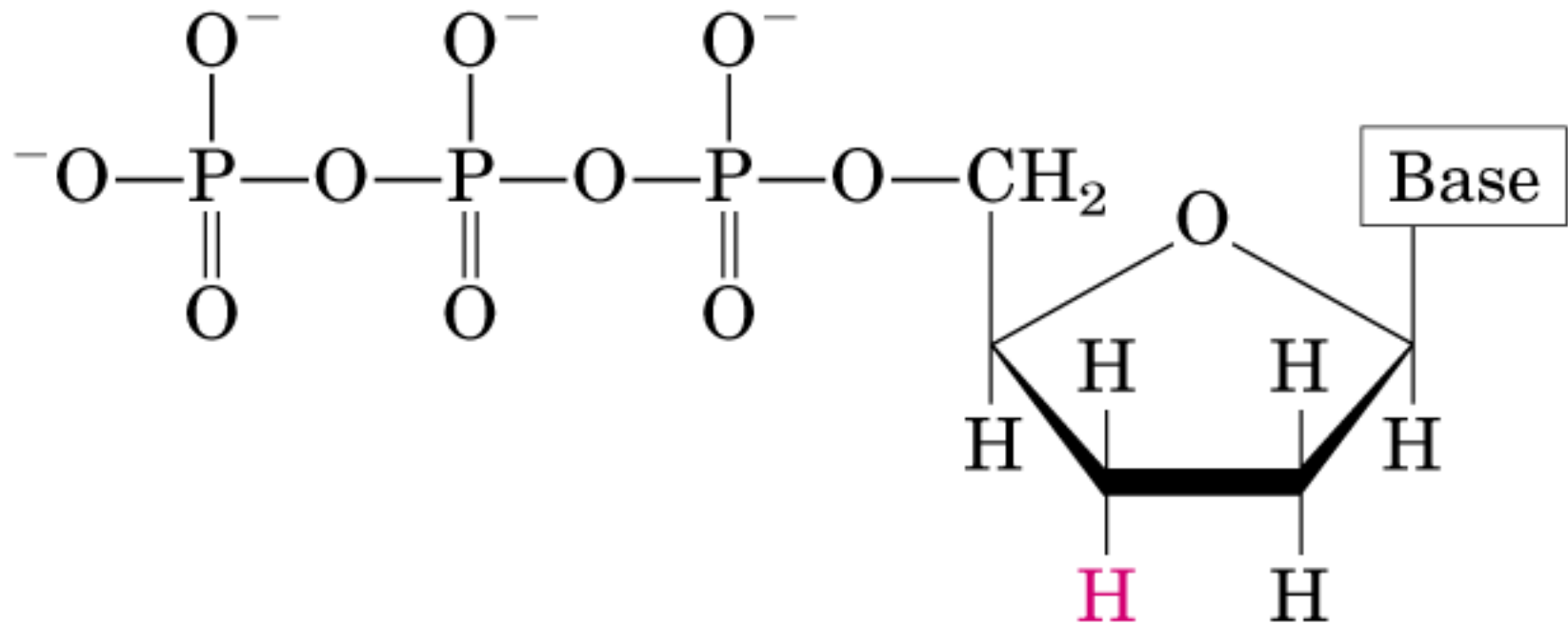
No Free form exists



- It is the order of these base pairs that determines genetic makeup
- One phosphate + one sugar + one base = one nucleotide
- Nucleotides are the building blocks of DNA – thus, each strand of DNA is a string of nucleotides



Sanger dideoxy sequencing incorporates dideoxy nucleotides, preventing further synthesis of the DNA strand



ddNTP analog

base (purine, pyrimidine) + ribose
(deoxyribos



N-glycosyl

linkage

nucleoside+phosphate



phosphoester

linkage



nucleotide

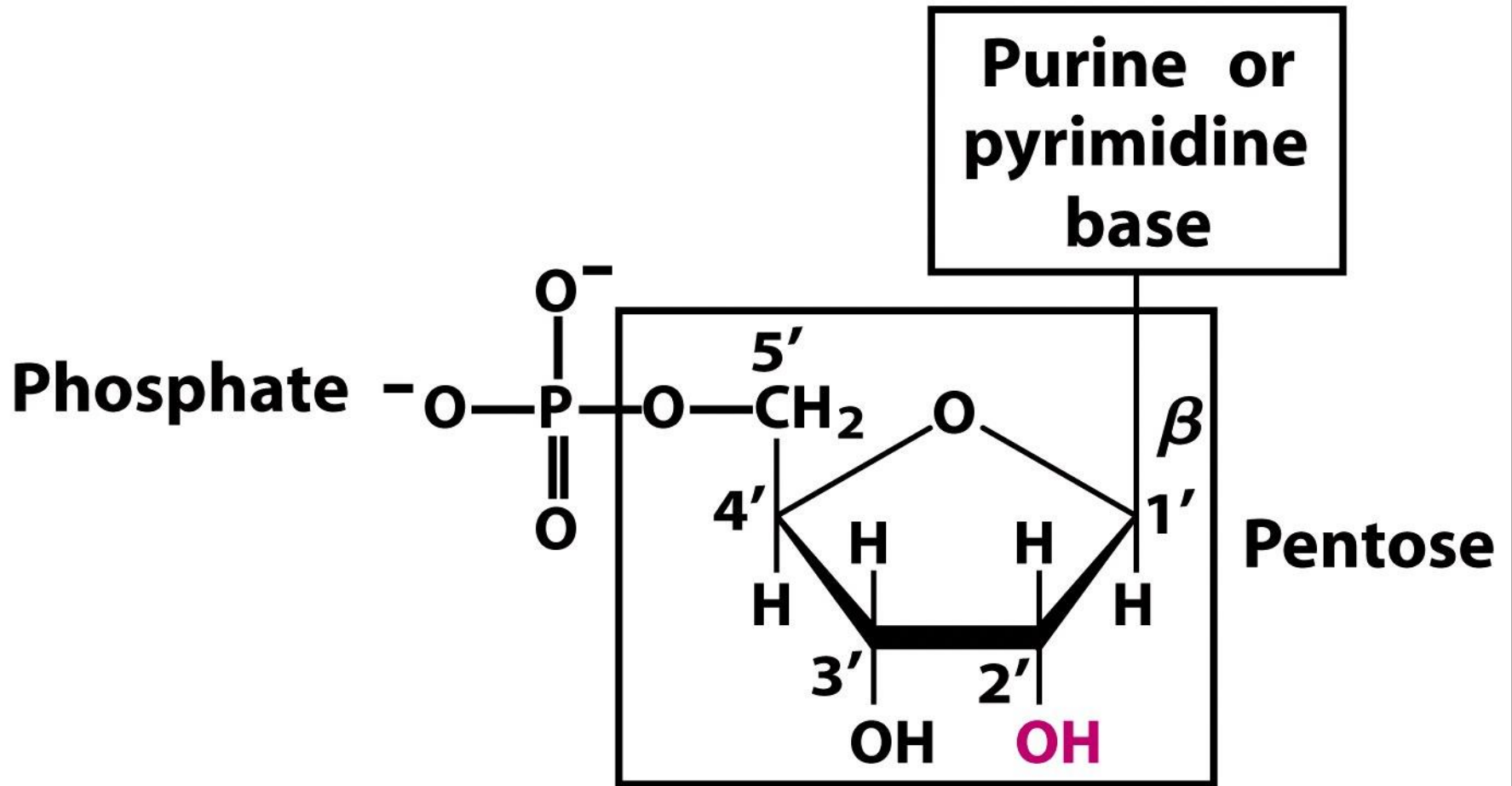
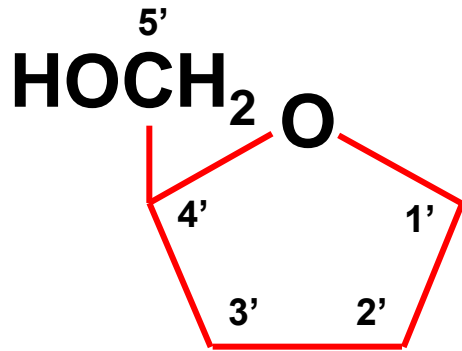


Figure 8-1a
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Nucleotide Structure - 1

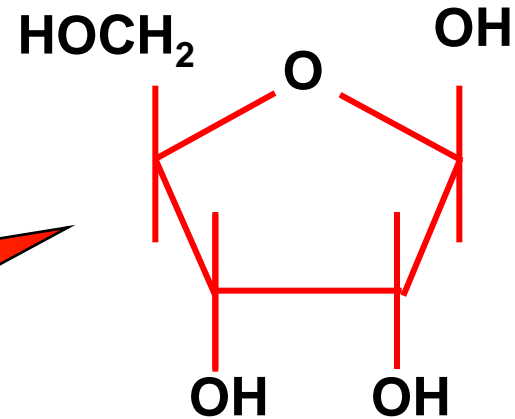
Sugars

Generic Ribose
Structure

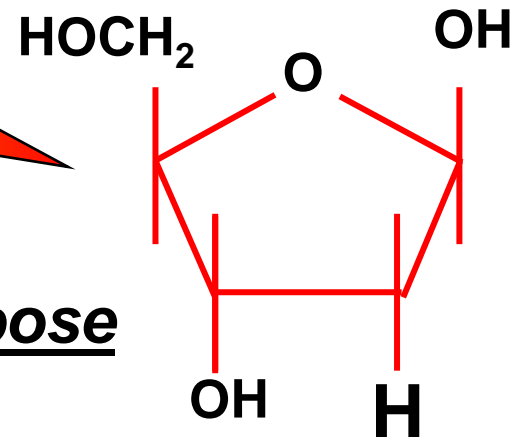


N.B. Carbons are given numberings
as a prime

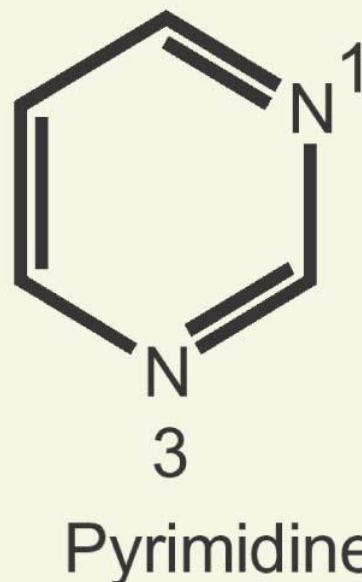
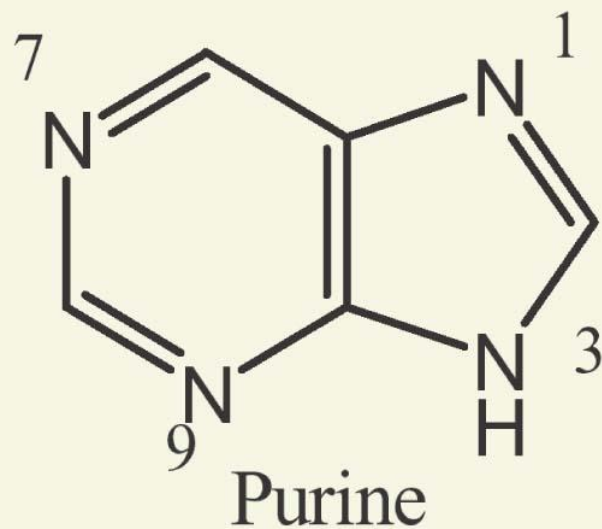
Ribose



Deoxyribose



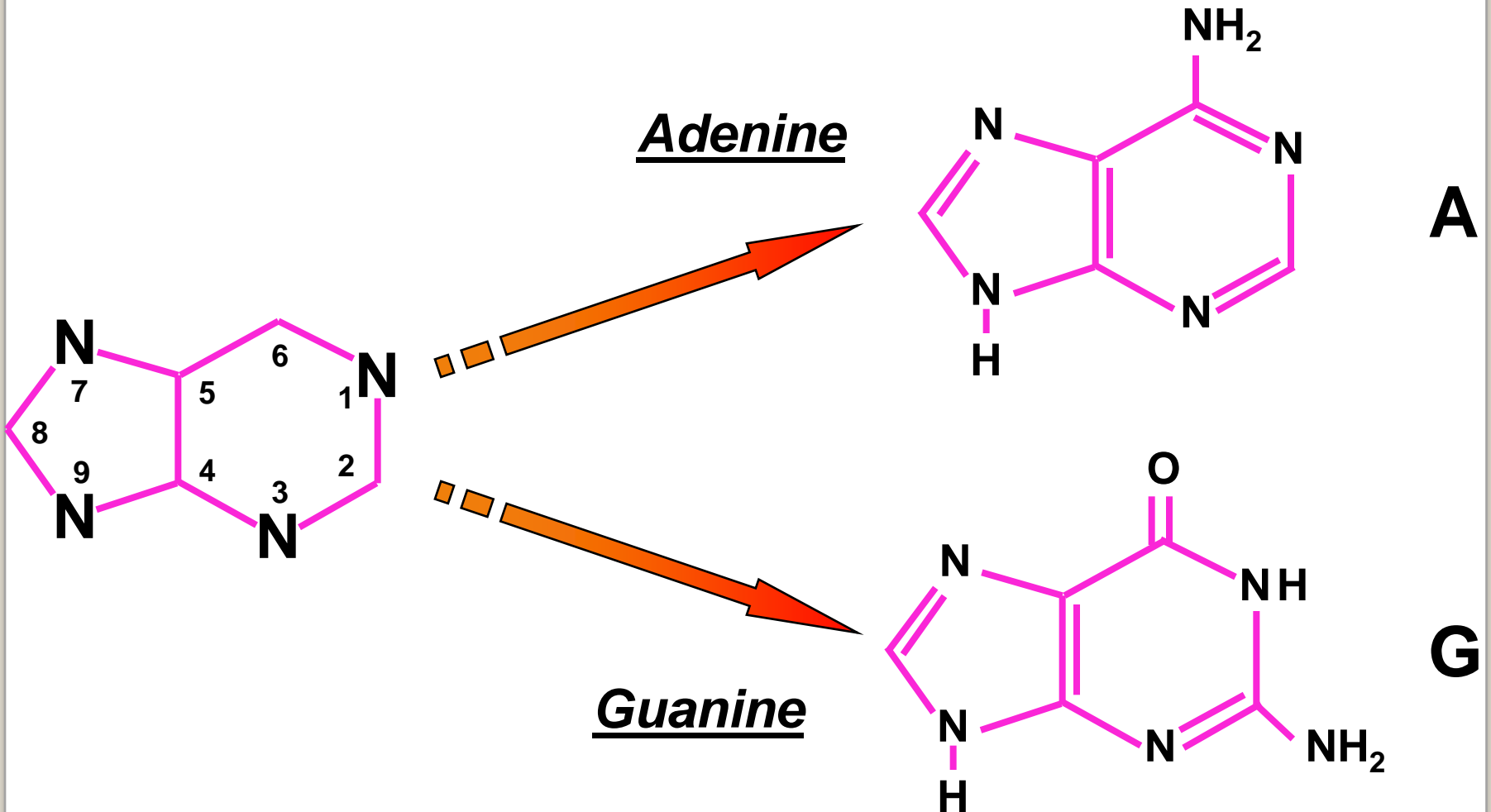
- Pyrimidine contains two pyridine-like nitrogens in a six-membered aromatic ring
- Purine has 4 N's in a fused-ring structure. Three are basic like pyridine-like and one is like that in pyrrole.

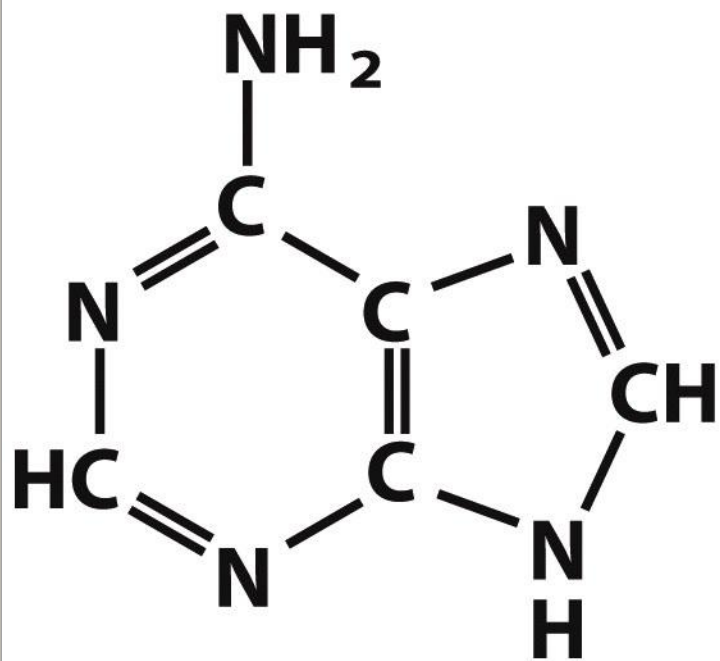


Puri

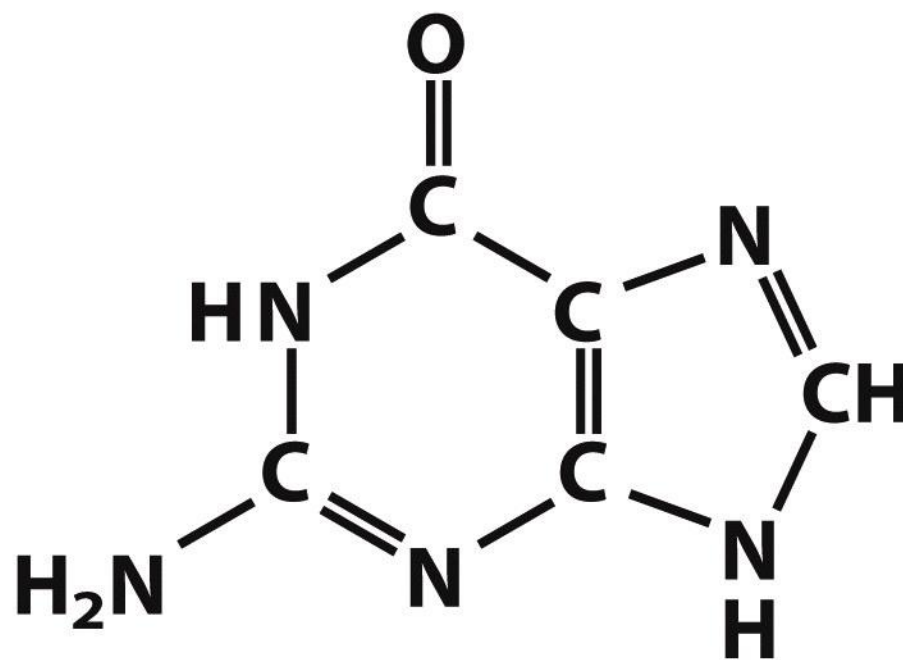
Nucleotide Structure - 2

Bases - Purines





Adenine



Guanine

Purines

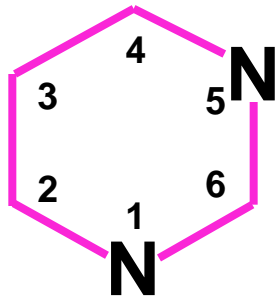
Figure 8-2 part 1

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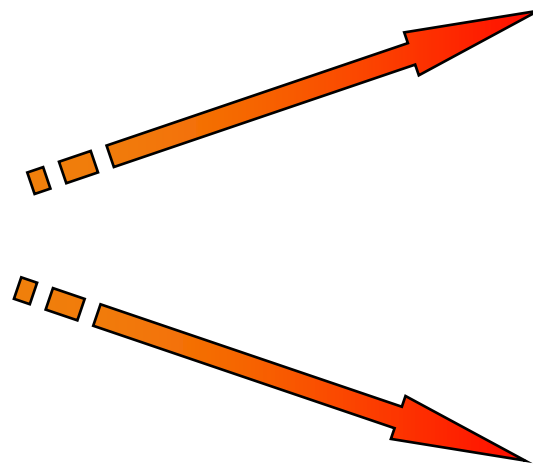
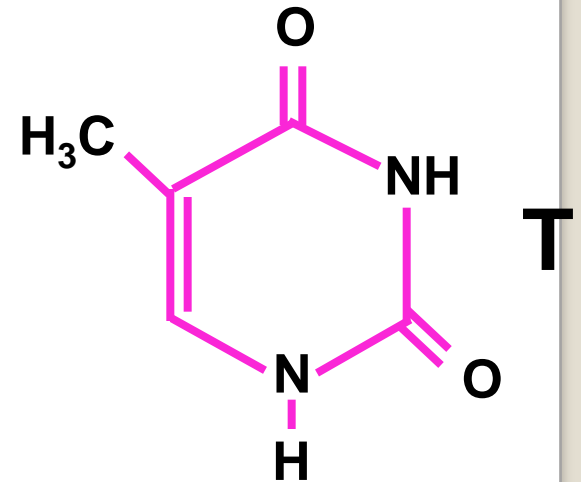
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Nucleotide Structure - 3

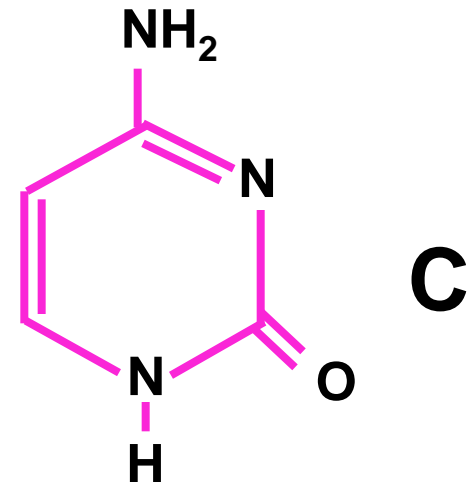
Bases - Pyrimidines

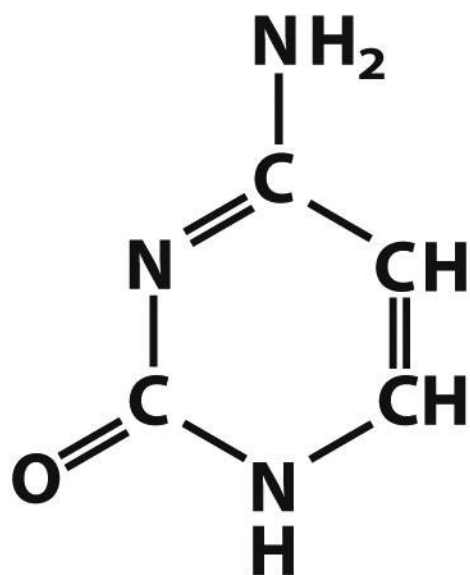


Thymine

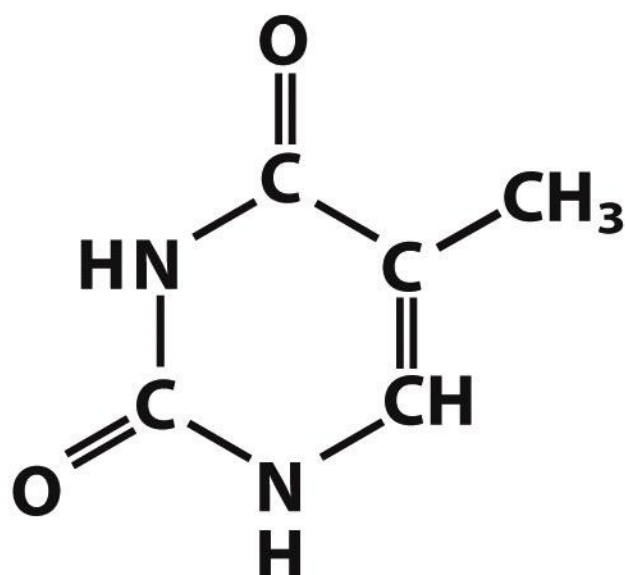


Cytosine

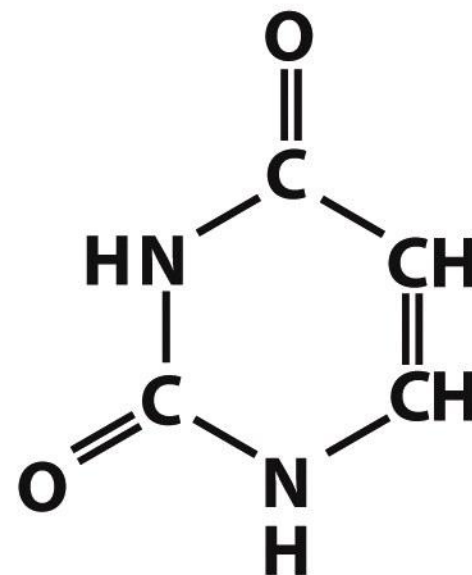




Cytosine



**Thymine
(DNA)**



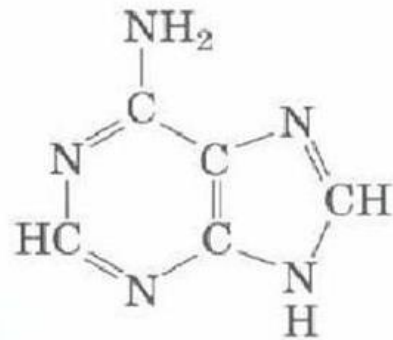
**Uracil
(RNA)**

Pyrimidines

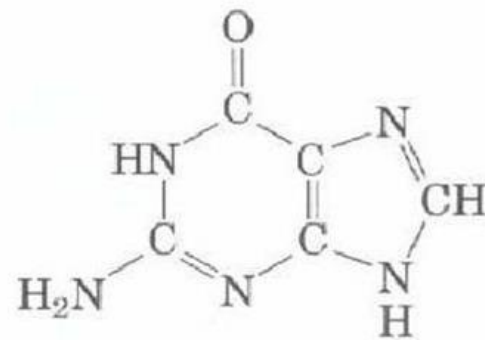
Figure 8-2 part 2

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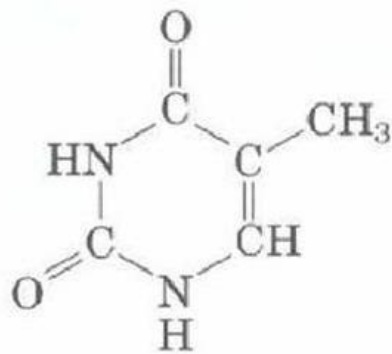


Adenine

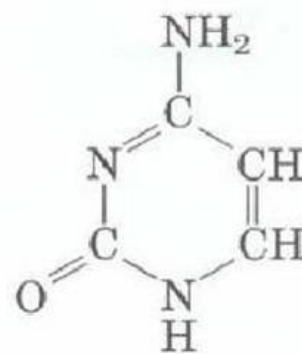


Guanine

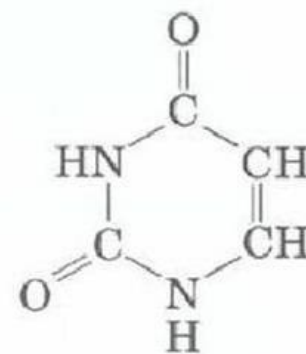
Purines



Thymine
(DNA)



Cytosine



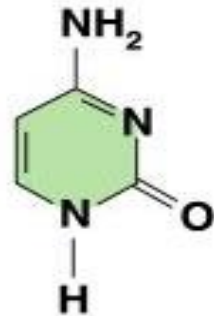
Uracil
(RNA)

Pyrimidines

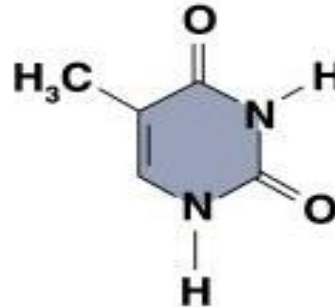
Nitrogen Bases

- The **nitrogen bases** in nucleotides consist of two general types:
 - **purines**: adenine (A) and guanine (G)
 - **pyrimidines**: cytosine (C), thymine (T) and Uracil (U)

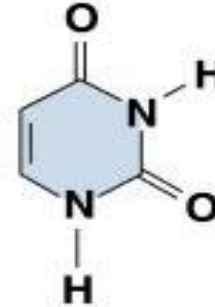
Pyrimidines



Cytosine (C)
(DNA and RNA)



Thymine (T)
(DNA only)

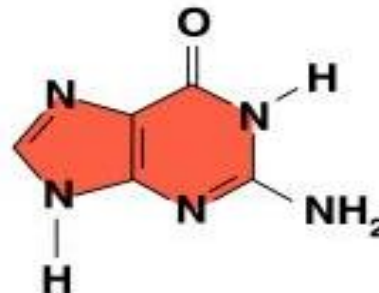


Uracil (U)
(RNA only)

Purines



Adenine (A)
(DNA and RNA)



Guanine (G)
(DNA and RNA)

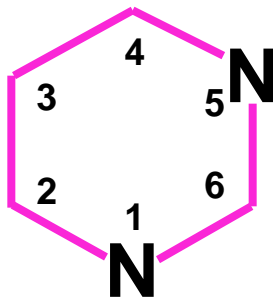
Nucleotide Structure - 4

Bases - Pyrimidines

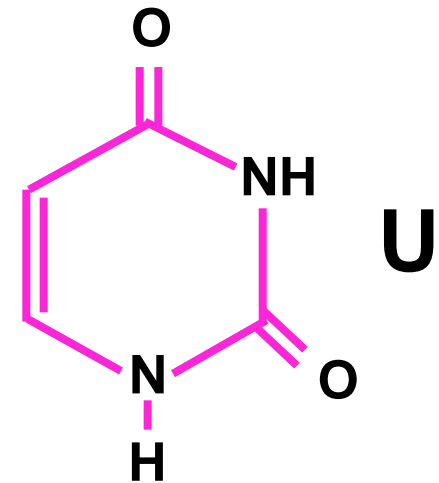
Thymine is found ONLY in DNA.

In RNA, thymine is replaced by uracil

Uracil and Thymine are structurally similar



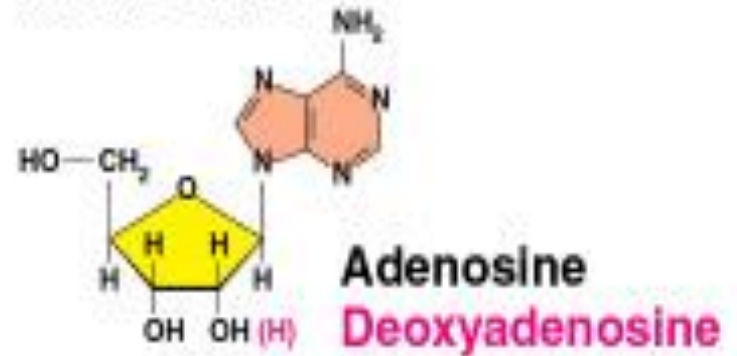
Uracil



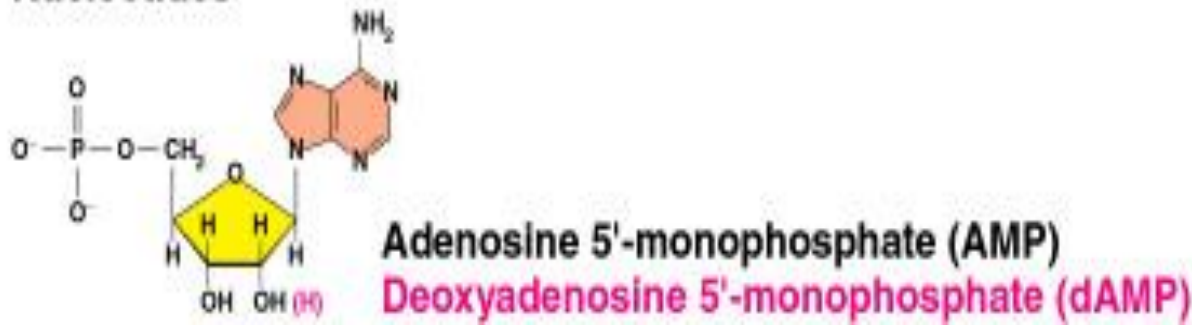
Nucleosides and Nucleotides

- A nucleoside consists of a nitrogen base linked by a glycosidic bond to C1' of a ribose or deoxyribose
- Nucleosides are named by changing the the nitrogen base ending to *-osine* for purines and *-idine* for pyrimidines
- A nucleotide is a nucleoside that forms a phosphate ester with the C5' OH group of ribose or deoxyribose
- Nucleotides are named using the name of the nucleoside followed by *5'-mono*

Nucleosides



Nucleotides

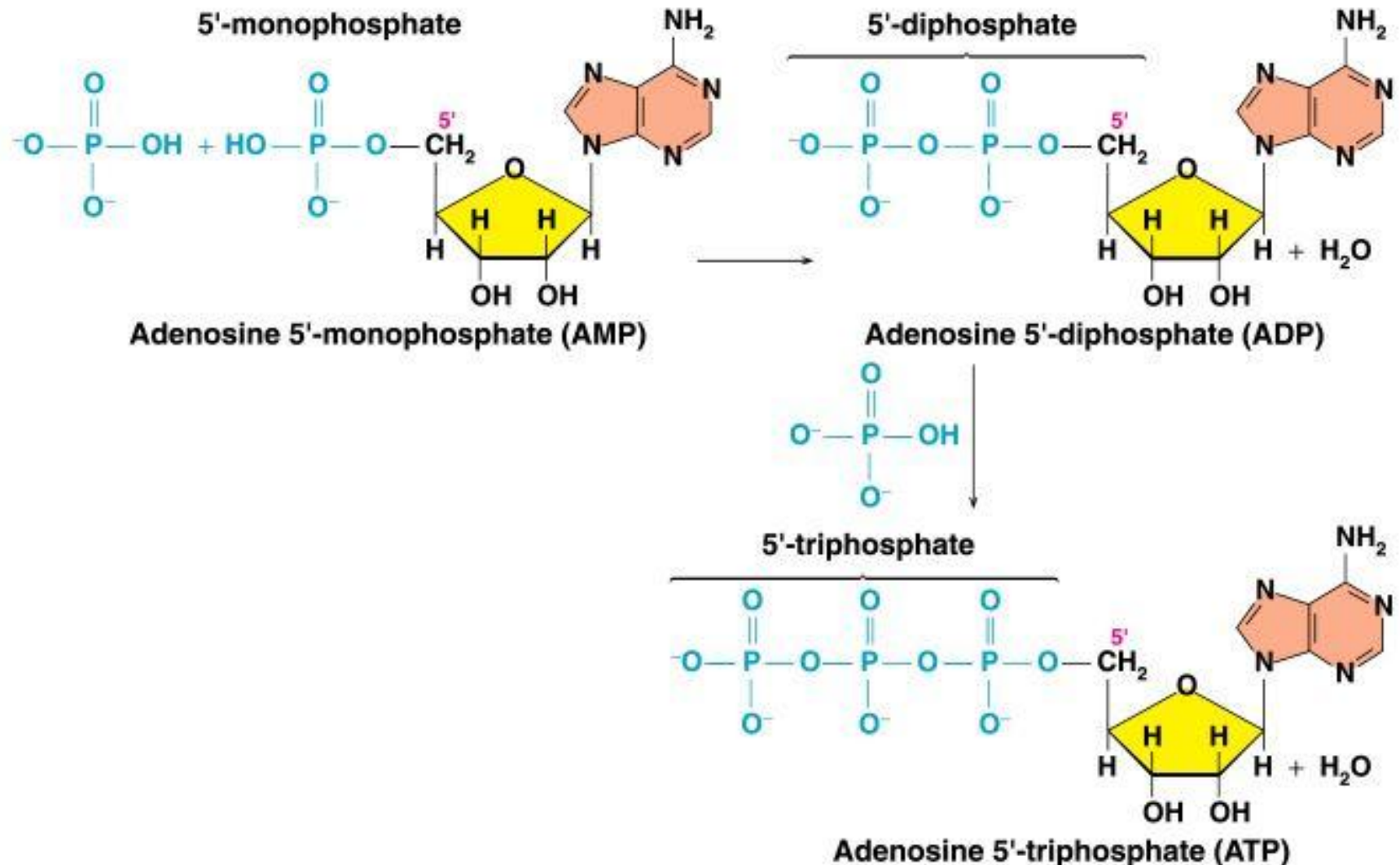


Names of Nucleosides and Nucleotides

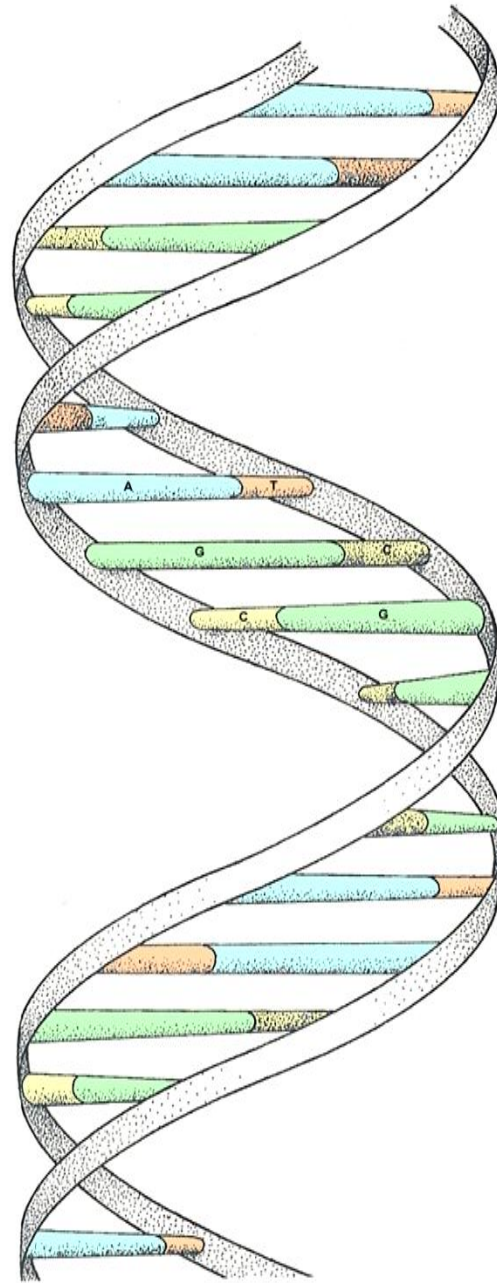
Base	Nucleosides	Nucleotides
RNA		
Adenine (A)	Adenosine (A)	Adenosine 5'-monophosphate (AMP)
Guanine (G)	Guanosine (G)	Guanosine 5'-monophosphate (GMP)
Cytosine (C)	Cytidine (C)	Cytidine 5'-monophosphate (CMP)
Uracil (U)	Uridine (U)	Uridine 5'-monophosphate (UMP)
DNA		
Adenine (A)	Deoxyadenosine (A)	Deoxyadenosine 5'-monophosphate (dAMP)
Guanine (G)	Deoxyguanosine (G)	Deoxyguanosine 5'-monophosphate (dGMP)
Cytosine (C)	Deoxycytidine (C)	Deoxycytidine 5'-monophosphate (dCMP)
Thymine (T)	Deoxythymidine (T)	Deoxythymidine 5'-monophosphate (dTMP)

AMP, ADP and ATP

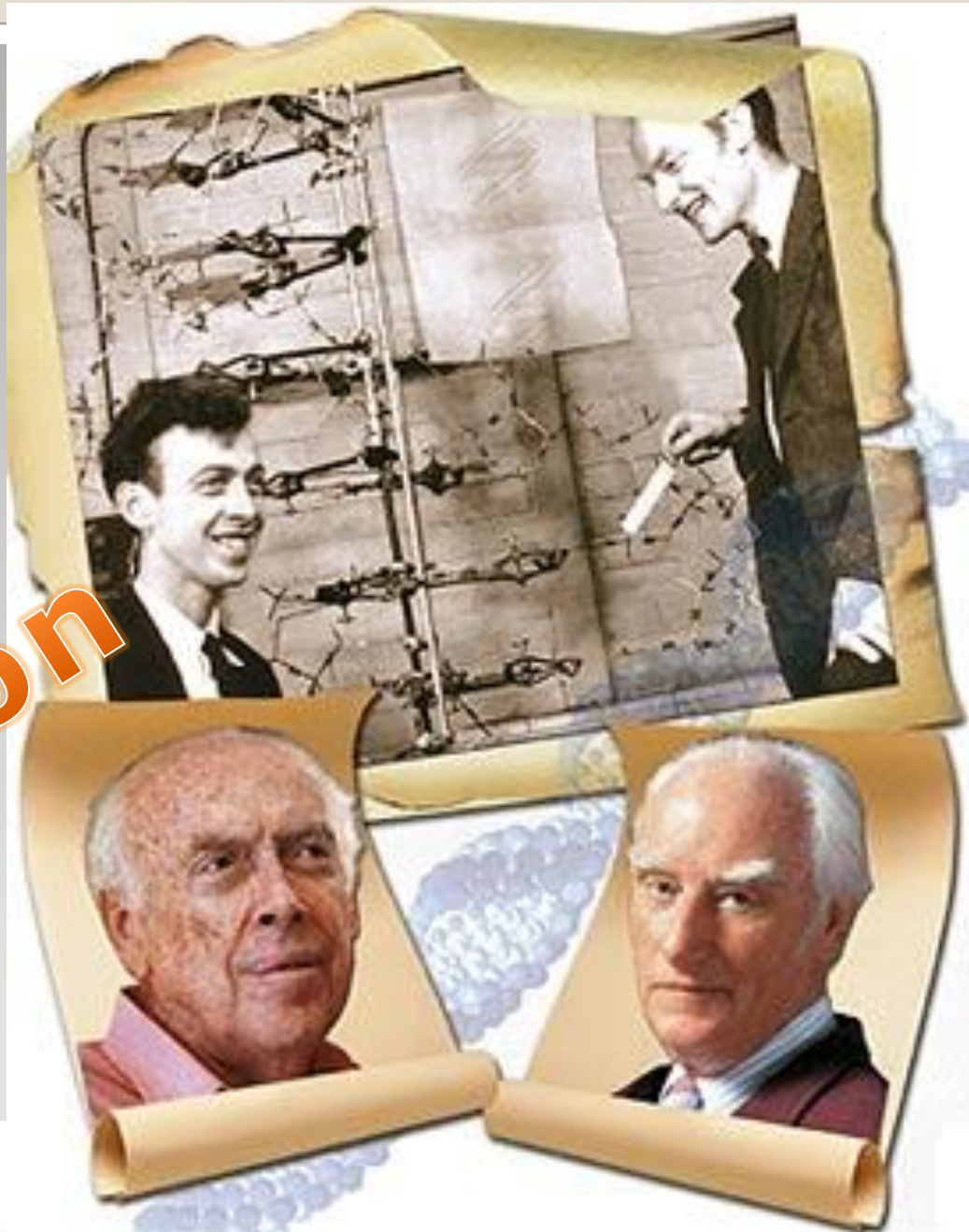
- Additional phosphate groups can be added to the nucleoside 5'-monophosphates to form **diphosphates** and **triphosphates**
- **ATP** is the major energy source for cellular activity



DNA



Watson



Crick

Died in
2004

DNA stands for **deoxyribose nucleic acid**

This chemical substance is present in the nucleus of all cells in all living organisms

DNA controls all the chemical changes which take place in cells

The kind of cell which is formed, (muscle, nerve etc) is controlled by DNA

DNA is a very large molecule made up of chain of sub-units

The sub-units are called **nucleotides**

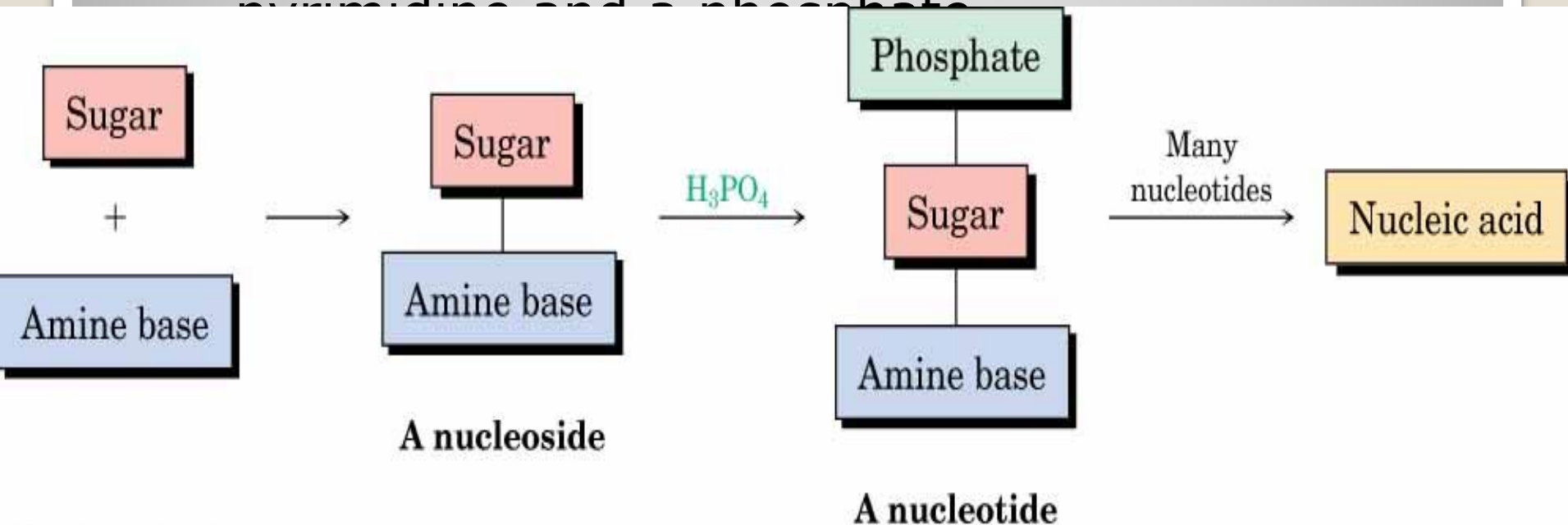
Each nucleotide is made up of

a sugar called **deoxyribose**

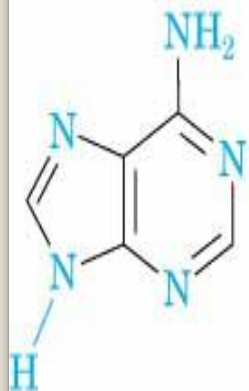
a phosphate group **-PO₄** and

an **organic base**

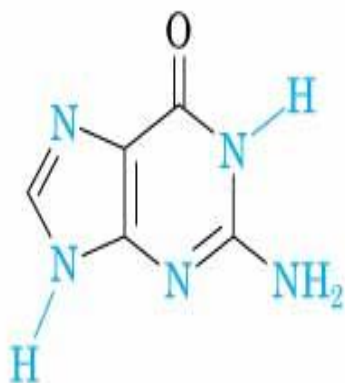
- Deoxyribonucleic acid (DNA) and ribonucleic acid (RNA), are the chemical carriers of genetic information
- Nucleic acids are biopolymers made of nucleotides, aldopentoses linked to a purine or pyrimidine and a phosphate



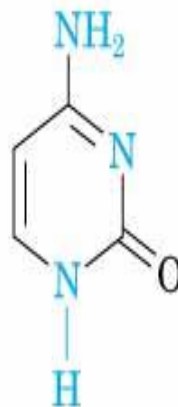
- Adenine, guanine, cytosine and thymine are in DNA



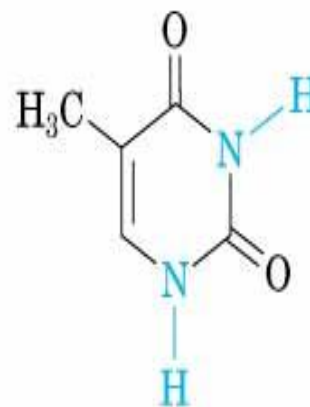
Adenine (A)
DNA
RNA



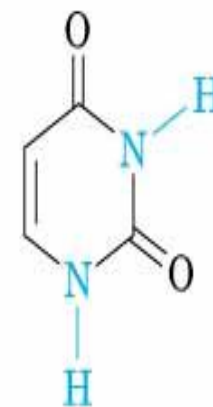
Guanine (G)
DNA
RNA



Cytosine (C)
DNA
RNA



Thymine (T)
DNA



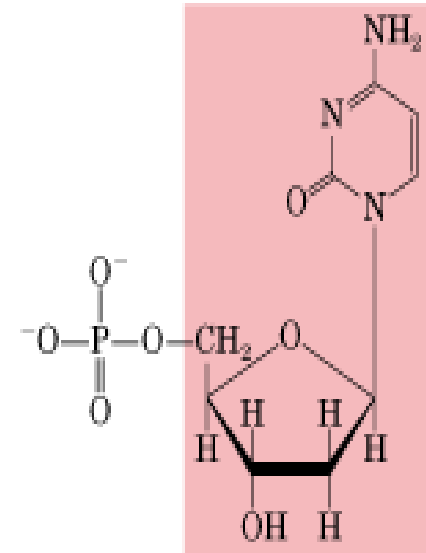
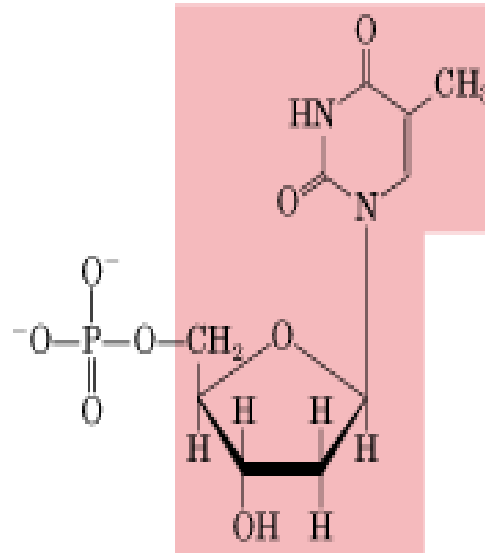
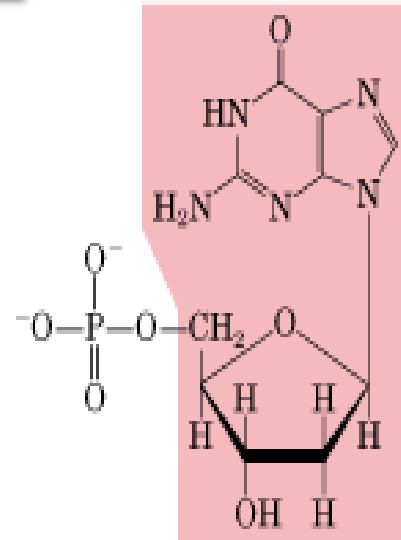
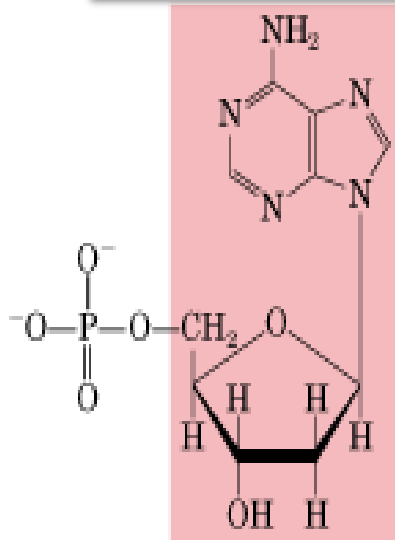
Uracil (U)
RNA

© 2004 Thomson/Brooks Cole

Heterocycles in DNA and RNA

Based on McMurry, Organic Chemistry, Chapter 28, 6th edition, (c) 2003

Deoxyribonucleotides found in DNA



Nucleotide: Deoxyadenylate
(deoxyadenosine
5'-monophosphate)

Nucleotide: Deoxyguanylate
(deoxyguanosine
5'-monophosphate)

Nucleotide: Deoxythymidylate
(deoxythymidine
5'-monophosphate)

Nucleotide: Deoxycytidylate
(deoxycytidine
5'-monophosphate)

Symbols: A, dA, dAMP

G, dG, dGMP

T, dT, dTMP

C, dC, dCMP

Nucleoside: Deoxyadenosine

Deoxyguanosine

Deoxythymidine

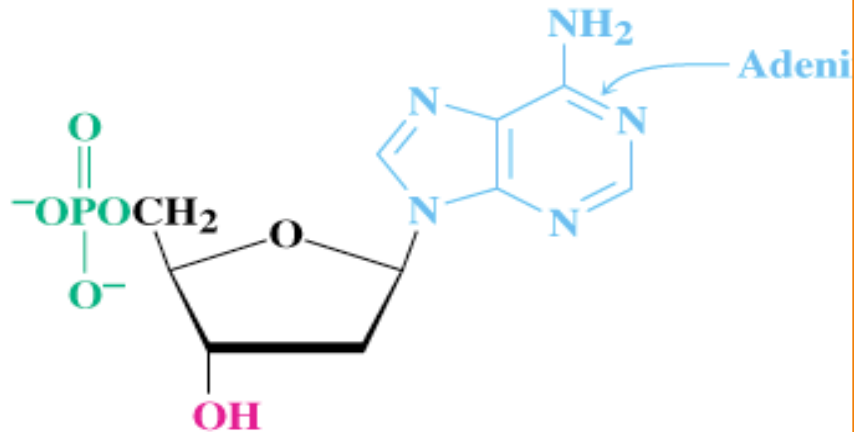
Deoxycytidine

dA

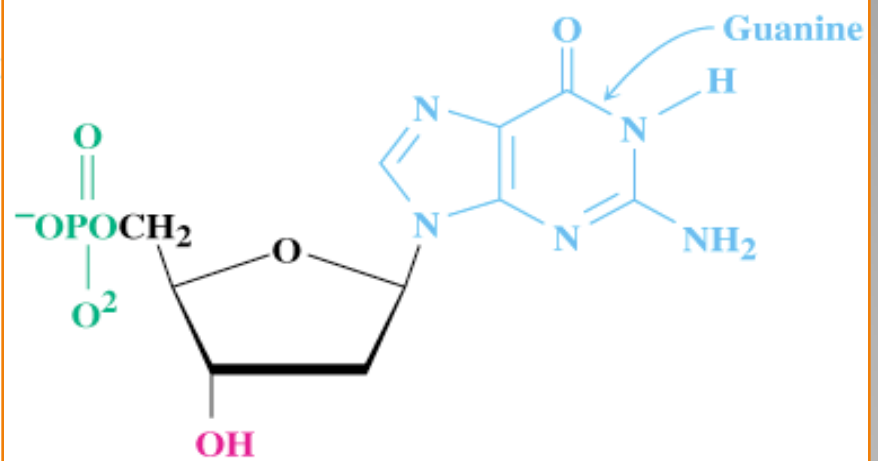
dG

dT

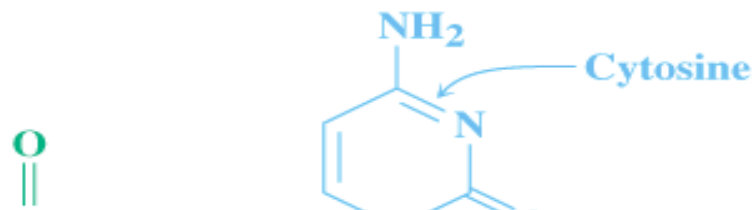
dC



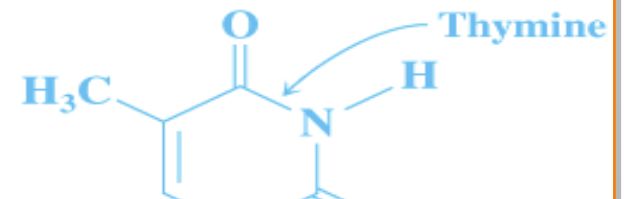
2'-Deoxyadenosine 5'-phosphate



2'-Deoxyguanosine 5'-phosphate



2'-Deoxycytidine 5'-phosphate



2'-Deoxythymidine 5'-phosphate

The Deoxyribonucleotides

Hydrogen Bonding Interactions

- Two bases can hydrogen bond to form a base pair
- For monomers, large number of base pairs is possible
- In polynucleotide, only few possibilities exist
- Watson-Crick base pairs predominate in double-stranded DNA
- A pairs with T
- C pairs with G
- Purine pairs with pyrimidine

– the building block molecule
of nucleic acid--**nucleotide**

In RNA:

AMP、CMP、GMP、TMP

In DNA:

dAMP、dCMP、dGMP、dUMP

Functions of Nucleotides and Nucleic Acids

- Nucleotide Functions:
 - Energy for metabolism (ATP)
 - Enzyme cofactors (NAD⁺)
 - Signal transduction (cAMP)
- Nucleic Acid Functions:
 - Storage of genetic info (DNA)
 - Transmission of genetic info (mRNA)
 - Processing of genetic information (ribozymes)
 - Protein synthesis (tRNA and rRNA)

二、the linkage ----

phosphodiester

bridge

3' terminal

5'terminal

Nucleotide

residues

DNA Nucleotides

Composition (3 parts):

- 1 - Deoxyribose sugar (no O in 3rd carbon)
- 2 - Phosphate group
- 3 - One of 4 types of bases (all containing nitrogen):
 - Adenine
 - Thymine (Only in DNA)
 - Cytosine
 - Guanine

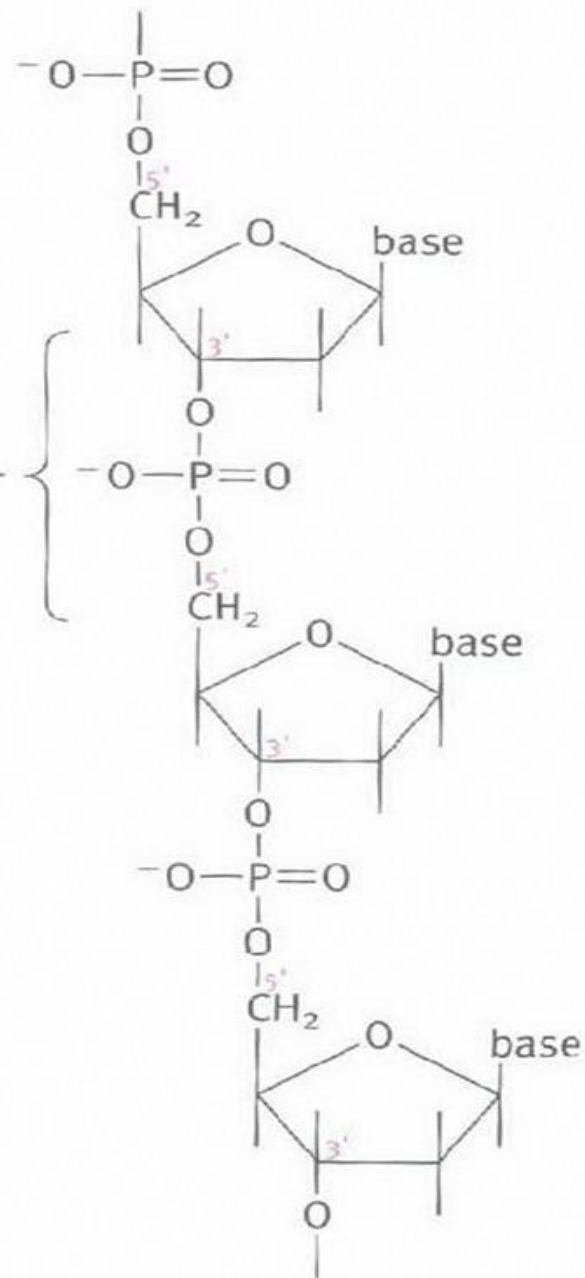
- In 1953 Watson and Crick noted that DNA consists of two polynucleotide strands, running in opposite directions and coiled around each other in a double helix
- Strands are held together by hydrogen bonds between specific pairs of bases
- Adenine (A) and thymine (T) form strong hydrogen bonds to each other but not to C or G
- (G) and cytosine (C) form strong hydrogen bonds to each other but not to A or T

28.10 Base Pairing in DNA: The Watson–Crick Model

- The strands of DNA are complementary because of H-bonding
- Whenever a G occurs in one strand, a C occurs opposite it in the other strand
- When an A occurs in one strand, a T occurs in the other

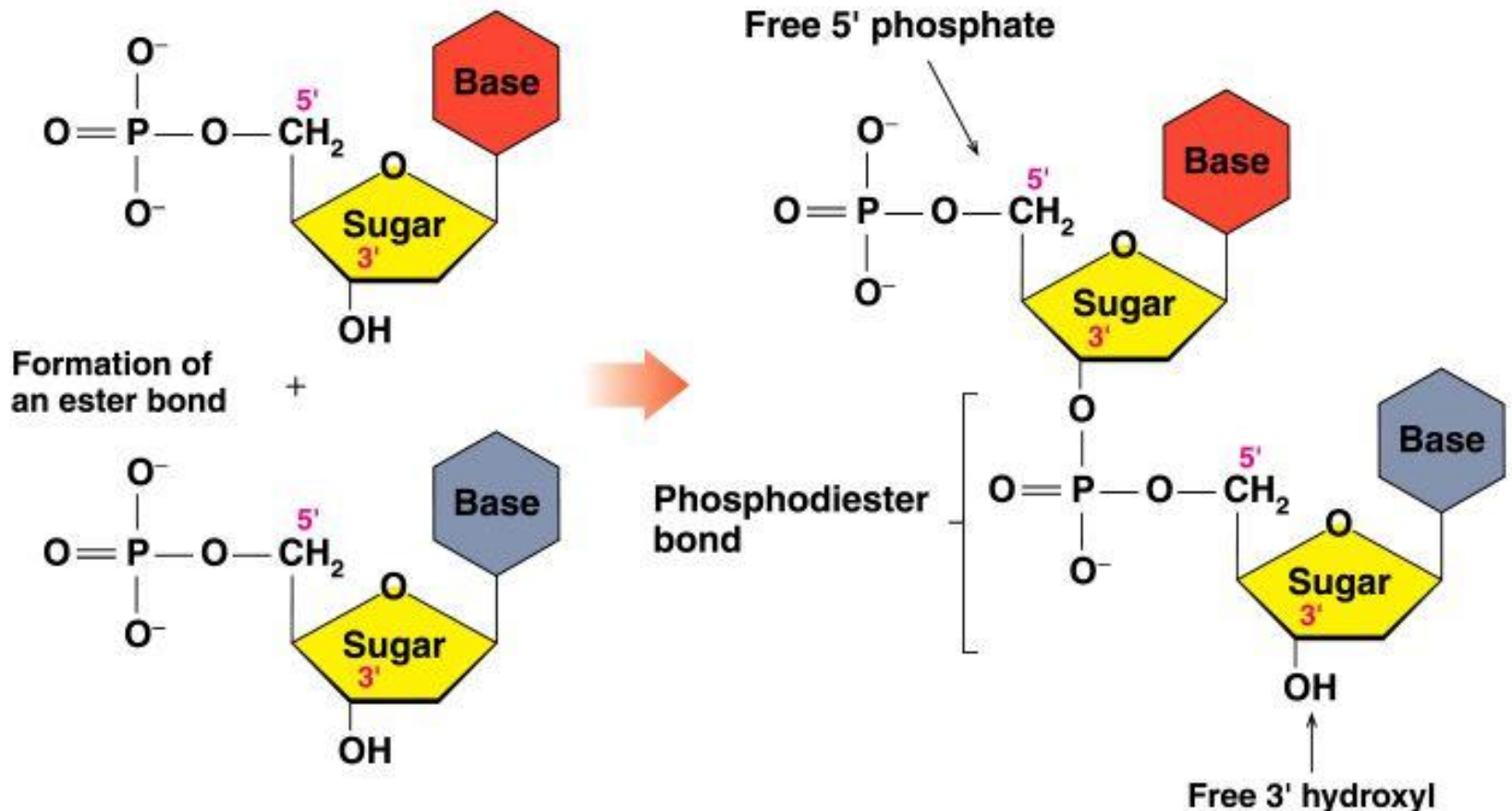
The Difference in the Strands

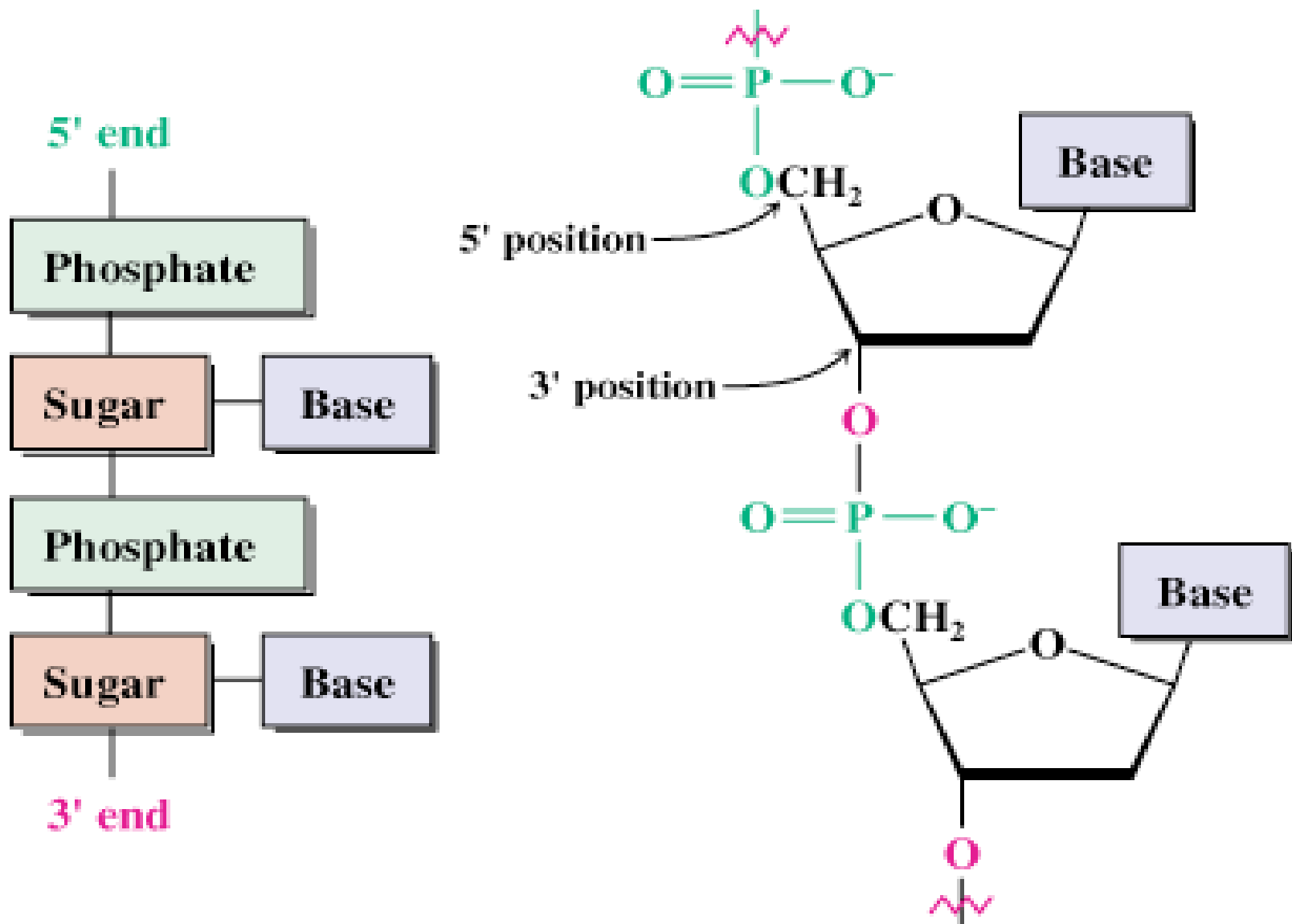
3'-5'
phosphodiester
link



Primary Structure of Nucleic Acids

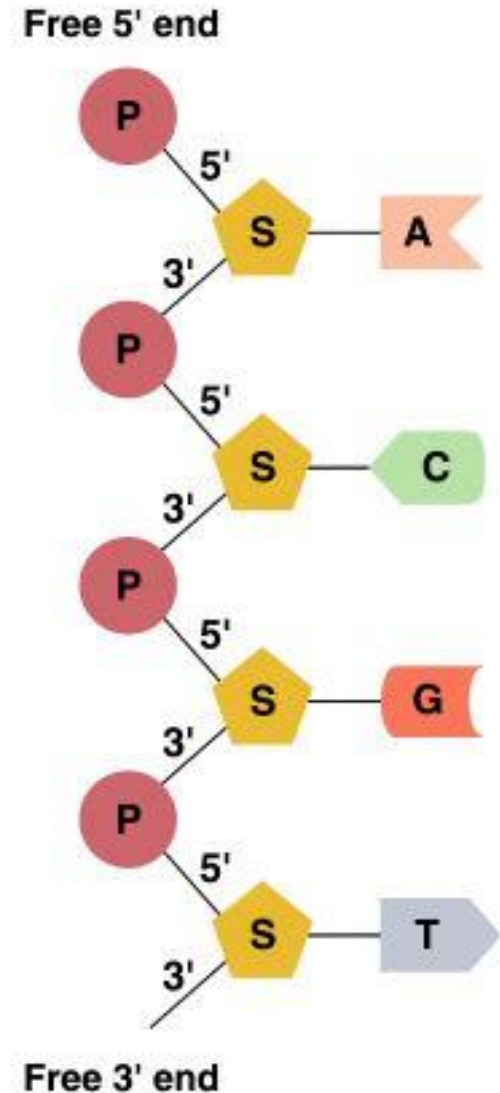
- The **primary structure** of a nucleic acid is the nucleotide sequence
- The nucleotides in nucleic acids are joined by phosphodiester bonds
- The 3'-OH group of the sugar in one nucleotide forms an ester bond to the phosphate group on the 5'-carbon of the sugar of the next nucleotide





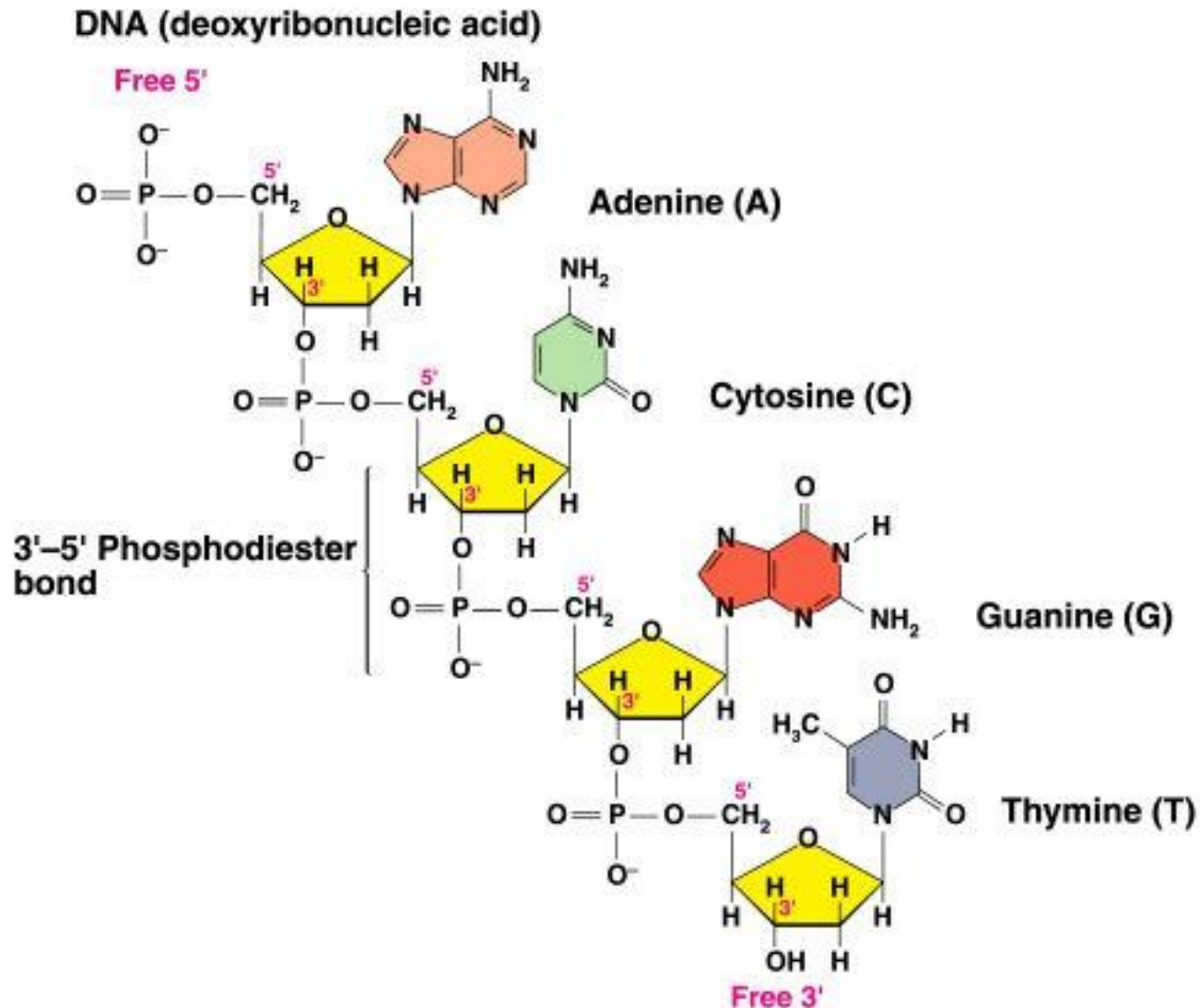
Reading Primary Structure

- A nucleic acid polymer has a free 5'-phosphate group at one end and a free 3'-OH group at the other end
- The sequence is read from the free 5'-end using the letters of the bases
- This example reads 5'—A—C—G—T—3'



Example of DNA Primary Structure

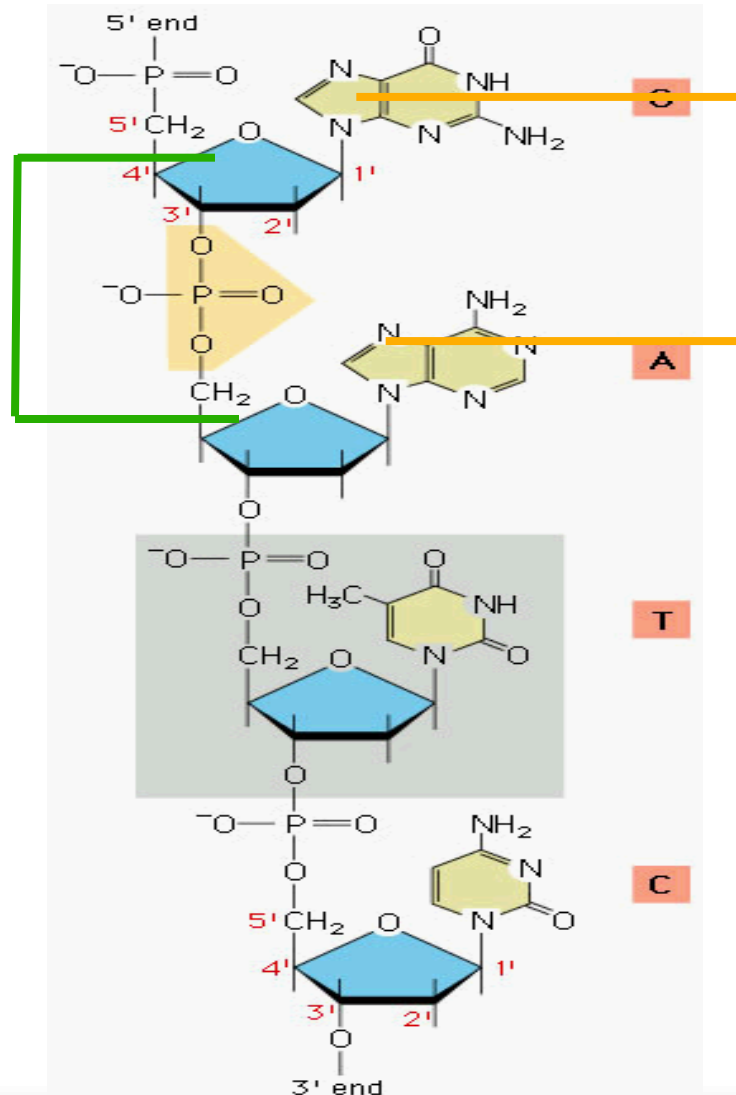
- In DNA, A, C, G, and T are linked by 3'-5' ester bonds between deoxyribose and phosphate



Nucleic Acid Structure

Polymerization

Sugar Phosphate
“backbone”



Nucleotide

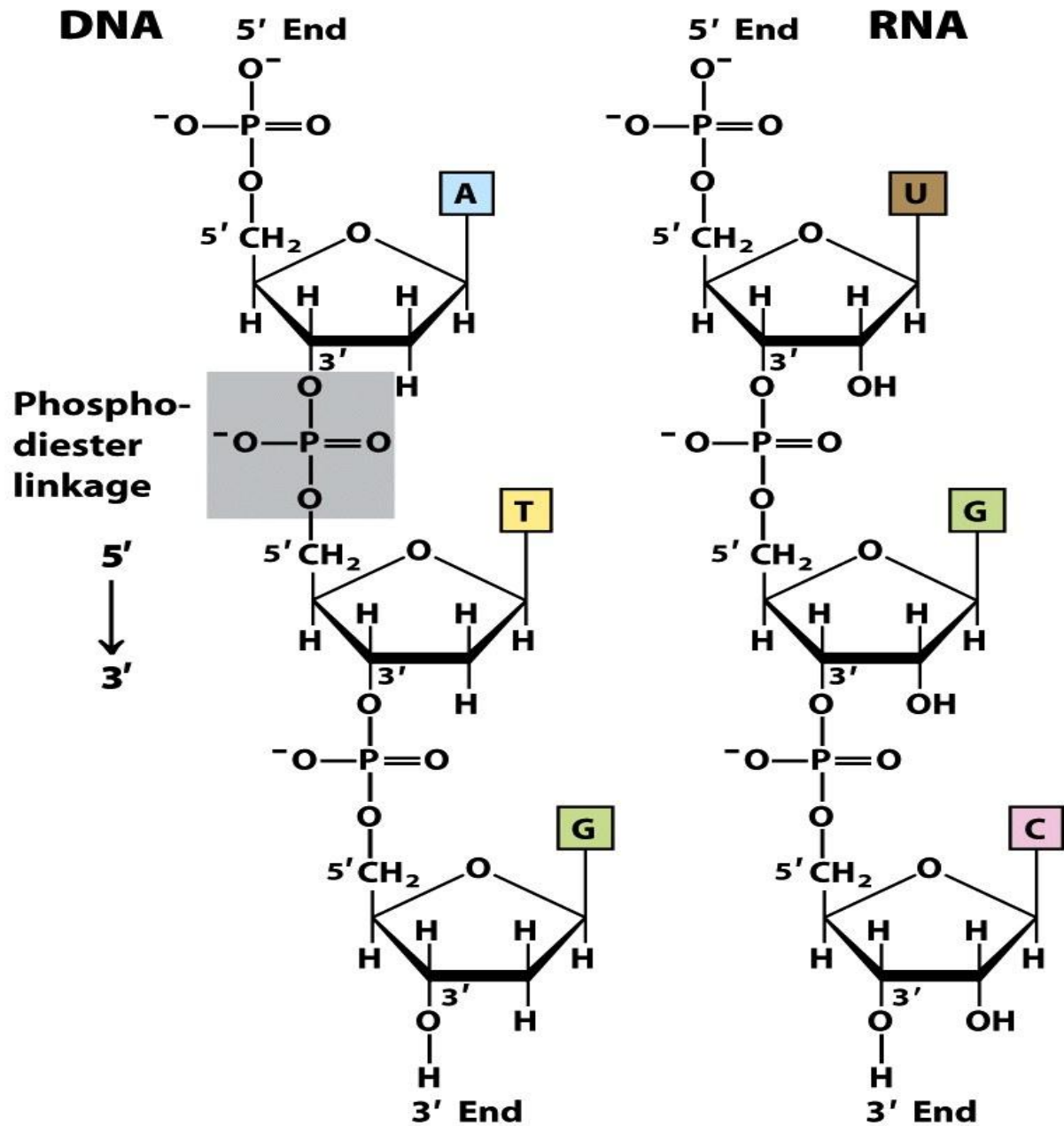


Figure 8-7

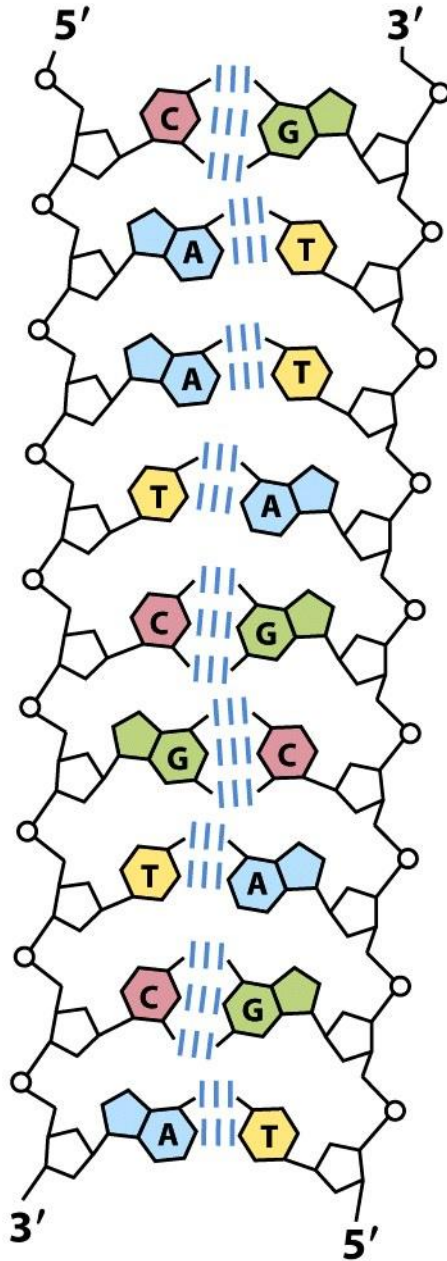
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- Chain is described from 5' end, identifying the bases in order of occurrence, using the abbreviations A for adenosine, G for guanosine, C for cytidine, and T for thymine (or U for uracil in RNA)

Describing a Sequence

- A typical sequence is written as TAGGCT



Properties of a DNA double helix

The strands of DNA are antiparallel

The strands are complementary

There are Hydrogen bond forces

There are base stacking interactions

There are 10 base pairs per turn

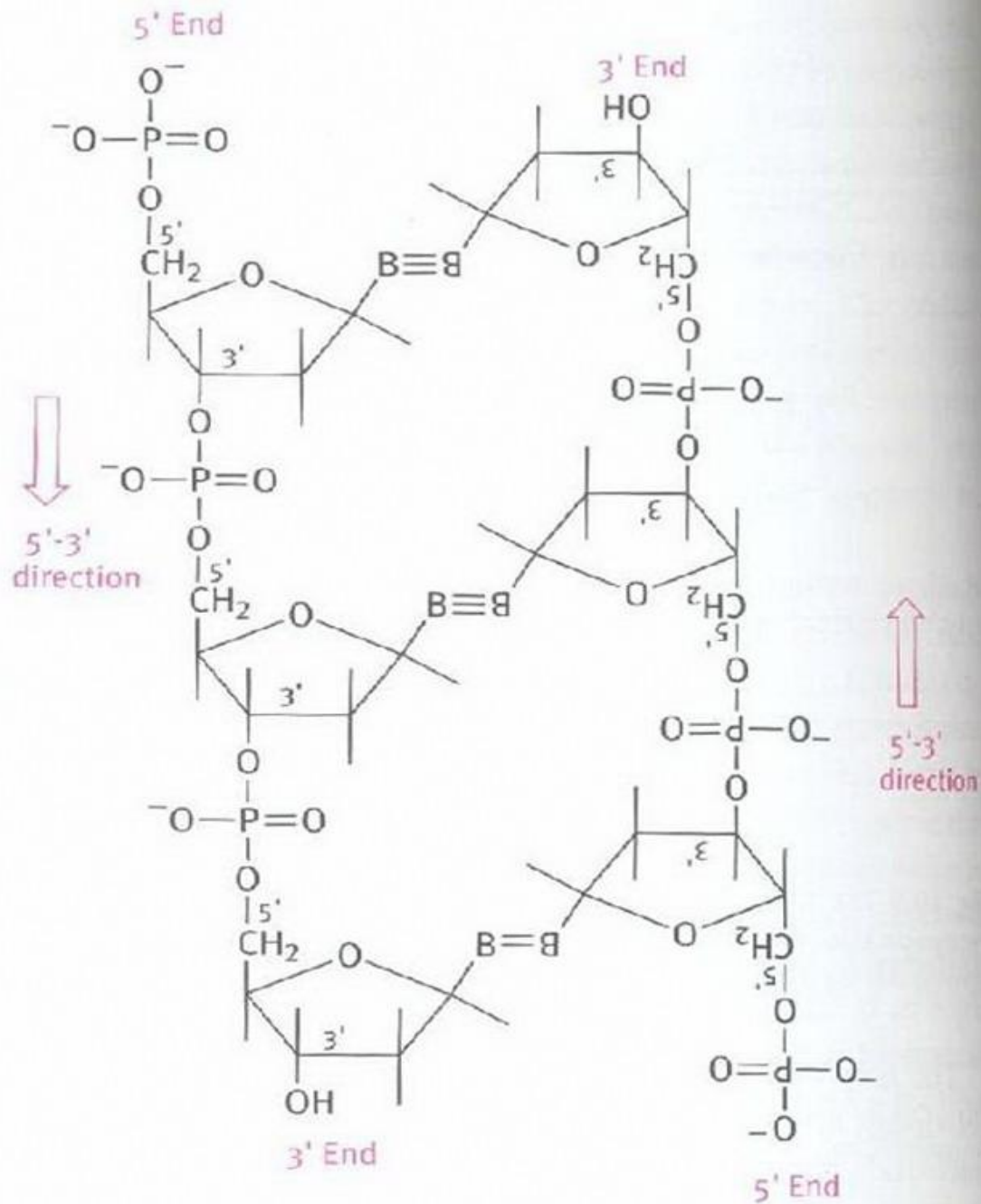


Fig. 20.6 Two antiparallel strands of DNA. B, Base.

The Double Helix (DNA)

Structural model:

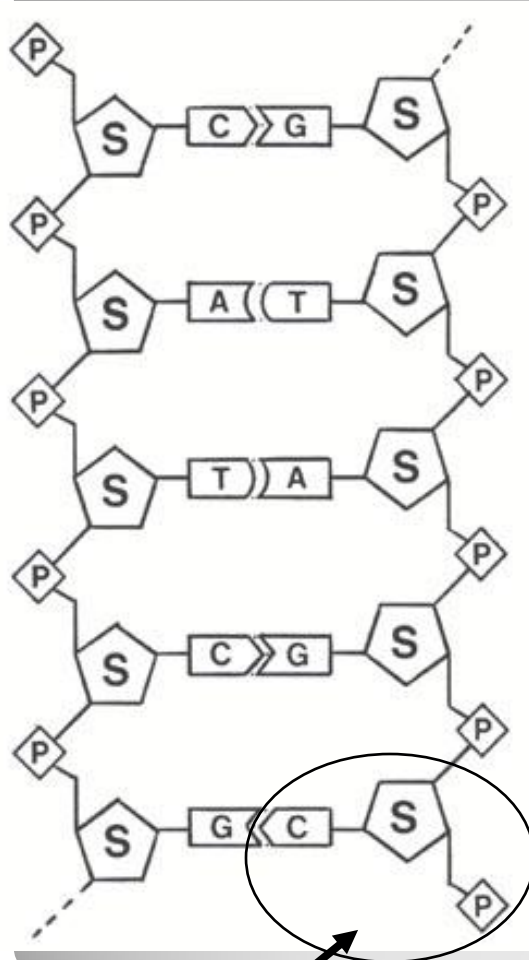
- Model proposed by Watson & Crick, 1953
- Two sugar-phosphate strands, next to each other, but running in opposite directions.
- **Specific Hydrogen bonds** occur among bases from one chain to the other:



Due to this specificity, a certain base on one strand indicates a certain base in the other.

- The 2 strands intertwine, forming a double-helix that winds around a central axis

Untwisted it
looks like this:



Nucleotide

- The sides of the ladder are:

P = phosphate

S = sugar molecule

- The steps of the ladder are C, G, T, A = nitrogenous bases
(Nitrogenous means containing the element nitrogen.)

A = Adenine (Apples are Tasty)

T = Thymine

A always pairs with T in DNA

C = Cytosine (Cookies are Good)

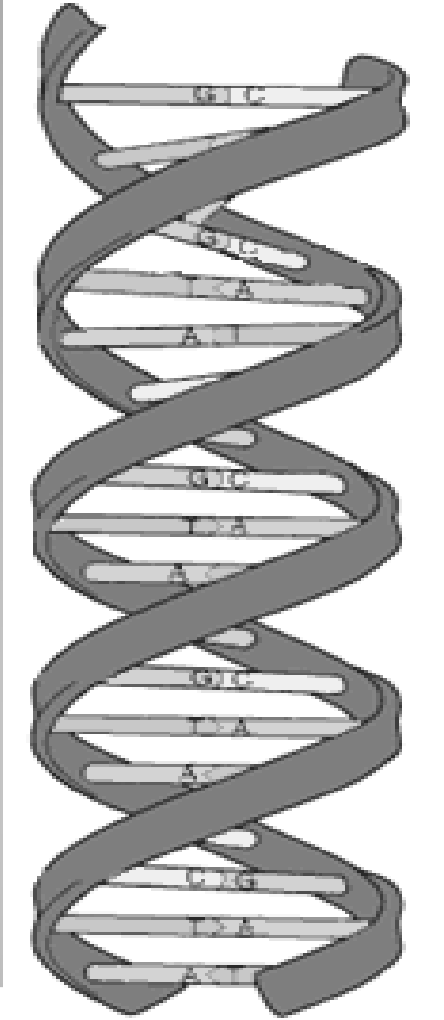
G = Guanine

C always pairs with G in DNA

Secondary Structure: DNA Double Helix

- In DNA there are two strands of nucleotides that wind together in a **double helix**
 - the strands run in opposite directions
 - the bases are arranged in step-like pairs
 - the **base pairs** are held together by hydrogen bonding
- The pairing of the bases from the two strands is very specific
- The **complimentary base pairs** are **A-T** and **G-C**
 - two hydrogen bonds form between A and T
 - three hydrogen bonds form between G and C
- Each pair consists of a purine and a pyrimidine, so they are the same width, keeping the two strands at equal distances from each other

Model of DNA:



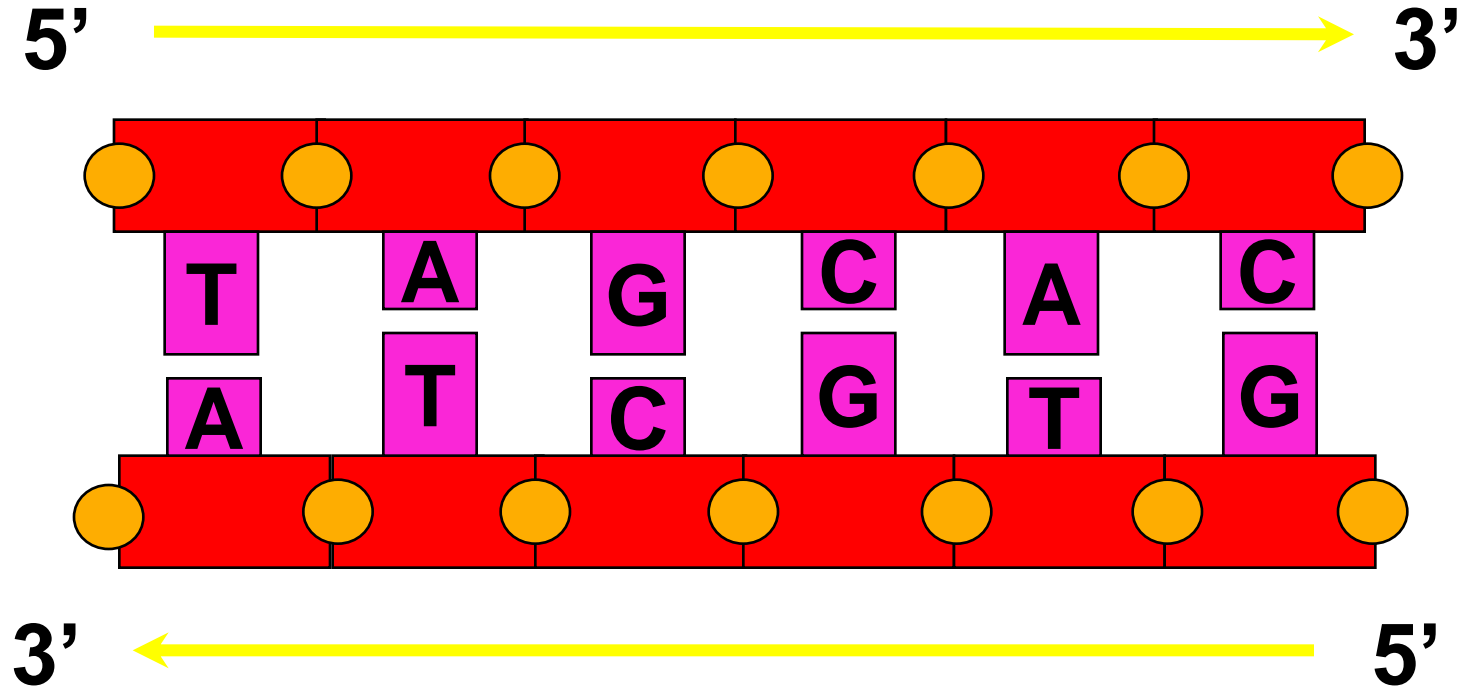
- The model was developed by Watson and Crick in 1953.
- They received a nobel prize in 1962 for their work.
- The model looks like a twisted ladder – double helix.

Nucleic Acid Structure

“Base Pairing”

DNA base-pairing is **antiparallel**

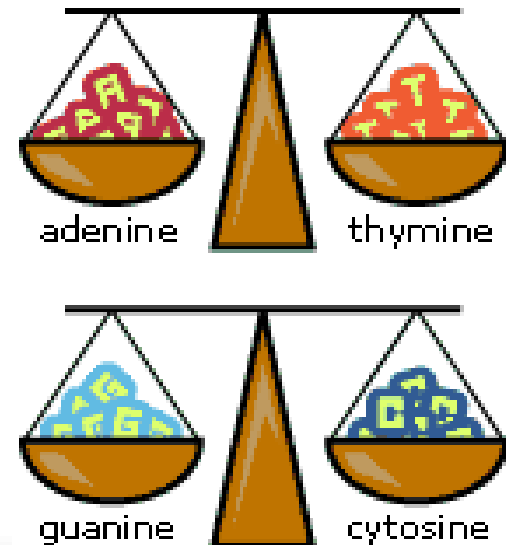
i.e. 5' - 3' (l-r) on top : 5' - 3' (r-l) on



Discovering the structure of DNA

Erwin Chargaff – (1905-2002)

- Columbia University, NY
- Investigated the composition of DNA
- His findings by 1950 strongly suggested the base-pairings of A-T & G-C
- Met with Watson and Crick in 1952 and shared his findings
- “Chargaff’s rule” $A = T$ & $C = G$



Nucleic Acid Structure

The double helix

First determined by Watson & Crick in 1953

Most energy favorable conformation for double stranded DNA to form

Shape and size is uniform for all life (i.e. DNA is identical)

Without anti-parallel base pairing this conformation could not exist

Structure consists of “major” grooves and “minor” grooves

Major grooves are critical for binding proteins that regulate DNA function

Discovering the structure of DNA

- **DNA = Deoxyribose nucleic acid**

- **Present in all living cells**

- **Contains all the information**

- **Nucleotides:**

- **a subunit that consist**

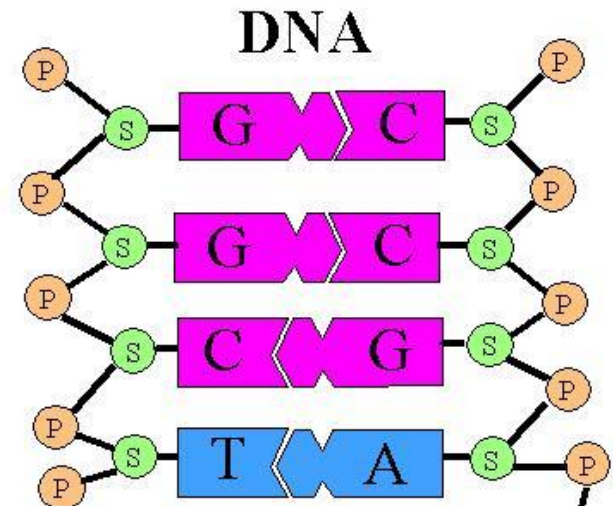
- **a sugar (deoxyribose)**

- **a phosphate**

- **and one nitrogen base – 4 different bases**

- **Adenine (A) and Thymine (T)**

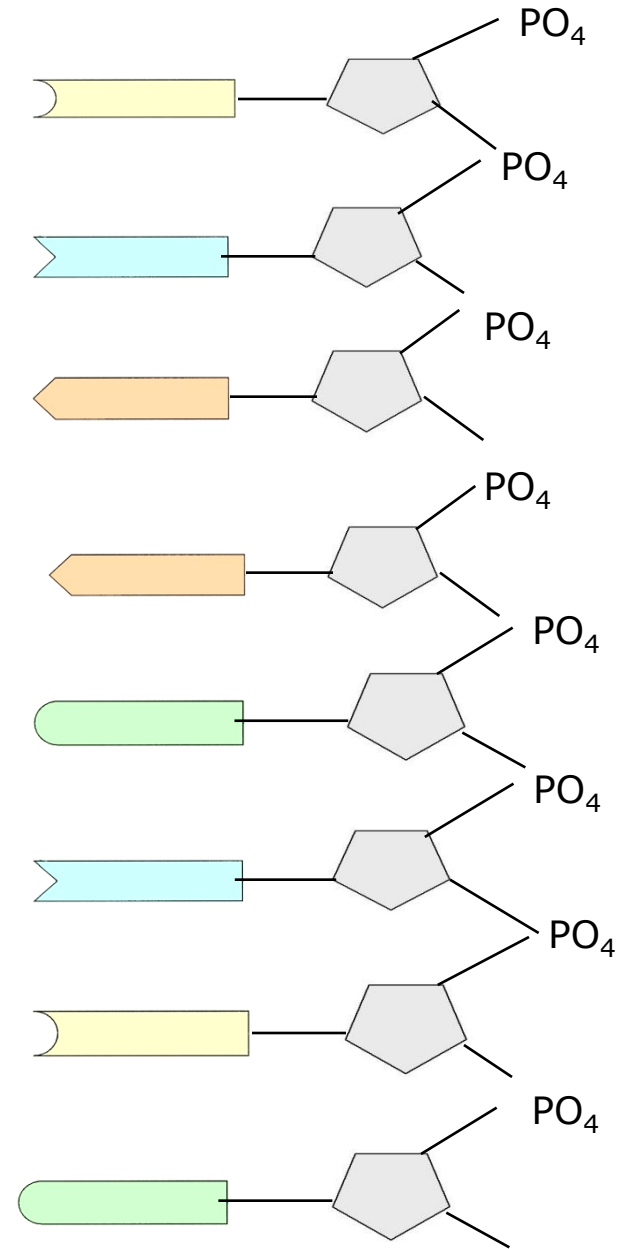
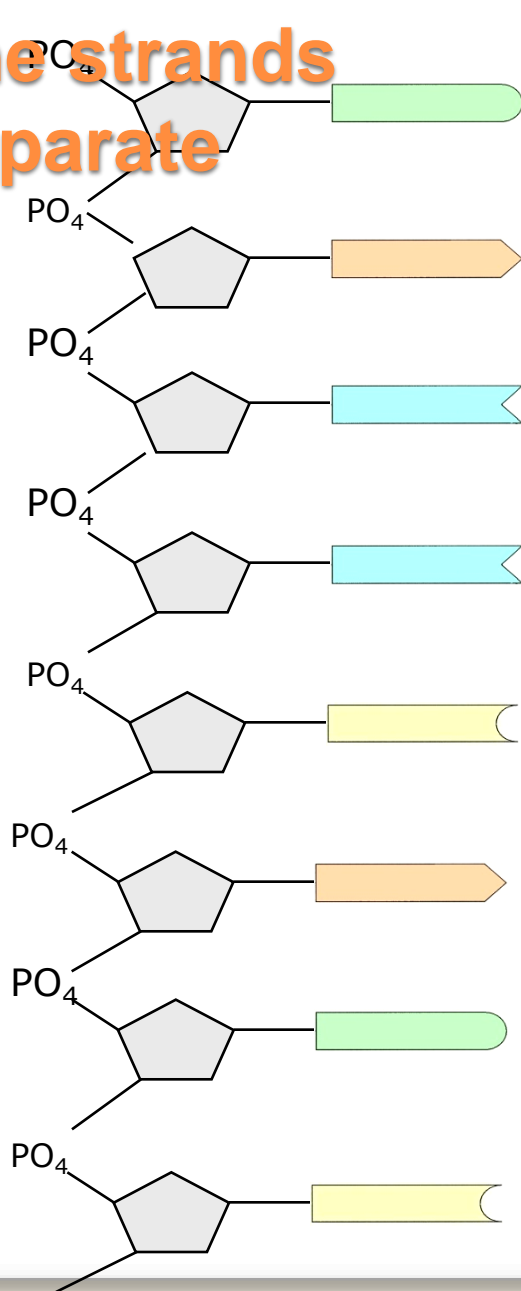
- **Guanine (G) and Cytosine (C)**



The paired strands are coiled into a spiral called

A DOUBLE HELIX

The strands
separate



Nucleic Acid Structure

“Base Pairing”

RNA [normally] exists as a single stranded polymer

DNA exists as a double stranded polymer

DNA double strand is created by hydrogen bonds between nucleotides

Nucleotides always bind to complementary nucleotides

A  **T** (2 H-bonds)

G  **C** (3 H-bonds)

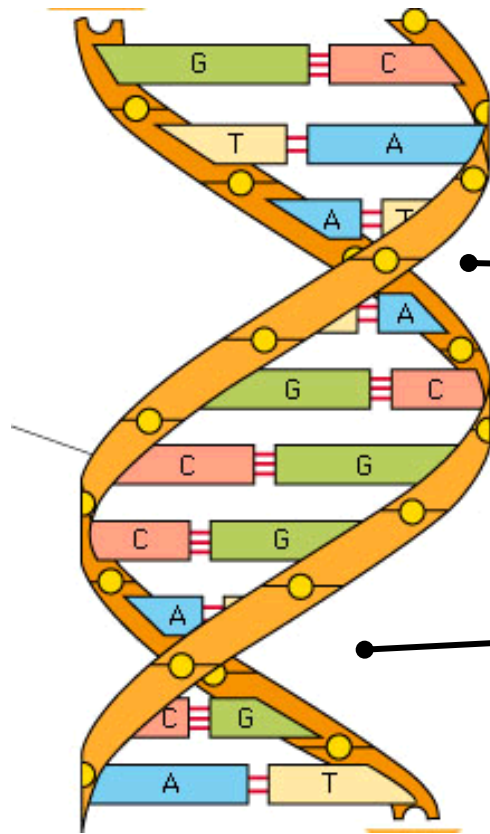
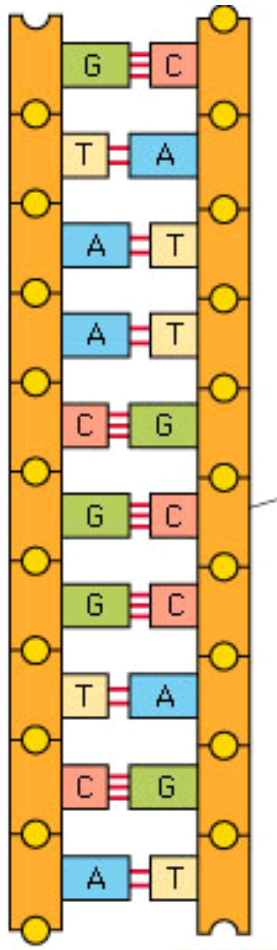


Practice DNA Base Pairs

G A T T A C A
C T A A T G T

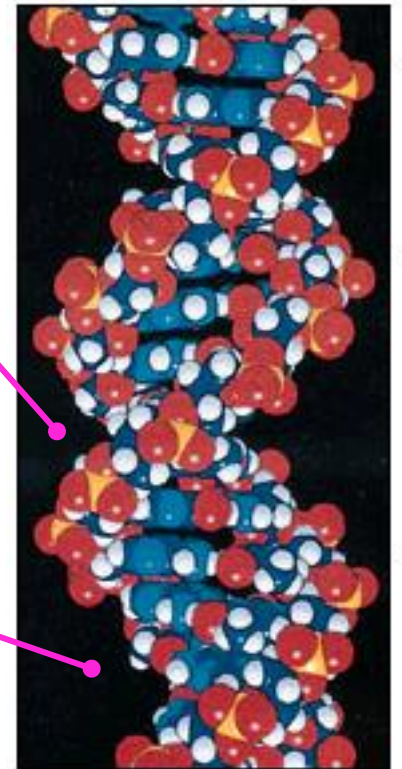
Nucleic Acid Structure

The double helix



Minor
Groove

Major
Groove



Before a cell divides, the DNA strands uncoil and separate

Each strand makes a new partner by adding the appropriate nucleotides

The result is that there are now two double-stranded DNA molecules in the nucleus

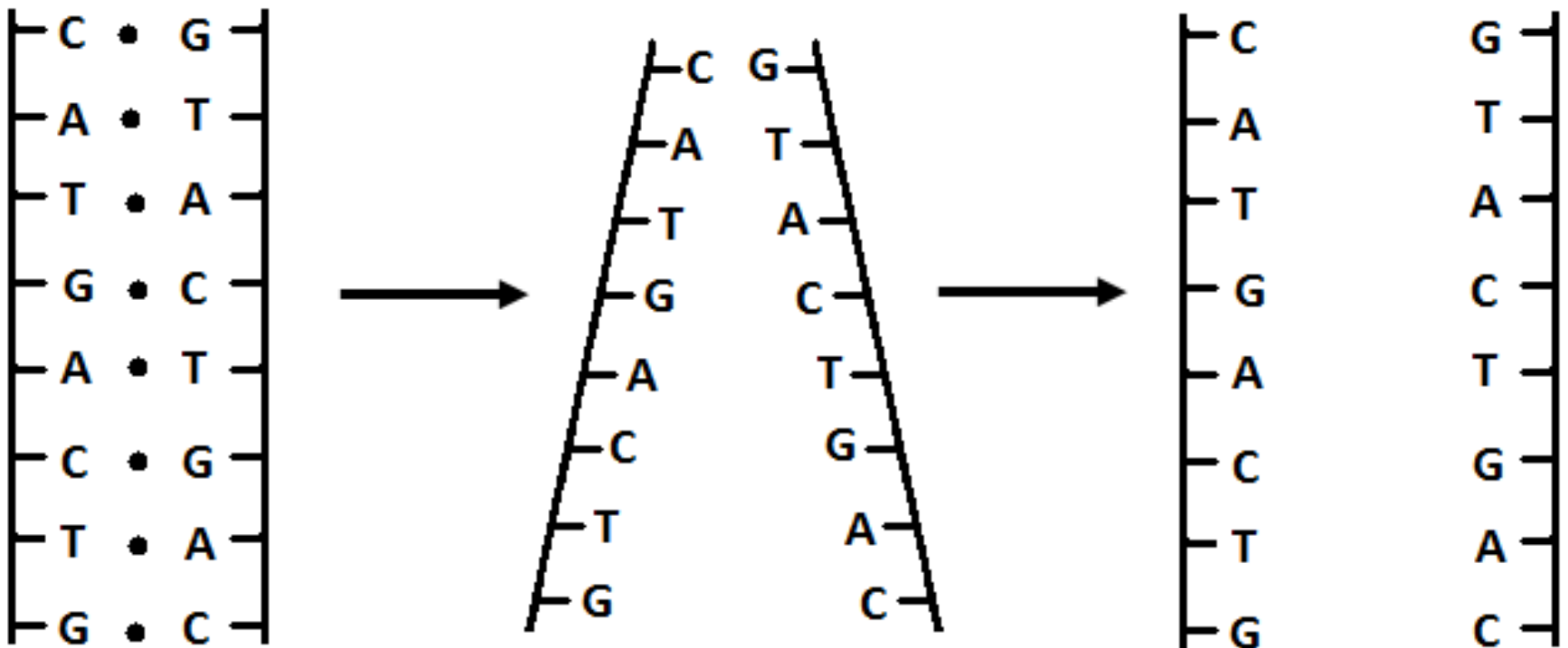
So that when the cell divides, each nucleus contains identical DNA

This process is called replication

STEP 1

Hydrogen bonds between base pairs are broken by the enzyme Helicase and DNA molecule unzips

DNA molecule separates into complementary halves



Complementarity of DNA strands

- Two chains differ in sequence
(sequence is read from 5' to 3')
- Two chains are **complementary**
- Two chains run antiparallel

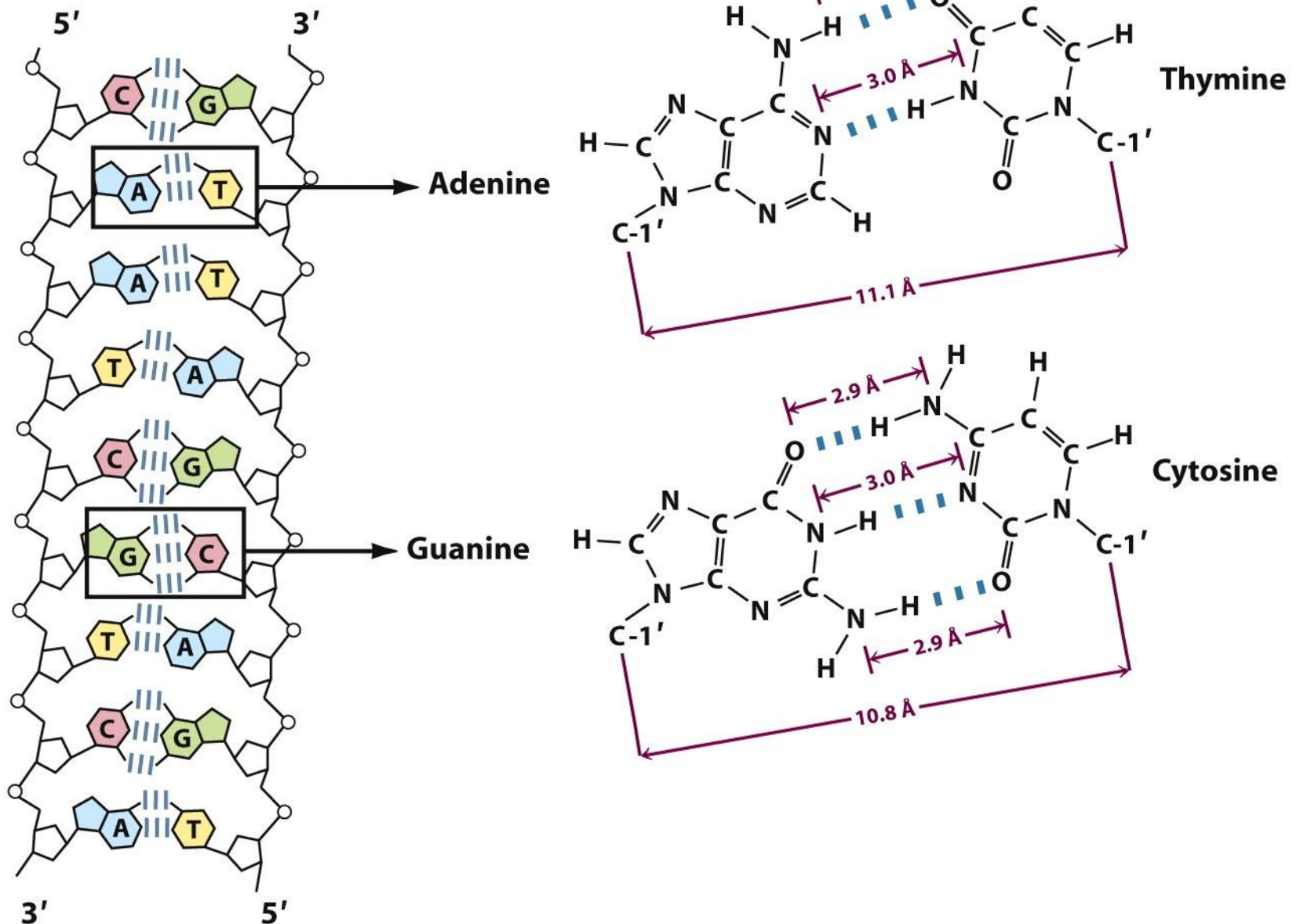


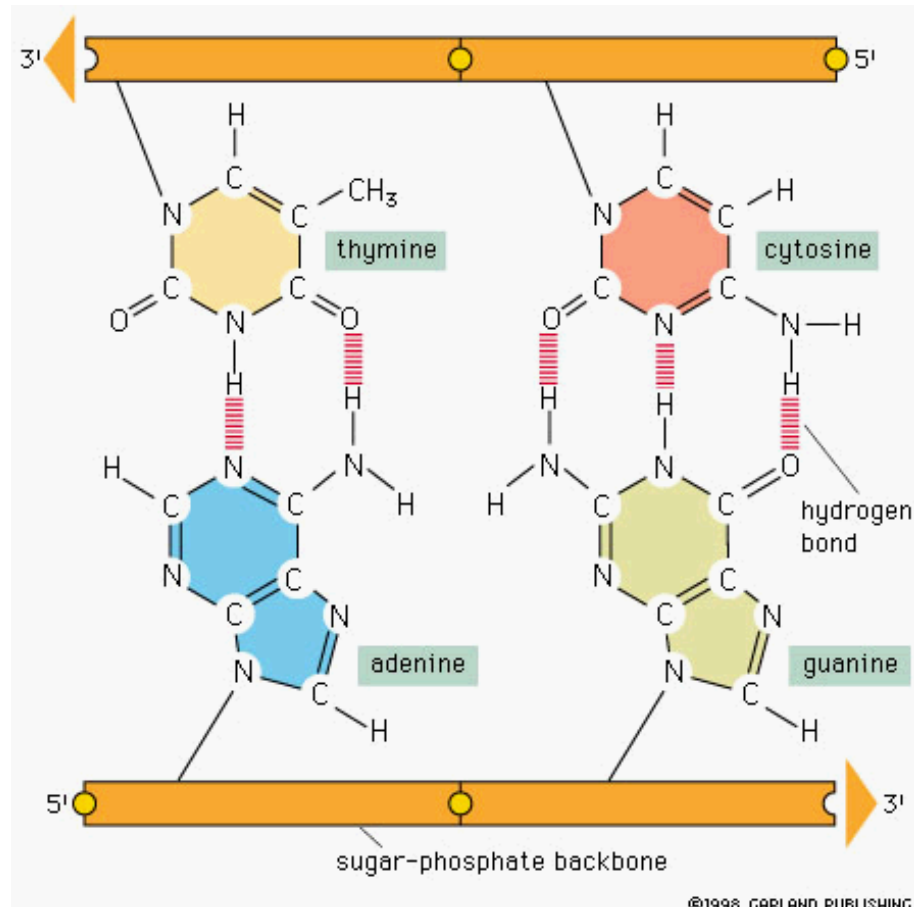
Figure 8-11

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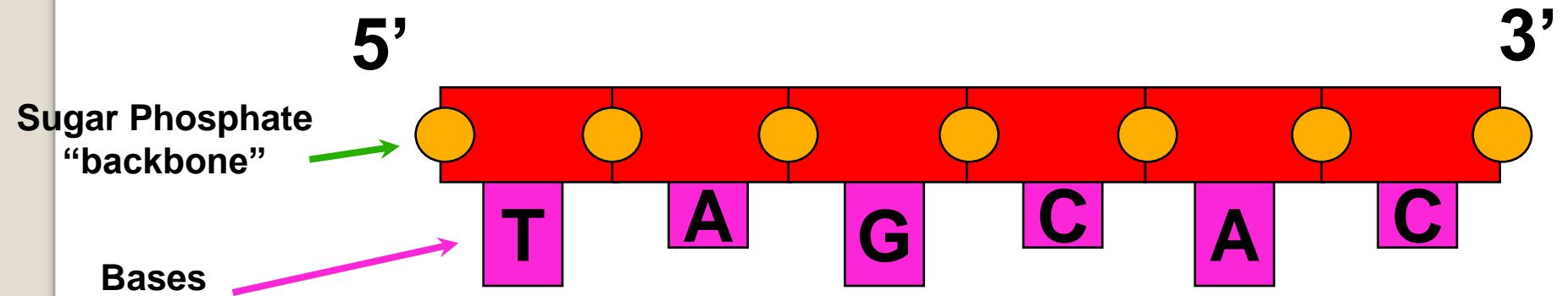
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Nucleic Acid Structure

“Base Pairing”

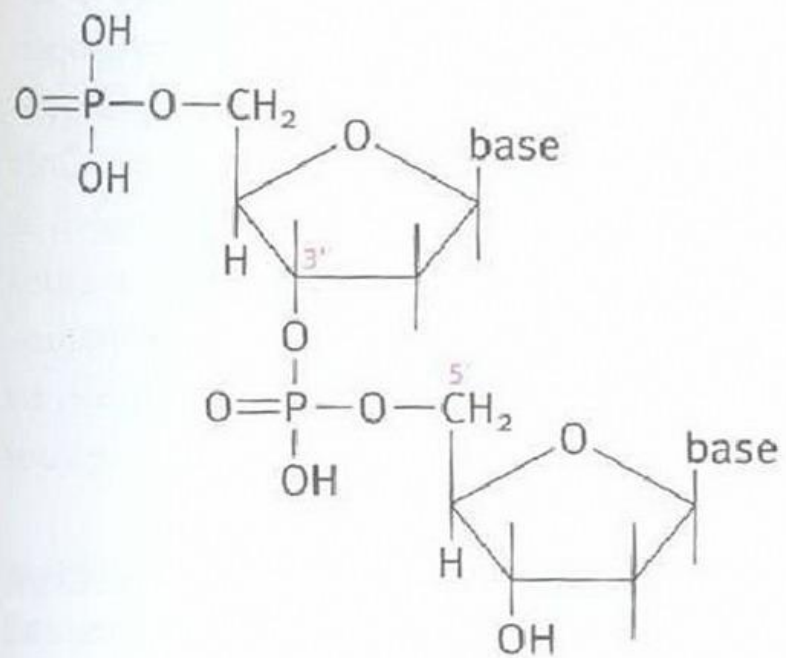
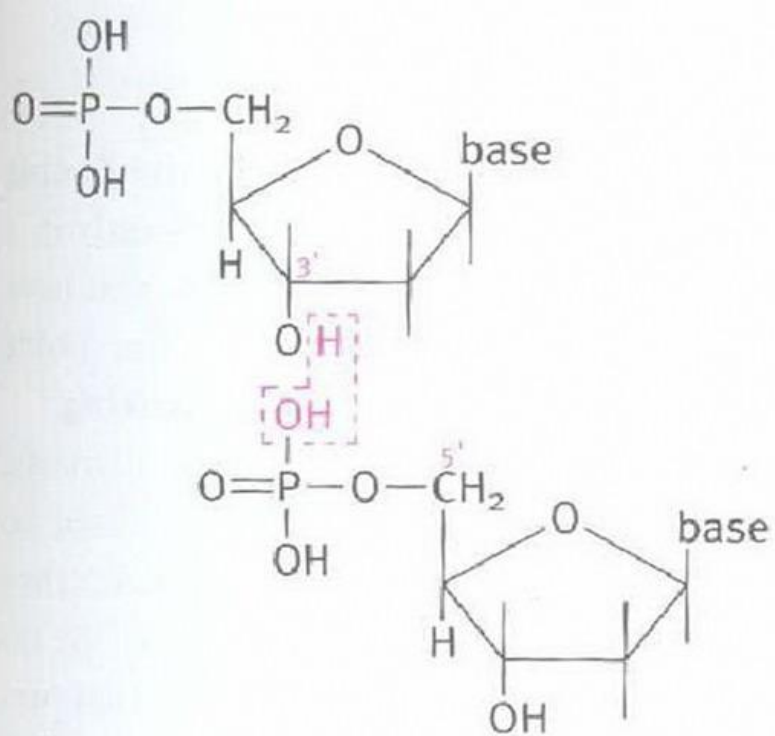


Nucleic Acid Structure Polymerization

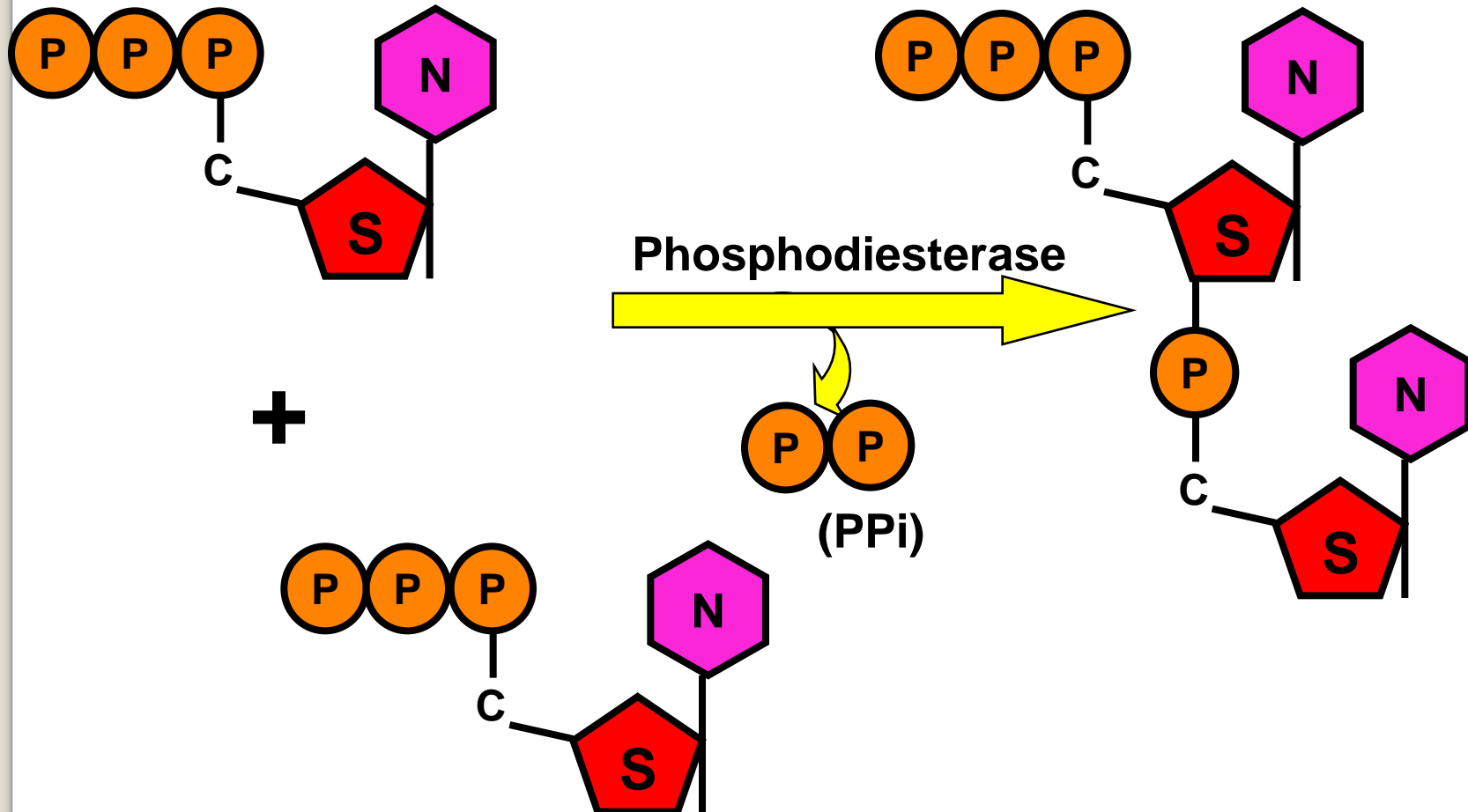


5' 3'

TAGCAC



Nucleic Acid Structure Polymerization



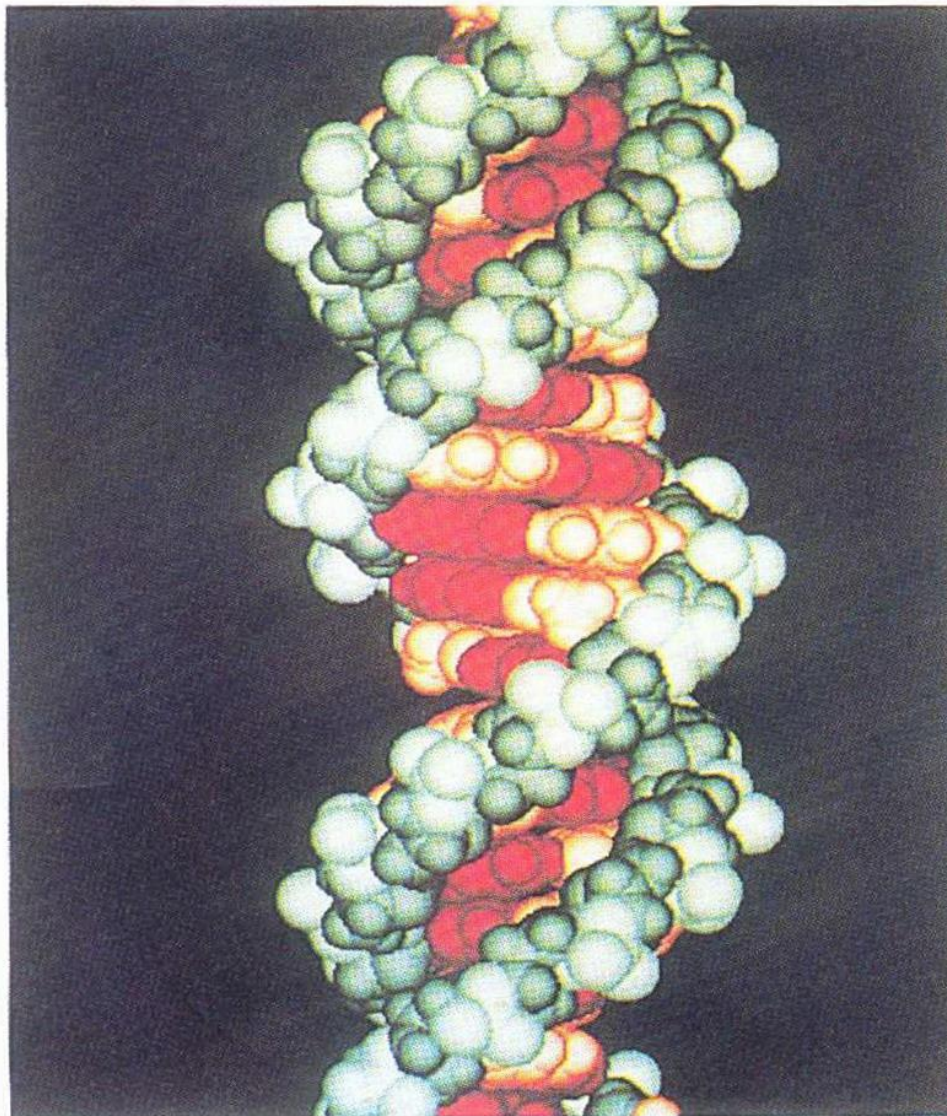
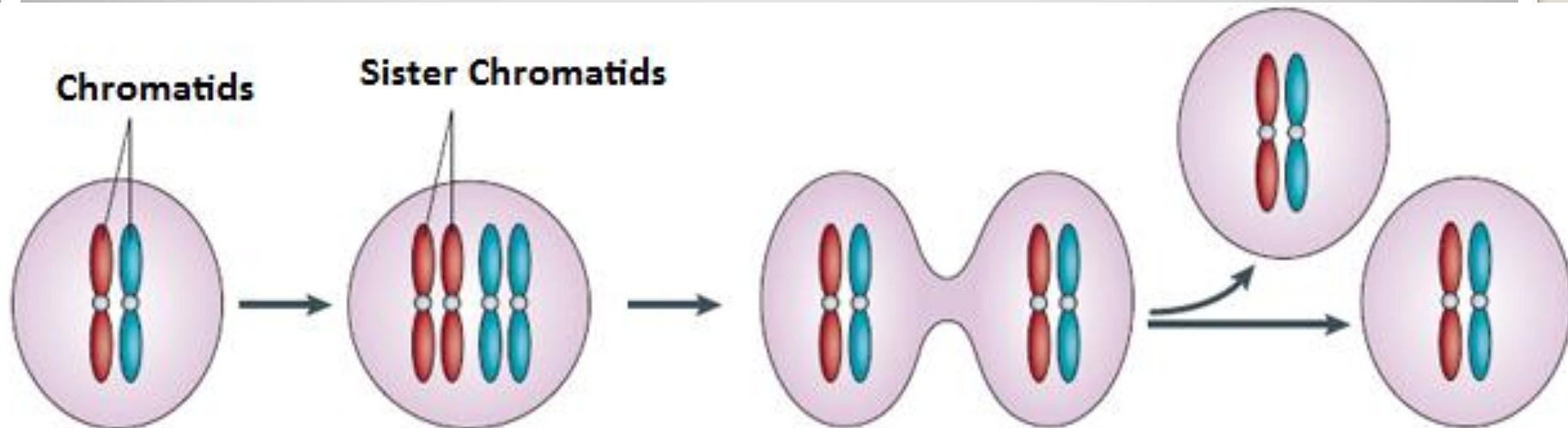


Fig. 20.5 A model of B DNA. Space-filling atomic model of a DNA segment with two major grooves and one minor groove.

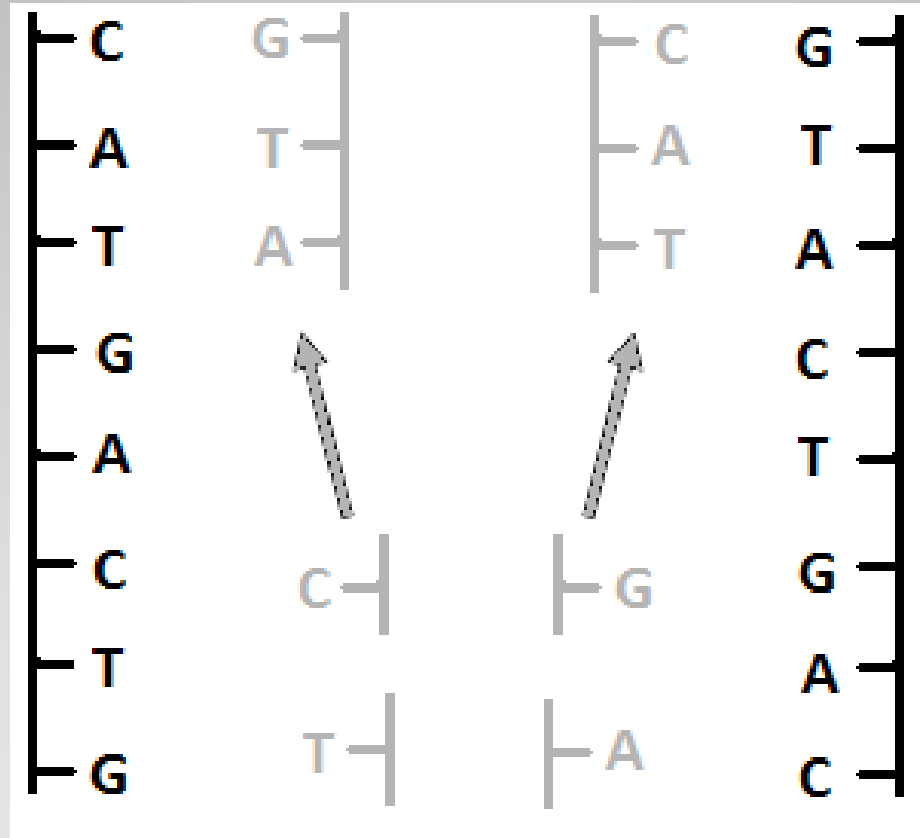
DNA Replication

- Cell division involving mitosis produces 2 daughter cells that are genetically identical to each other and genetically identical to the parent cell
- Remember that for this to happen, DNA in the parent cell must be replicated (copied) before the cell divides – this process occurs during Interphase in the cell cycle



STEP 2

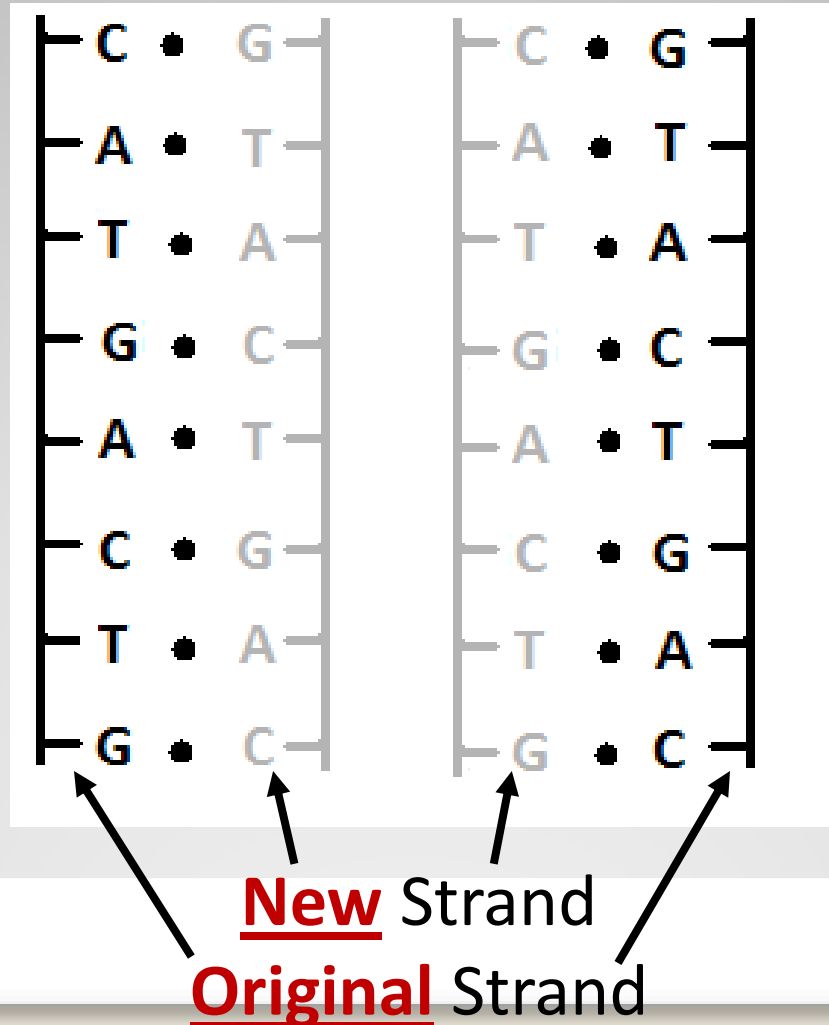
Nucleotides match up with complementary bases



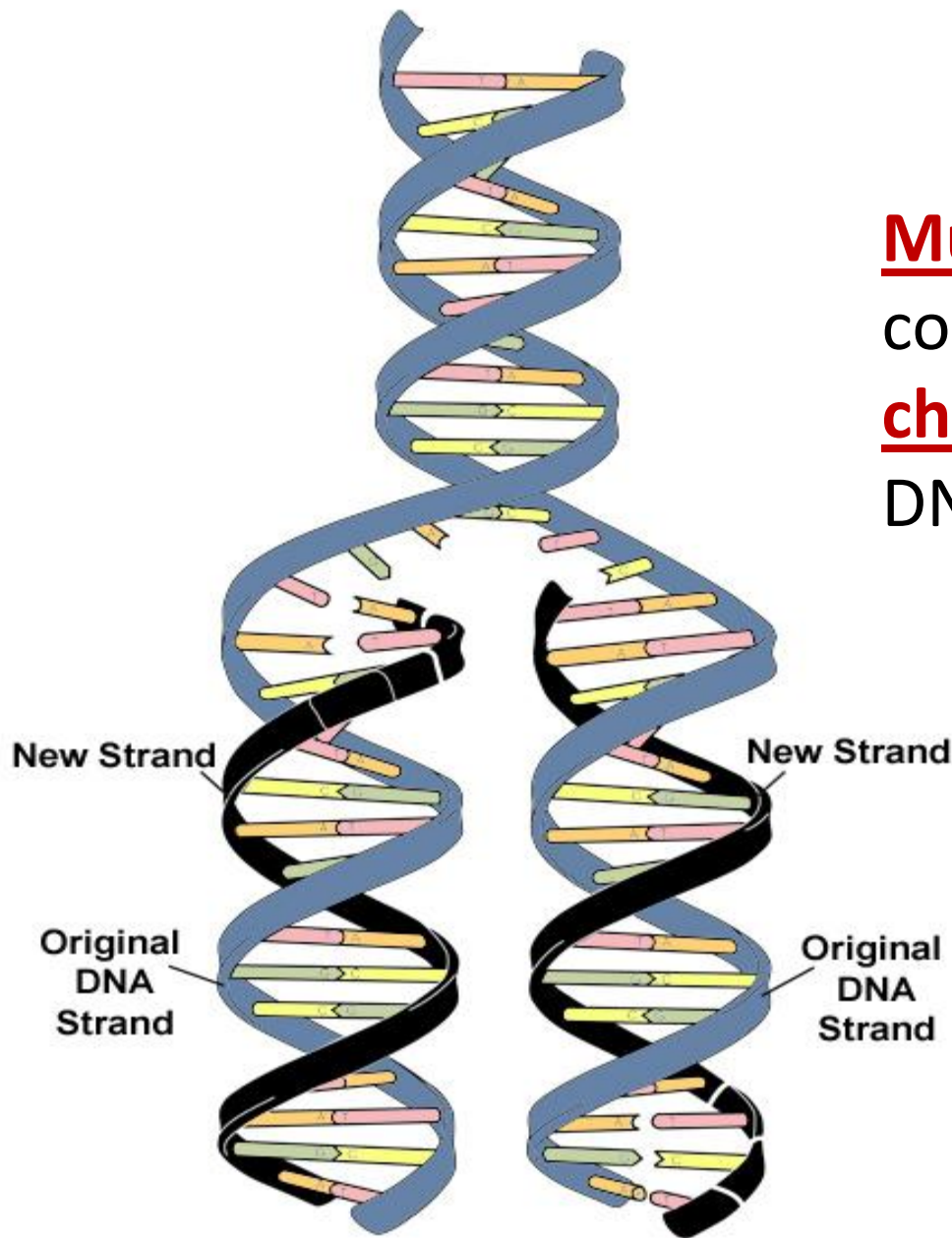
Free nucleotides abundant in nucleus

STEP 3

Nucleotides are linked into 2 new strands of DNA by the enzyme, polymerase—DNA polymerase also proofreads for copying errors



Mutations occur when copying **errors** cause a **change** in the **sequence** of DNA nucleotide bases



DNA Replication

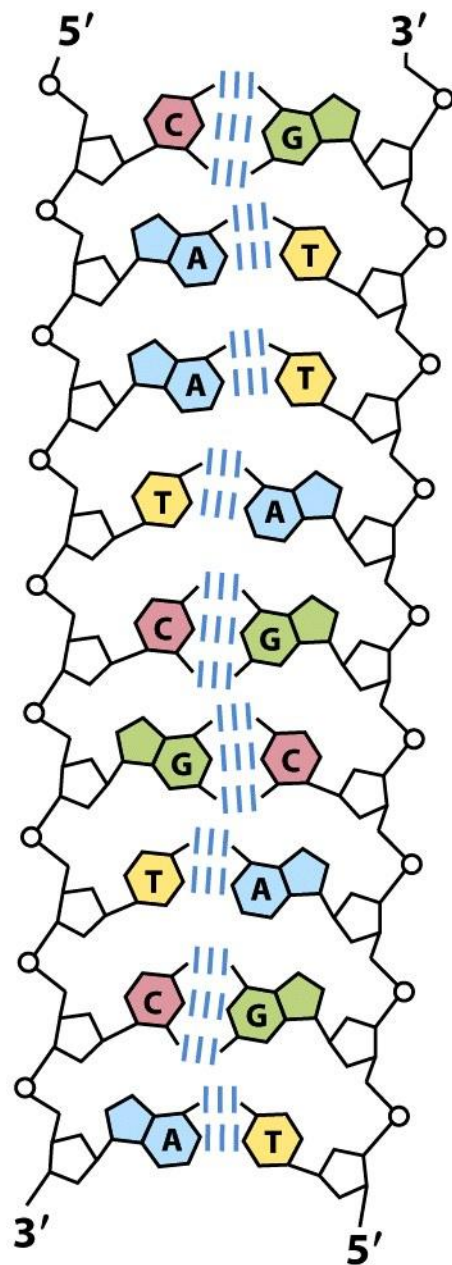


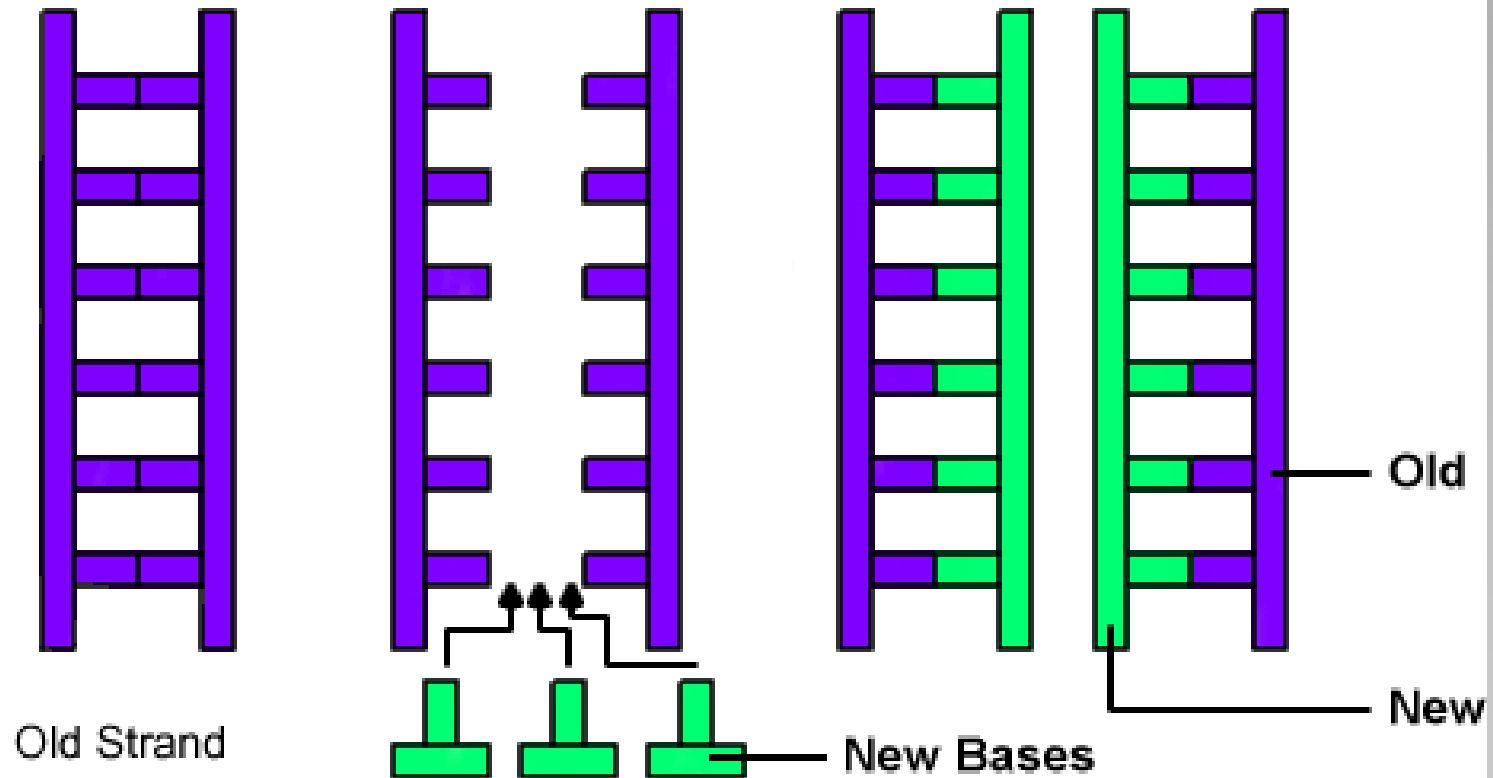
Figure 8-14

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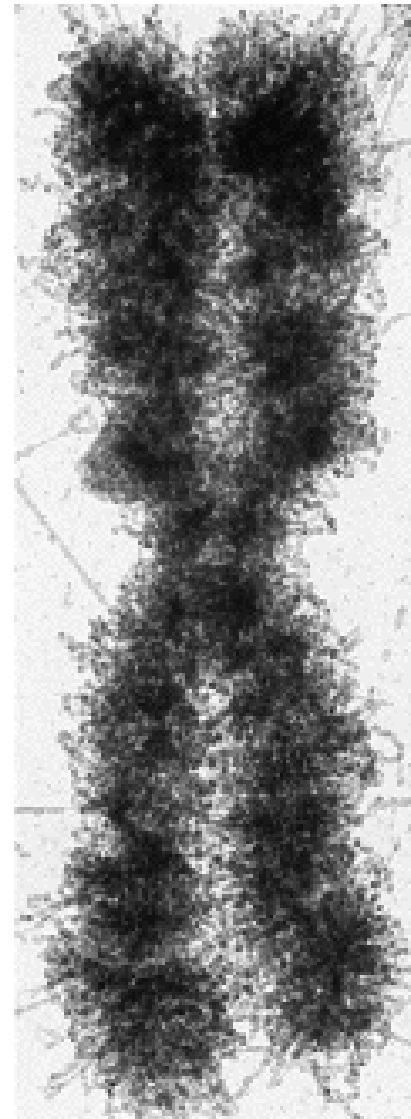
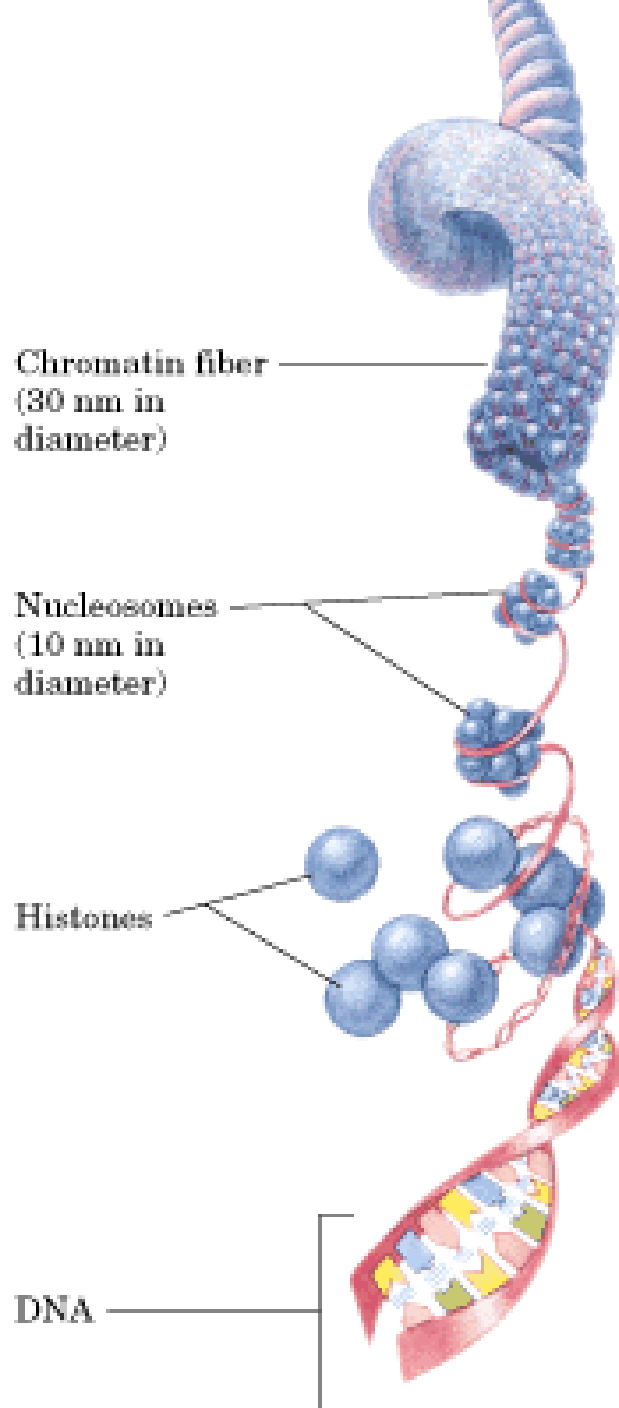
Diagram Examples of DNA Replication:

(You could see DNA replication represented different ways.)



Double helix unzips
New bases (A,T,G,C)
are added

Two new strands are
created, each
contain half of the
original strand.



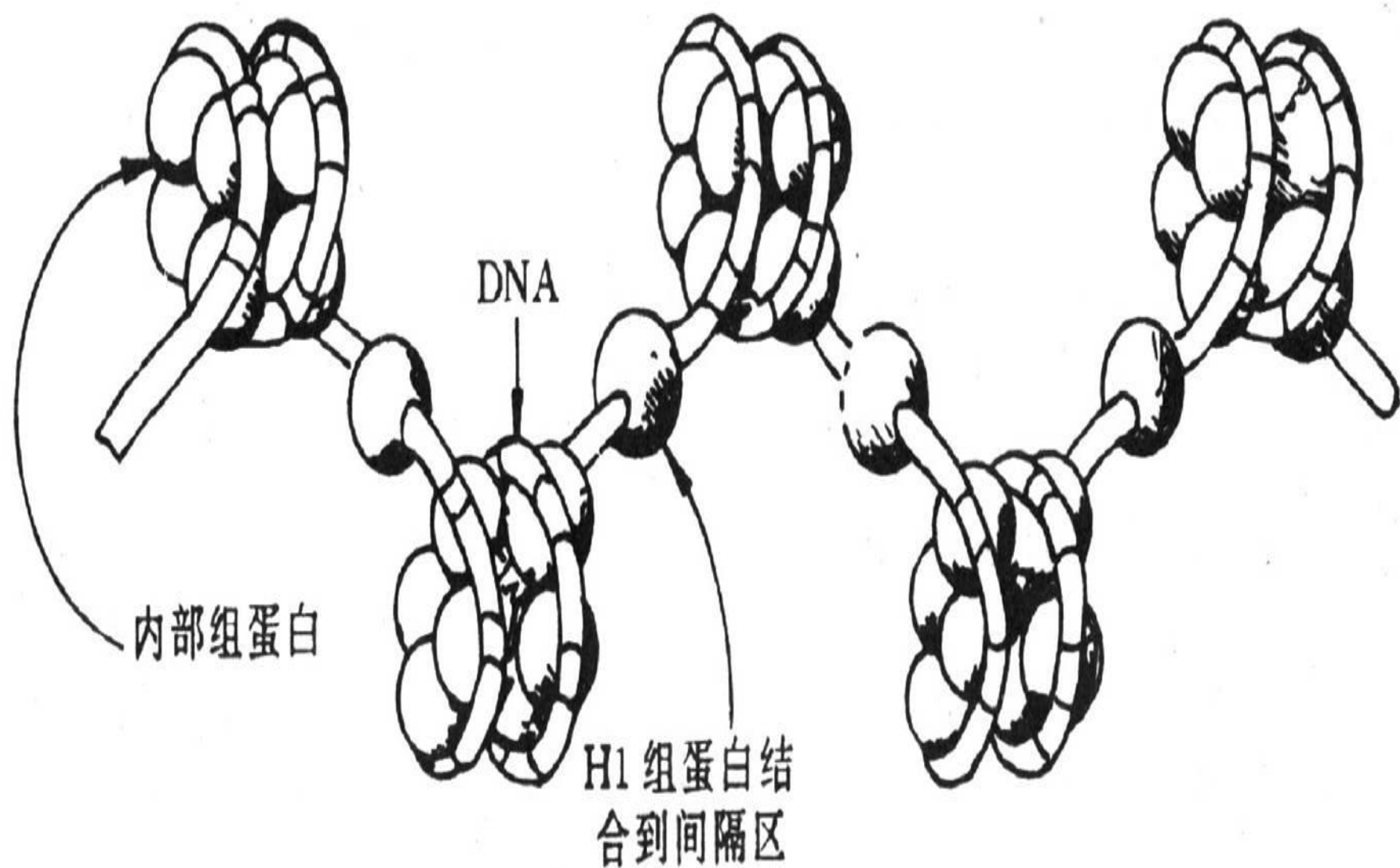
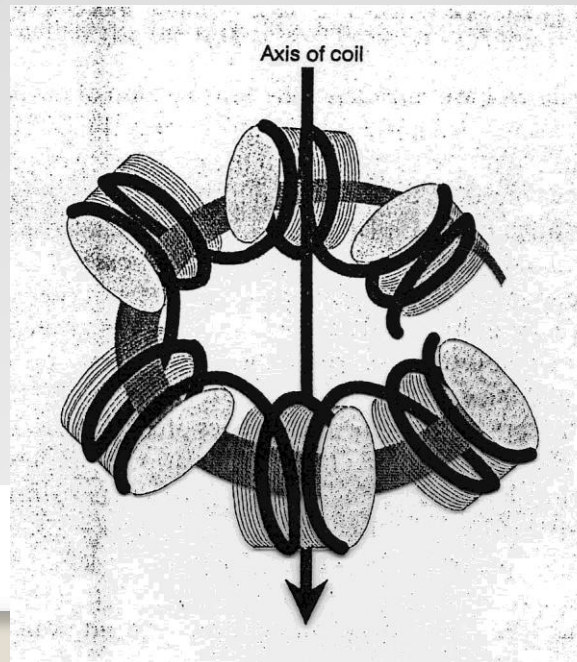


图 3-14 核小体结构

Storage of DNA

- In **eukaryotic** cells (animals, plants, fungi) DNA is stored in the **nucleus**, which is separated from the rest of the cell by a semipermeable membrane
- The DNA is only organized into **chromosomes** during cell replication
- Between replications, the DNA is stored in a compact ball called **chromatin**, and is wrapped around proteins called **histones** to form **nucleosomes**



DNA Replication

- When a eukaryotic cell divides, the process is called **mitosis**
 - the cell splits into two identical daughter cells
 - the DNA must be replicated so that each daughter cell has a copy
- **DNA replication** involves several processes:
 - first, the DNA must be unwound, separating the two strands
 - the single strands then act as templates for synthesis of the new strands, which are complimentary in sequence
 - bases are added one at a time until two new DNA strands that exactly duplicate the original DNA are produced
- The process is called **semi-conservative replication** because one strand of each daughter DNA comes from the parent DNA and one strand is new
- The energy for the synthesis comes from hydrolysis of phosphate groups as the phosphodiester bonds form between the bases

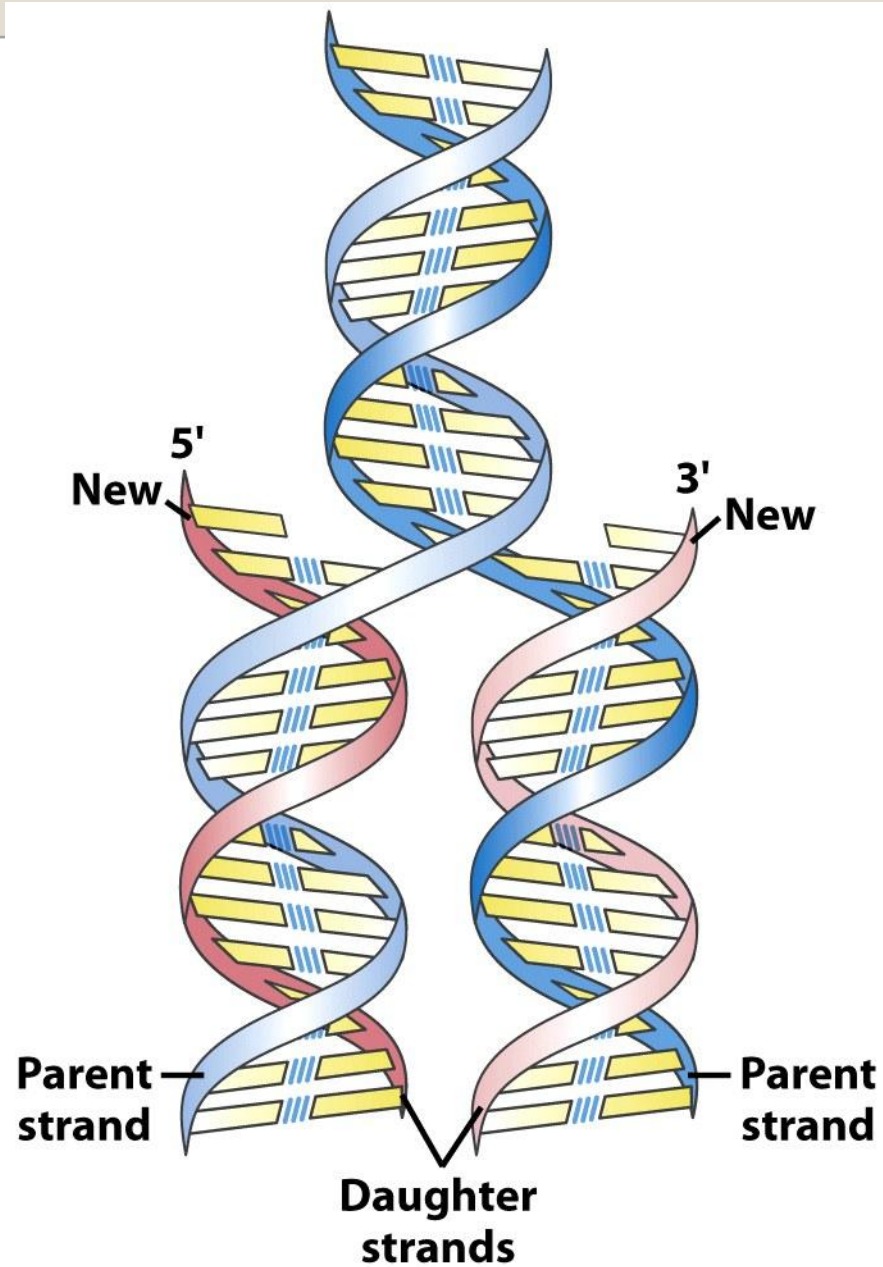
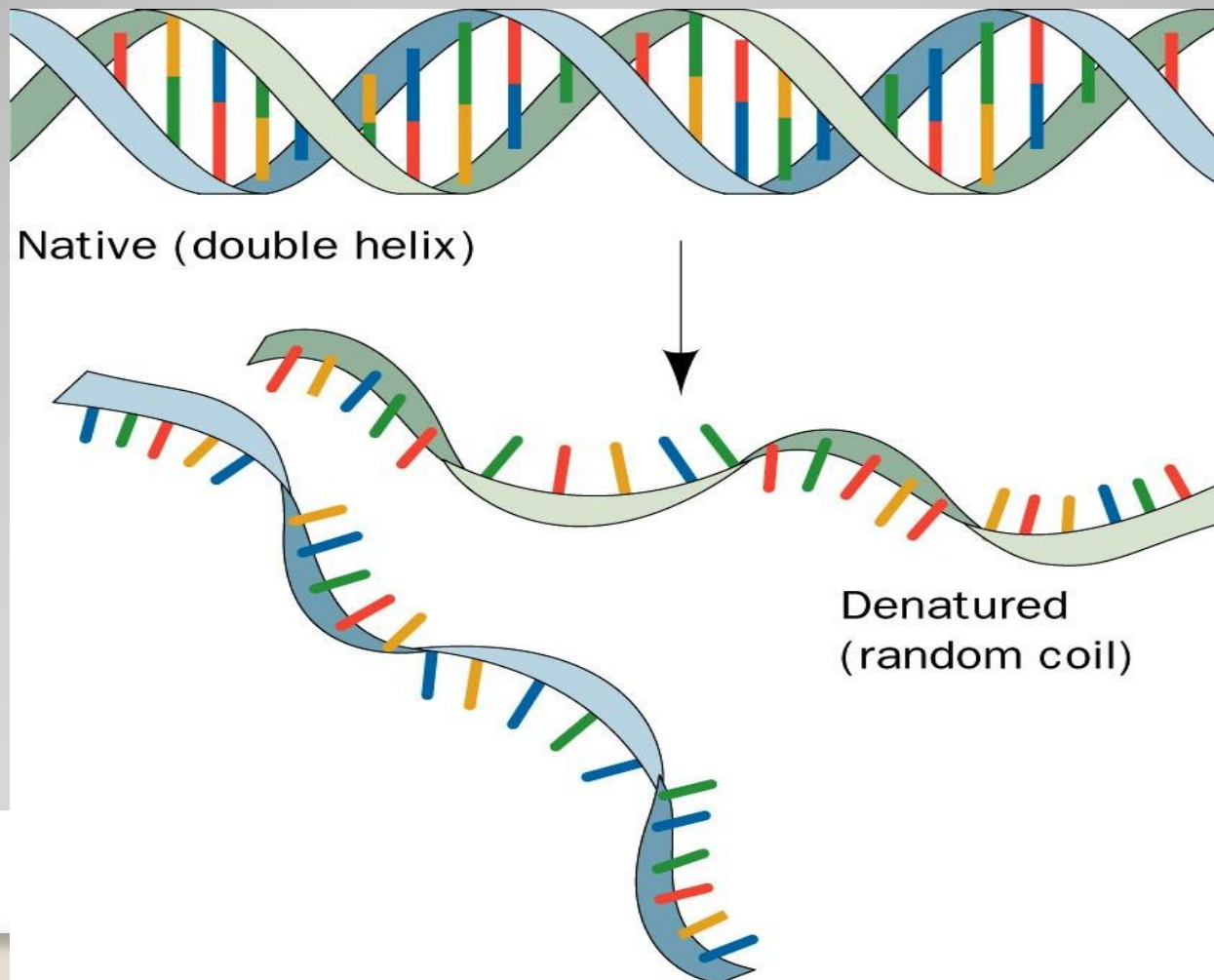


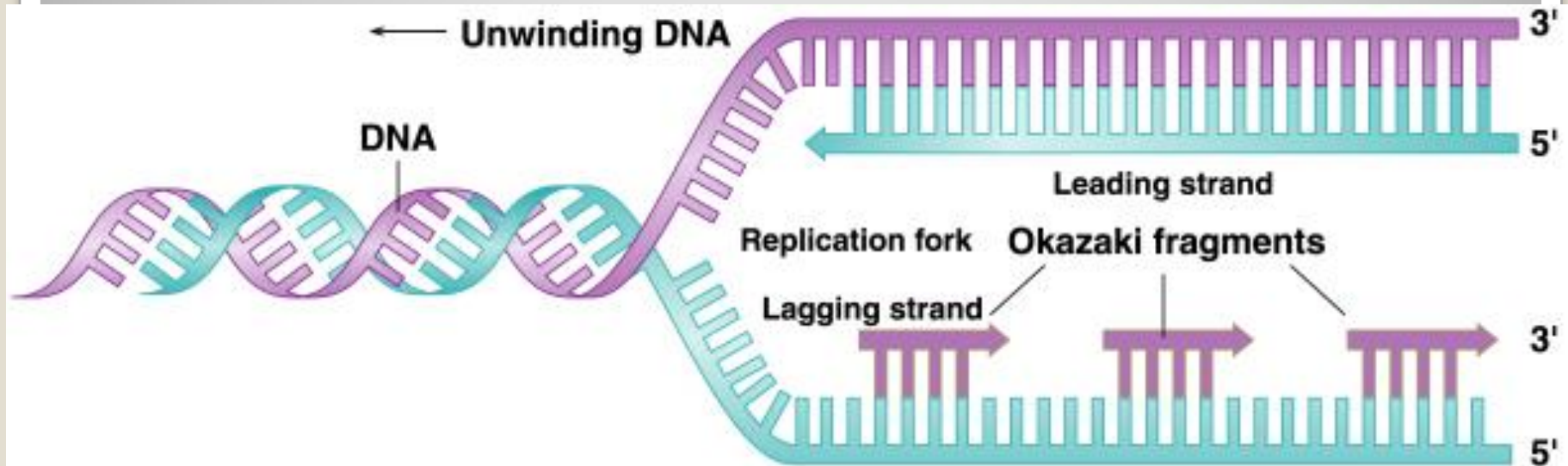
Figure 8-15
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Figure 5-14 Schematic representation of the strand separation in duplex DNA resulting from its heat denaturation.



Direction of Replication

- The enzyme *helicase* unwinds several sections of parent DNA
- At each open DNA section, called a **replication fork**, DNA *polymerase* catalyzes the formation of 5'-3' ester bonds of the **leading strand**
- The **lagging strand**, which grows in the 3'-5' direction, is synthesized in short sections called **Okazaki fragments**
- The Okazaki fragments are joined by DNA *ligase* to give a single 3'-5' DNA strand



Replication

DNA

Transcription

RNA

Translation

Protein

DNA	mRNA	Polypeptide
5' C G 3'	5' C 3'	↑ Amino terminus
G C	G }	Arg
T A	U }	
G C	G }	Gly
G C	G }	
A T	A }	
T A	U }	Tyr
A T	A }	
C G	C }	Thr
A T	A }	
C G	C }	
T A	U }	Phe
T A	U }	
T A	U }	
G C	G }	Ala
C G	C }	
C G	C }	
G C	G }	Val
T A	U }	
T A	U }	
T A	U }	
C G	C }	Ser
T A	U }	↓ Carboxyl terminus
3' T A 5'	3' U 3'	

Template strand

RNA Nucleotides

Composition (3 parts):

- 1- Ribose sugar (with O in 3rd carbon)
- 2- Phosphate group
- 3- One of 4 types of bases (all containing nitrogen):
 - Adenine
 - Uracyl (only in RNA)
 - Cytosine
 - Guanine

Ribonucleic Acid (RNA)

- RNA is much more abundant than DNA
- There are several important differences between RNA and DNA:
 - the pentose sugar in RNA is ribose, in DNA it's deoxyribose
 - in RNA, uracil replaces the base thymine (U pairs with A)
 - RNA is single stranded while DNA is double stranded
 - RNA molecules are much smaller than DNA molecules
- There are three main types of RNA:
 - ribosomal (rRNA), messenger (mRNA) and transfer (tRNA)

Types of RNA

Table 22.3 Types of RNA Molecules

Type	Abbreviation	Percentage of Total RNA	Function in the Cell
Ribosomal RNA	rRNA	75	Major component of the ribosomes
Messenger RNA	mRNA	5–10	Carries information for protein synthesis from the DNA in the nucleus to the ribosomes
Transfer RNA	tRNA	10–15	Brings amino acids to the ribosomes for protein synthesis

- Its sequence is copied from genetic DNA
- It travels to ribosomes, small granular particles in the cytoplasm of a cell where protein synthesis takes place

Messenger RNA (mRNA)

- Ribosomes are a complex of proteins and rRNA
- The synthesis of proteins from amino acids and ATP occurs in the ribosome
- The rRNA provides both structure and catalysis

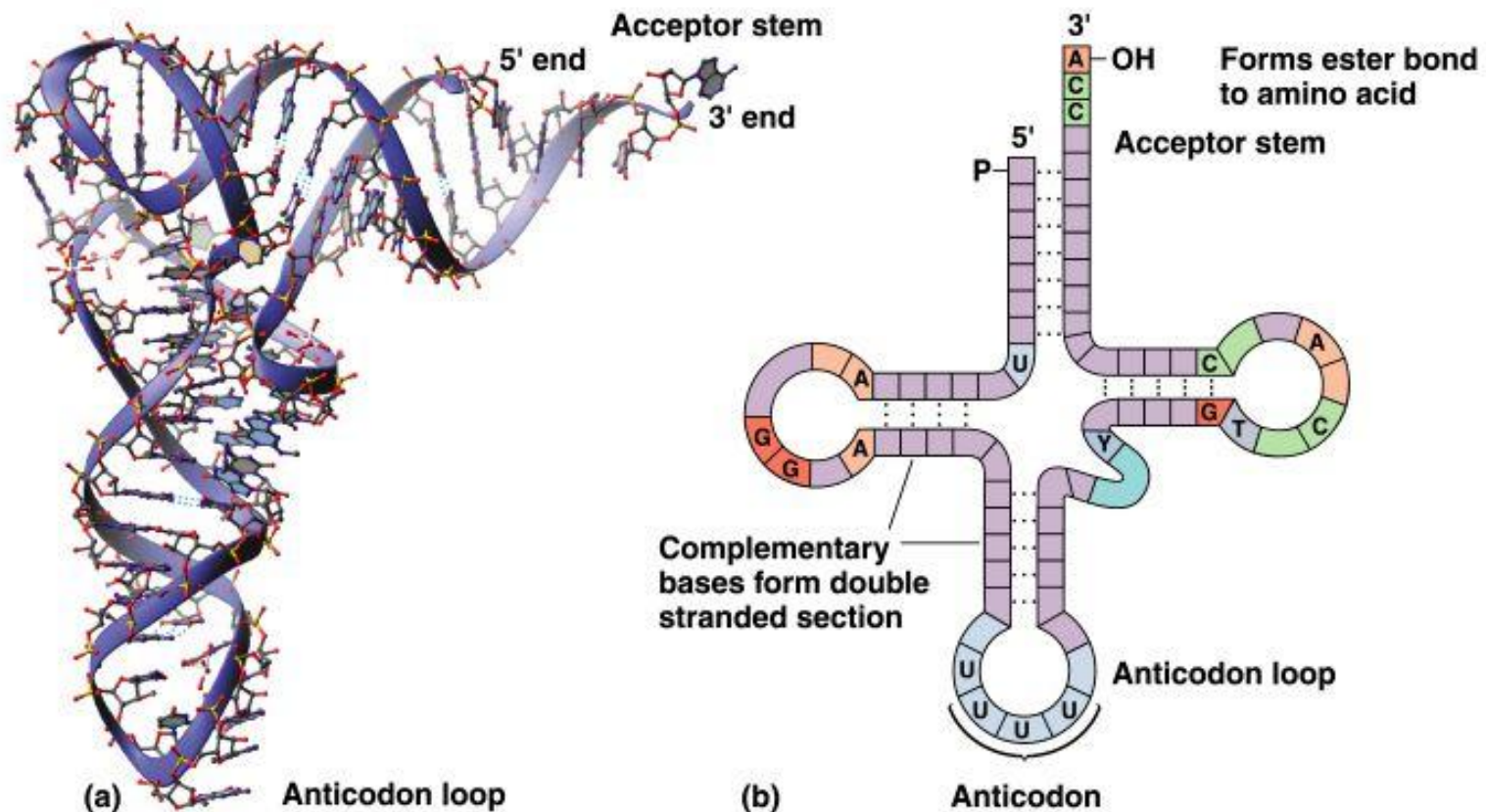
Ribosomal RNA (rRNA)

- Transports amino acids to the ribosomes where they are joined together to make proteins
- There is a specific tRNA for each amino acid
- Recognition of the tRNA at the anti-codon communicates which amino acid is attached

Transfer RNA (tRNA)

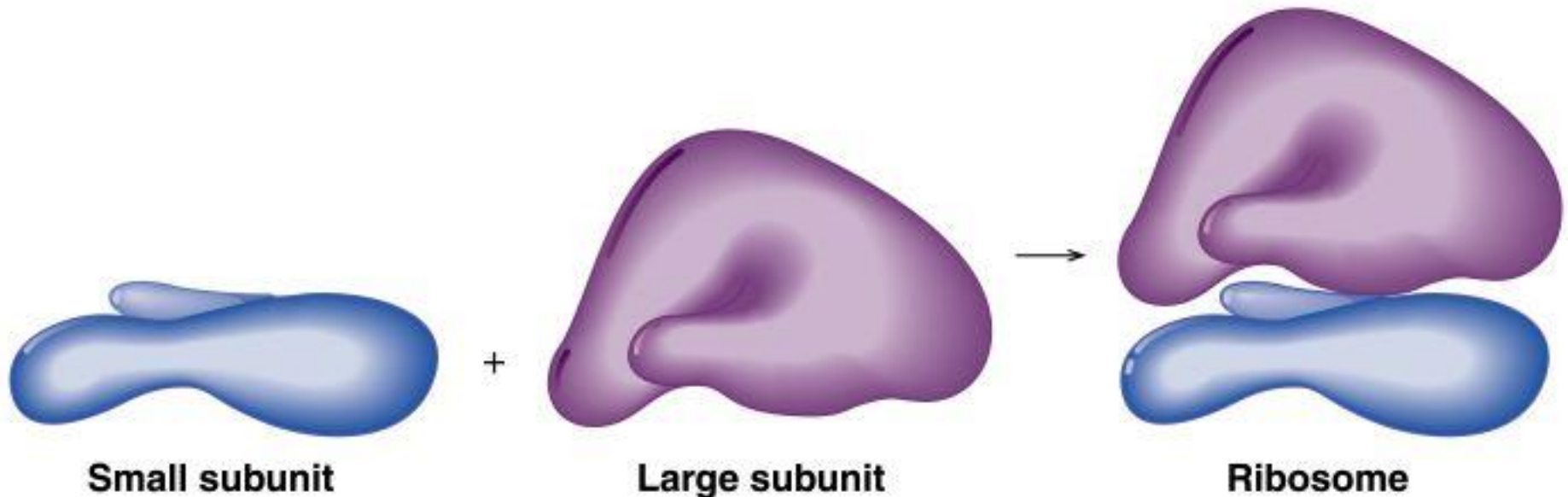
Transfer RNA

- **Transfer RNA** translates the genetic code from the messenger RNA and brings specific amino acids to the ribosome for protein synthesis
- Each amino acid is recognized by one or more specific tRNA
- tRNA has a tertiary structure that is L-shaped
 - one end attaches to the amino acid and the other binds to the mRNA by a 3-base complementary sequence



Ribosomal RNA and Messenger RNA

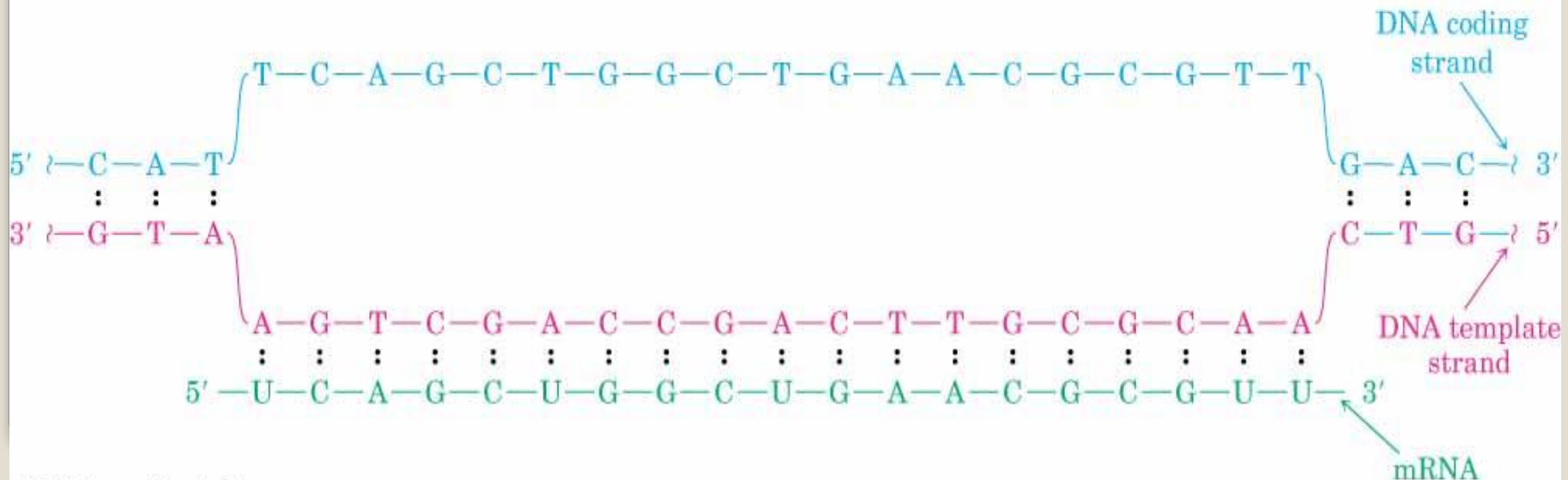
- **Ribosomes** are the sites of protein synthesis
 - they consist of **ribosomal DNA** (65%) and proteins (35%)
 - they have two subunits, a large one and a small one
- **Messenger RNA** carries the genetic code to the ribosomes
 - they are strands of RNA that are complementary to the DNA of the gene for the protein to be synthesized



How DNA Works

- 1- DNA stores genetic information in segments called **genes**
- 2- The DNA code is in **Triplet Codons** (short sequences of **3** nucleotides each)
- 3- Certain **codons** are translated by the cell into **certain Amino acids**.
4. Thus, the **sequence of nucleotides** in DNA indicate a **sequence of Amino acids** in a **protein**.

- Several turns of the DNA double helix unwind, exposing the bases of the two strands
- Ribonucleotides line up in the proper order by hydrogen bonding to their complementary bases on DNA
- Bonds form in the 5' → 3' direction,



RNA—Ribonucleic Acid

- RNA is a messenger that allows the instruction of DNA to be delivered to the rest of the cell
- RNA is different than DNA:
 1. The sugar in RNA is ribose; the sugar in DNA is deoxyribose
 2. RNA is a single strand of nucleotides; DNA is a double strand of nucleotides
 3. RNA has Uracil (U) instead of Thymine (T) which is in DNA
 4. RNA is found inside and outside of the nucleus; DNA is found only inside the nucleus

G

U

A

C

U

G

A

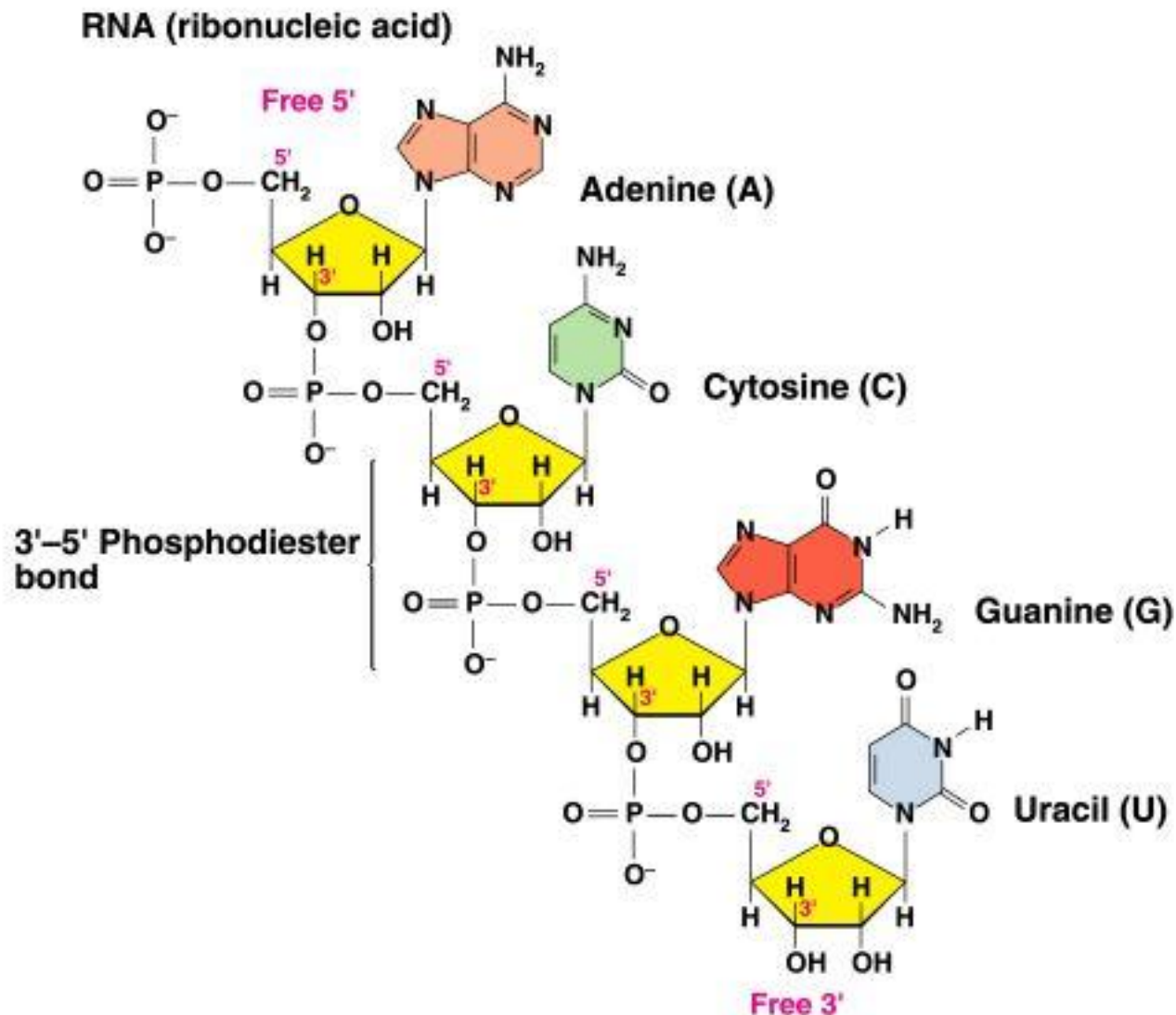
C

- Only one of the two DNA strands is transcribed into mRNA
- The strand that contains the gene is the coding or sense strand
- The strand that gets transcribed is the template or antisense strand
- The RNA molecule produced during transcription is a copy of the coding strand (with U in place of T)

Transcription of RNA from DNA

Example of RNA Primary Structure

- In RNA, A, C, G, and U are linked by 3'-5' ester bonds between ribose and phosphate

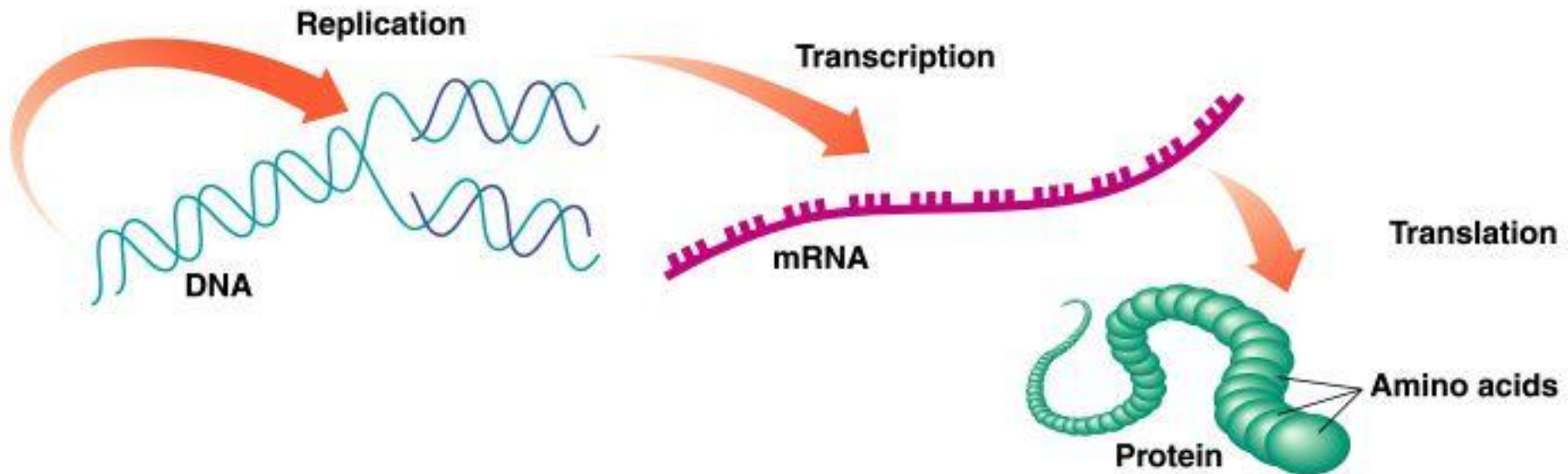


- There are 61 different tRNAs, one for each of the 61 codons that specifies an amino acid
- tRNA has 70-100 ribonucleotides and is bonded to a specific amino acid by an ester linkage through the 3' hydroxyl on ribose at the 3' end of the tRNA
- Each tRNA has a segment called an anticodon, a sequence of three ribonucleotides complementary to the codon sequence

The Parts of Transfer RNA

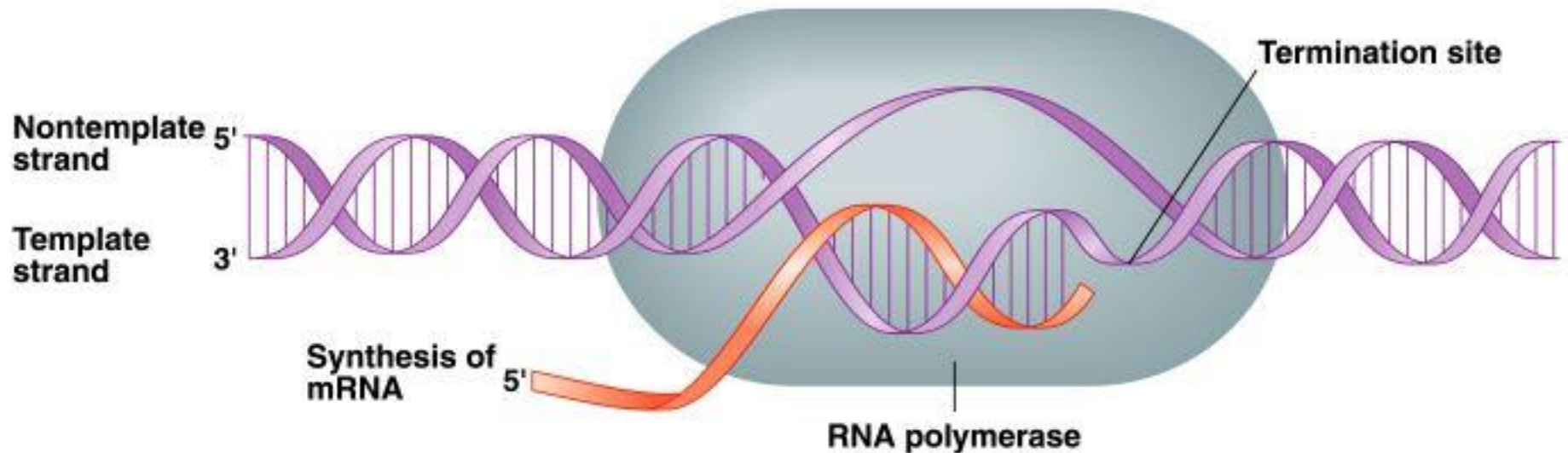
Protein Synthesis

- The two main processes involved in **protein synthesis** are
 - the formation of mRNA from DNA (**transcription**)
 - the conversion by tRNA to protein at the ribosome (**translation**)
- Transcription takes place in the nucleus, while translation takes place in the cytoplasm
- Genetic information is transcribed to form mRNA much the same way it is replicated during cell division



RNA Polymerase

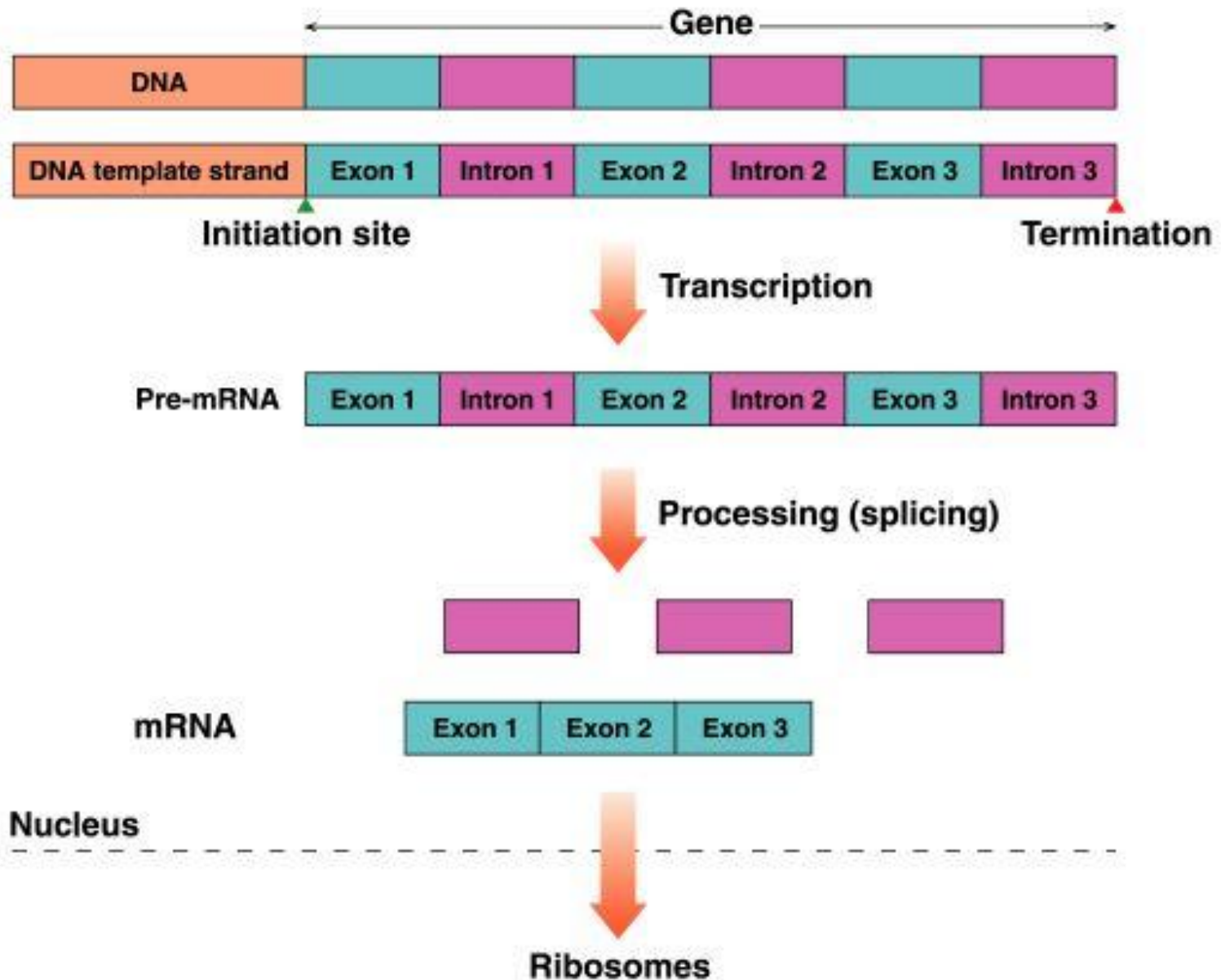
- During transcription, RNA *polymerase* moves along the DNA template in the 3'-5'direction to synthesize the corresponding mRNA
- The mRNA is released at the termination point



Processing of mRNA

- **Genes** in the DNA of eukaryotes contain **exons** that code for proteins along with **introns** that do not
- Because the initial mRNA, called a **pre-RNA**, includes the noncoding introns, it must be processed before it can be read by the tRNA
- While the mRNA is still in the nucleus, the introns are removed from the pre-RNA
- The exons that remain are joined to form the mRNA that leaves the nucleus with the information for the synthesis of protein

Removing Introns from mRNA



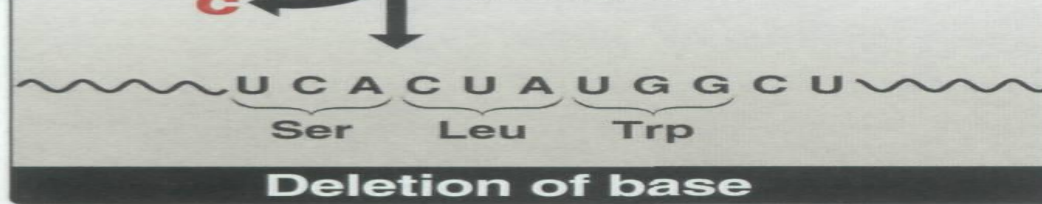


Figure 31.5

Frame-shift mutations as a result of addition or deletion of a base can cause an alteration in the reading frame of mRNA.

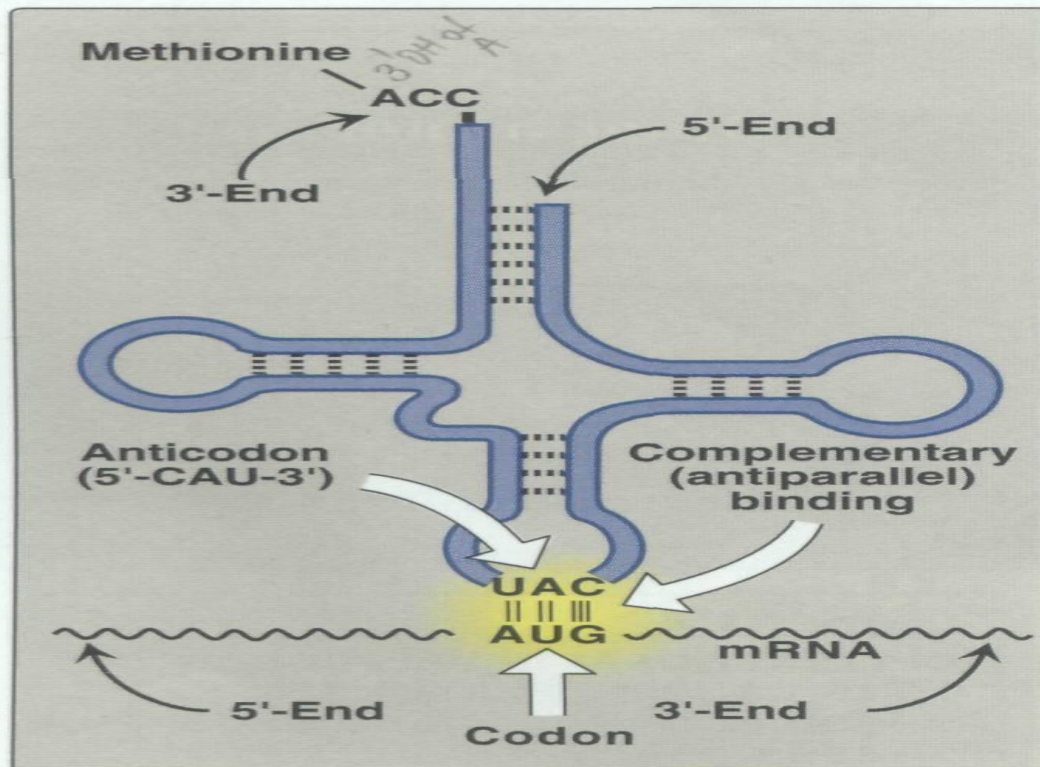
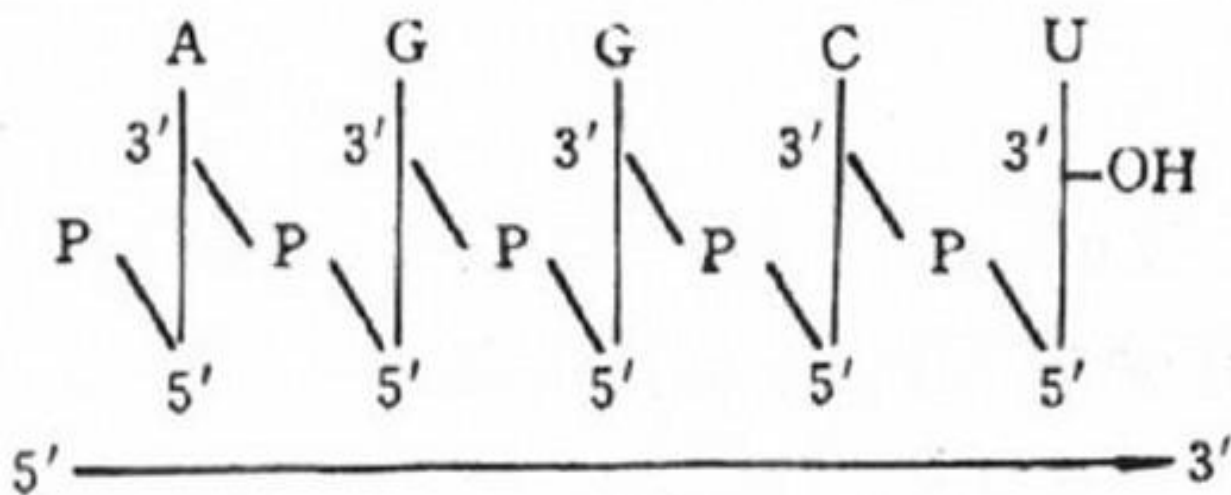


Figure 31.6

Complementary, antiparallel binding of the anticodon for methionyl-tRNA (CAU) to the mRNA codon for methionine (AUG).



5' pApGpGpCpU 3'

A → G → G → C → U

5' AGGCU 3'

图 3-8 核酸简式

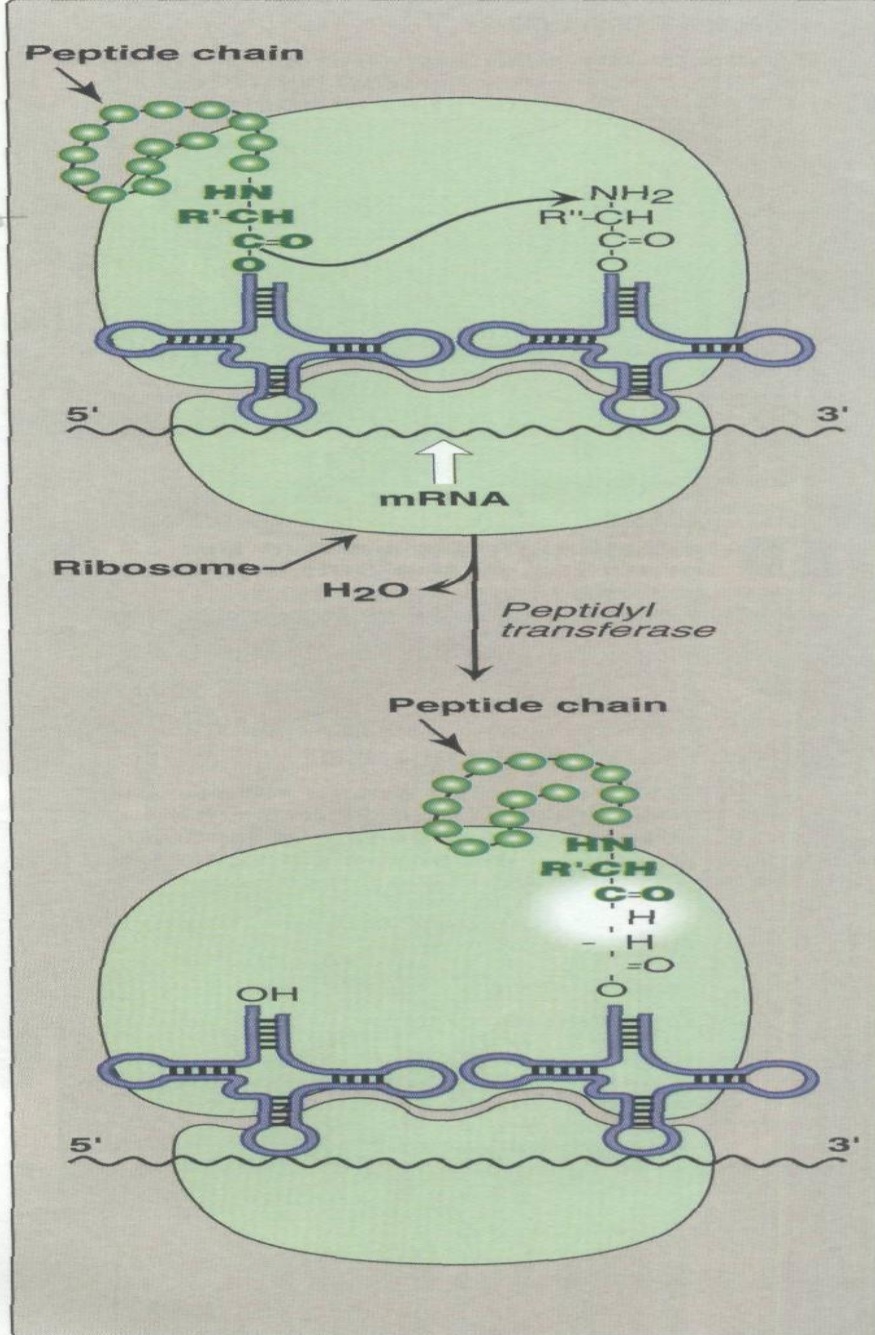


Figure 31.12
Formation of a peptide bond.

Transcription

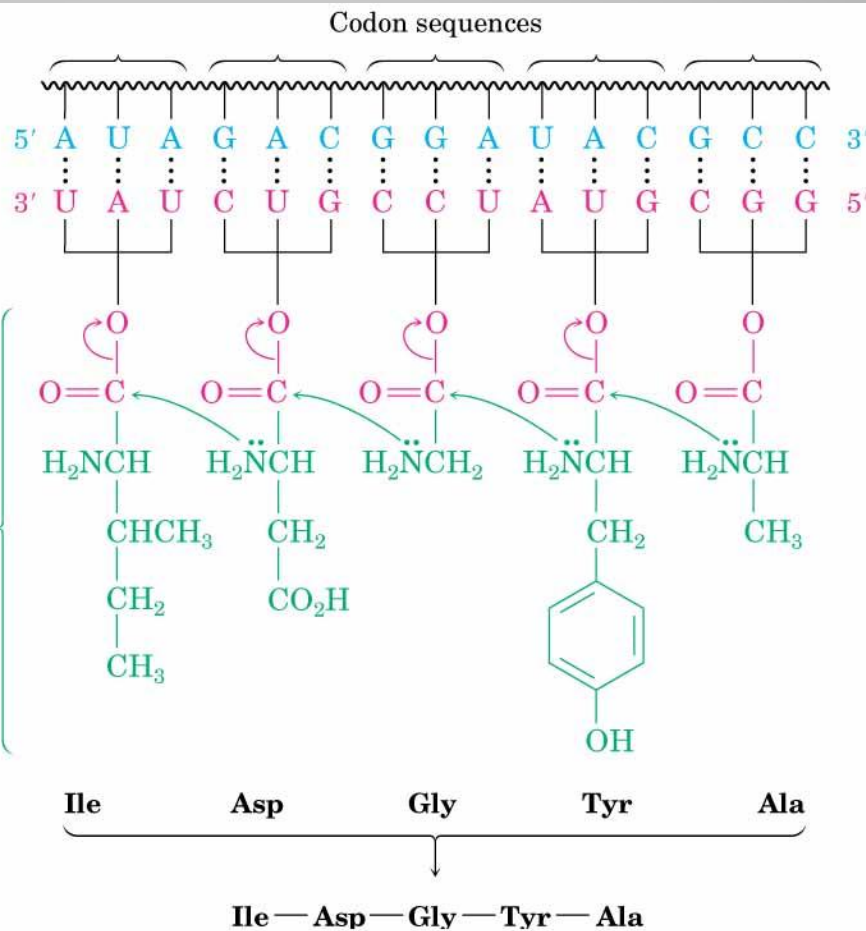
- Several steps occur during **transcription**:
 - a section of DNA containing the gene unwinds
 - one strand of DNA is copied starting at the initiation point, which has the sequence TATAAA
 - an mRNA is synthesized using complementary base pairing with uracil (U) replacing thymine (T)
 - the newly formed mRNA moves out of the nucleus to ribosomes in the cytoplasm and the DNA re-winds

mRNA chain

Codon on mRNA chain

Anticodon on tRNA

Bound amino acid residue



The

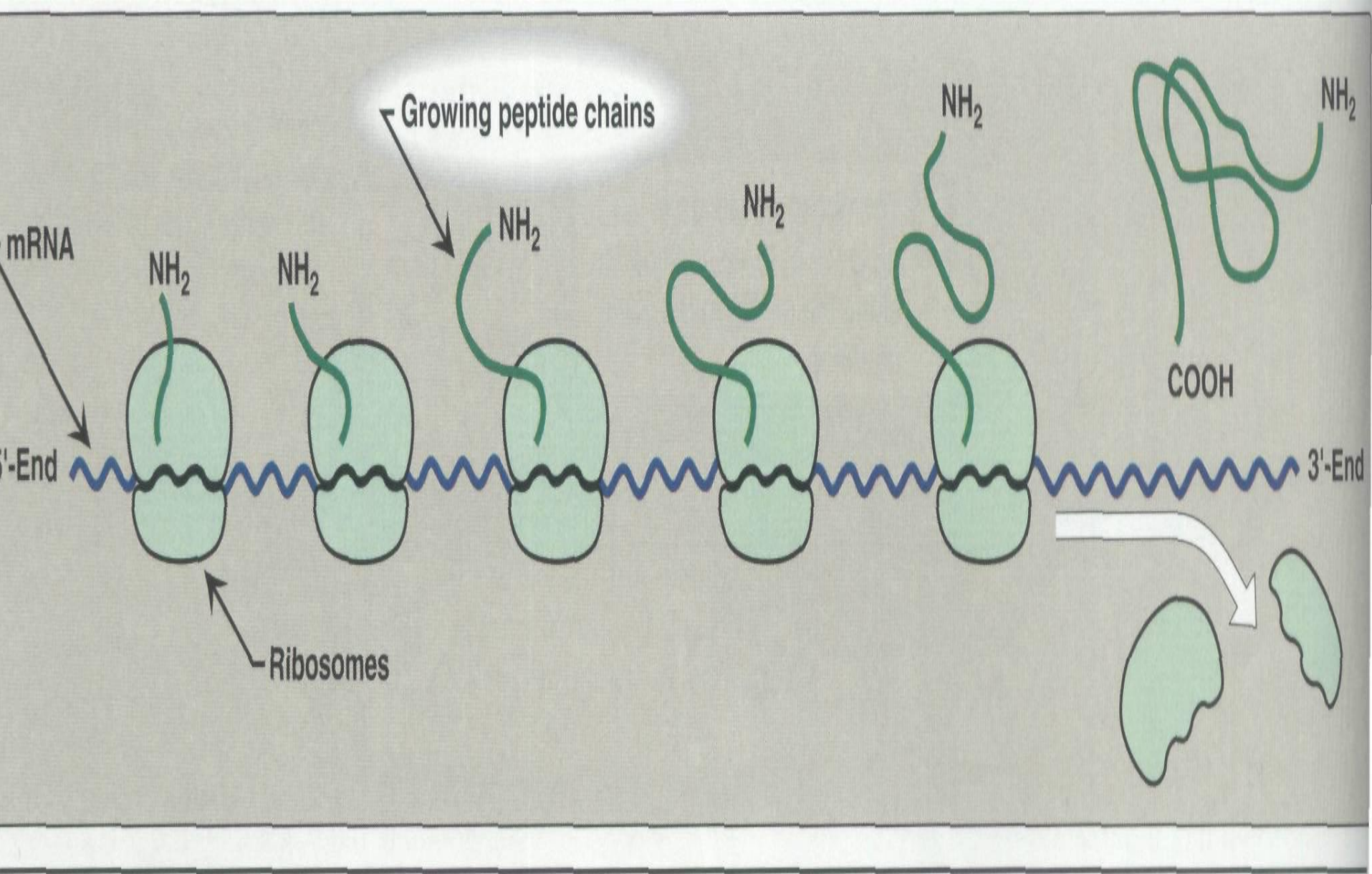
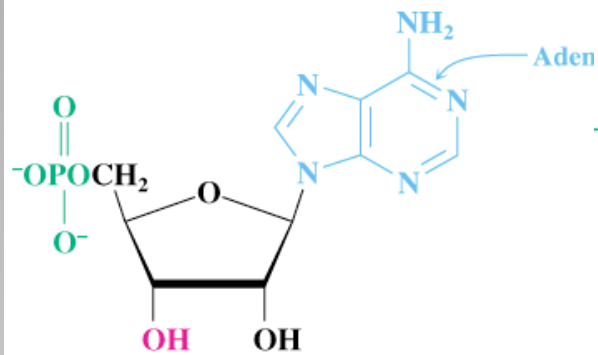


Figure 31.14

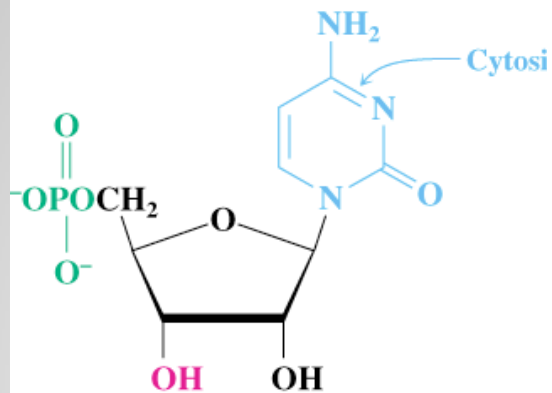
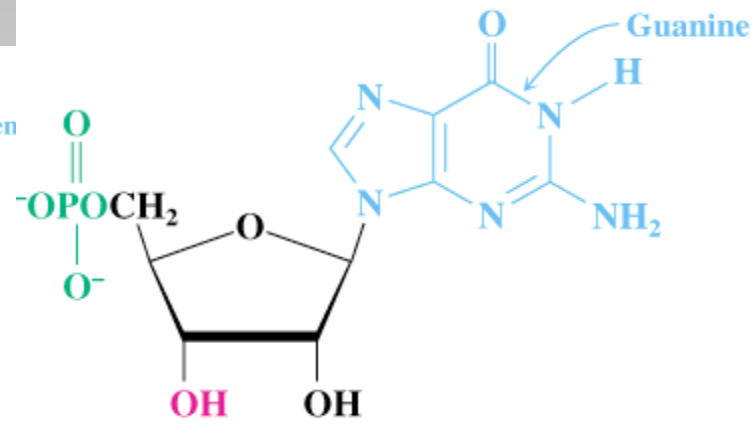
A polyribosome consists of several ribosomes simultaneously translating one mRNA.

Regulation of Transcription

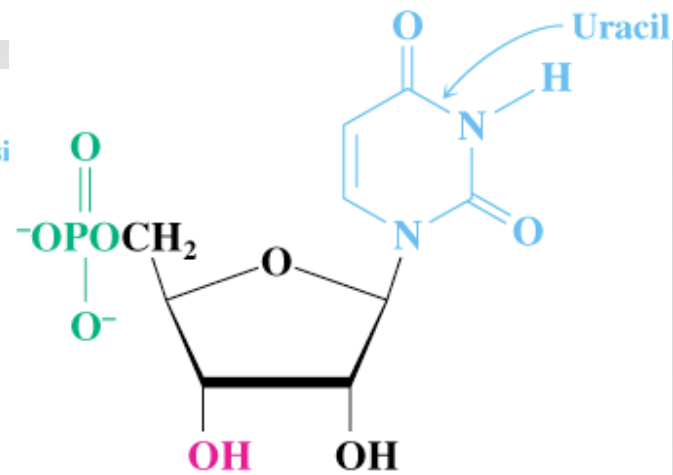
- A specific mRNA is synthesized when the cell requires a particular protein
- The synthesis is regulated at the transcription level:
 - **feedback control**, where the end products speed up or slow the synthesis of mRNA
 - **enzyme induction**, where a high level of a reactant induces the transcription process to provide the necessary enzymes for that reactant
- Regulation of transcription in eukaryotes is complicated and we will not study it here



Adenosine 5'-phosphate



Cytidine 5'-phosphate



Uridine 5'-phosphate

The

The Genetic Code

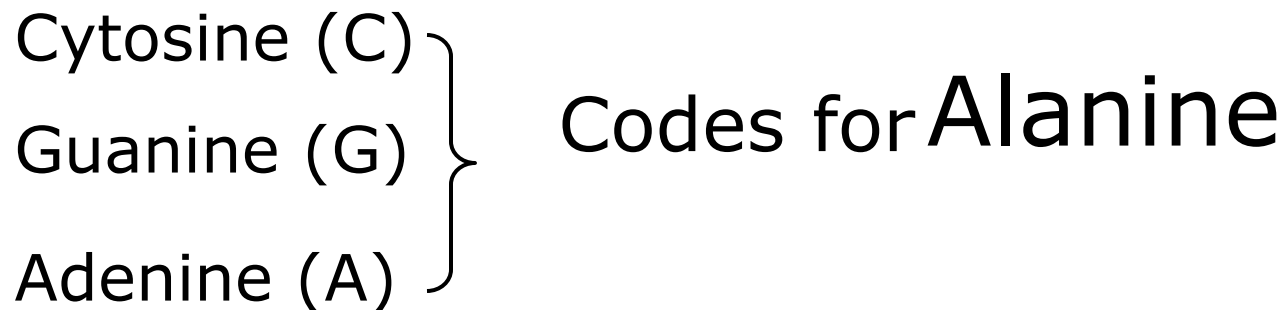
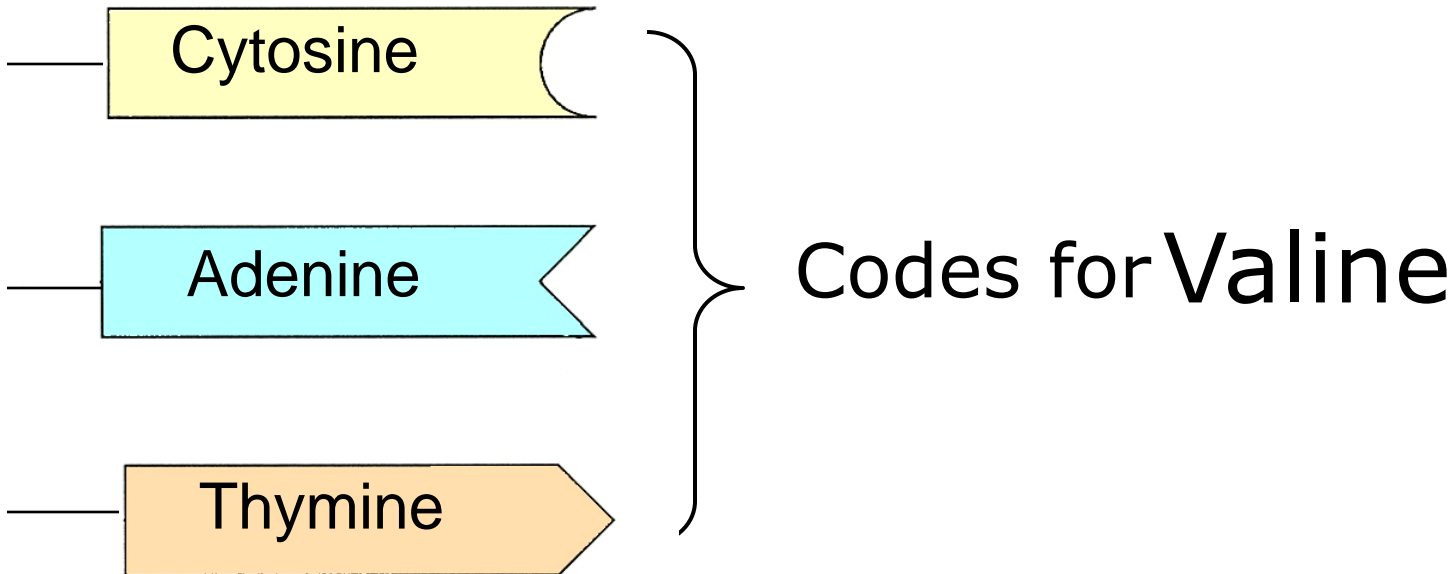
- The **genetic code** is found in the sequence of nucleotides in mRNA that is translated from the DNA
- A **codon** is a **triplet** of bases along the mRNA that codes for a particular amino acid
- Each of the 20 amino acids needed to build a protein has at least 2 codons
- There are also codons that signal the “start” and “end” of a polypeptide chain
- The amino acid sequence of a protein can be determined by reading the triplets in the DNA sequence that are complementary to the codons of the mRNA, or directly from the mRNA sequence
- The entire DNA sequence of several organisms, including humans, have been determined, however,
 - only primary structure can be determined this way
 - doesn't give tertiary structure or protein function

he sequence of bases in DNA forms the Genetic Code

**A group of three bases (a
triplet) controls
the production of a particular
amino acid in
the cytoplasm of the cell**

**The different amino acids and the
order in which they are joined up
determines the sort of protein
being produced**

For example



This is known as the **triplet code**

Each triplet codes for a specific amino acid

CGA	-	CAA	-	CCA	-	CCA	-	GCT	-	GGG	-	GAG	-	CCA	-
↓		↓		↓		↓		↓		↓		↓			
Ala		Val		Gly		Gly		Arg		Pro		Leu		Gly	

The amino acids are joined together in the sequence to make part of a protein

—Ala—Val— Gly— Gly—Arg— Pro— Leu—Gly—

mRNA Codons and Associated Amino Acids

Second Letter					
First Letter	U	C	A	G	Third Letter
U	UUU } Phe	UCU } Ser	UAU } Tyr	UGU } Cys	U
	UUC } Phe	UCC } Ser	UAC } Tyr	UGC } Cys	C
	UUA } Leu	UCA } Ser	UAA STOP	UGA STOP	A
	UUG } Leu	UCG } Ser	UAG STOP	UGG Trp	G
C	CUU } Leu	CCU } Pro	CAU } His	CGU } Arg	U
	CUC } Leu	CCC } Pro	CAC } His	CGC } Arg	C
	CUA } Leu	CCA } Pro	CAA } Gln	CGA } Arg	A
	CUG } Leu	CCG } Pro	CAG } Gln	CGG } Arg	G
A	AUU } Ile	ACU } Thr	AAU } Asn	AGU } Ser	U
	AUC } Ile	ACC } Thr	AAC } Asn	AGC } Ser	C
	AUA } Ile	ACA } Thr	AAA } Lys	AGA } Arg	A
	^a AUG Met/start	ACG } Thr	AAG } Lys	AGG } Arg	G
G	GUU } Val	GCU } Ala	GAU } Asp	GGU } Gly	U
	GUC } Val	GCC } Ala	GAC } Asp	GGC } Gly	C
	GUA } Val	GCA } Ala	GAA } Glu	GGA } Gly	A
	GUG } Val	GCG } Ala	GAG } Glu	GGG } Gly	G

^a Codon that signals the start of a peptide chain.

STOP codons signal the end of a peptide chain.

Reading the Genetic Code

- Suppose we want to determine the amino acids coded for in the following section of a mRNA

5'—CCU —AGC—GGA—CUU—3'

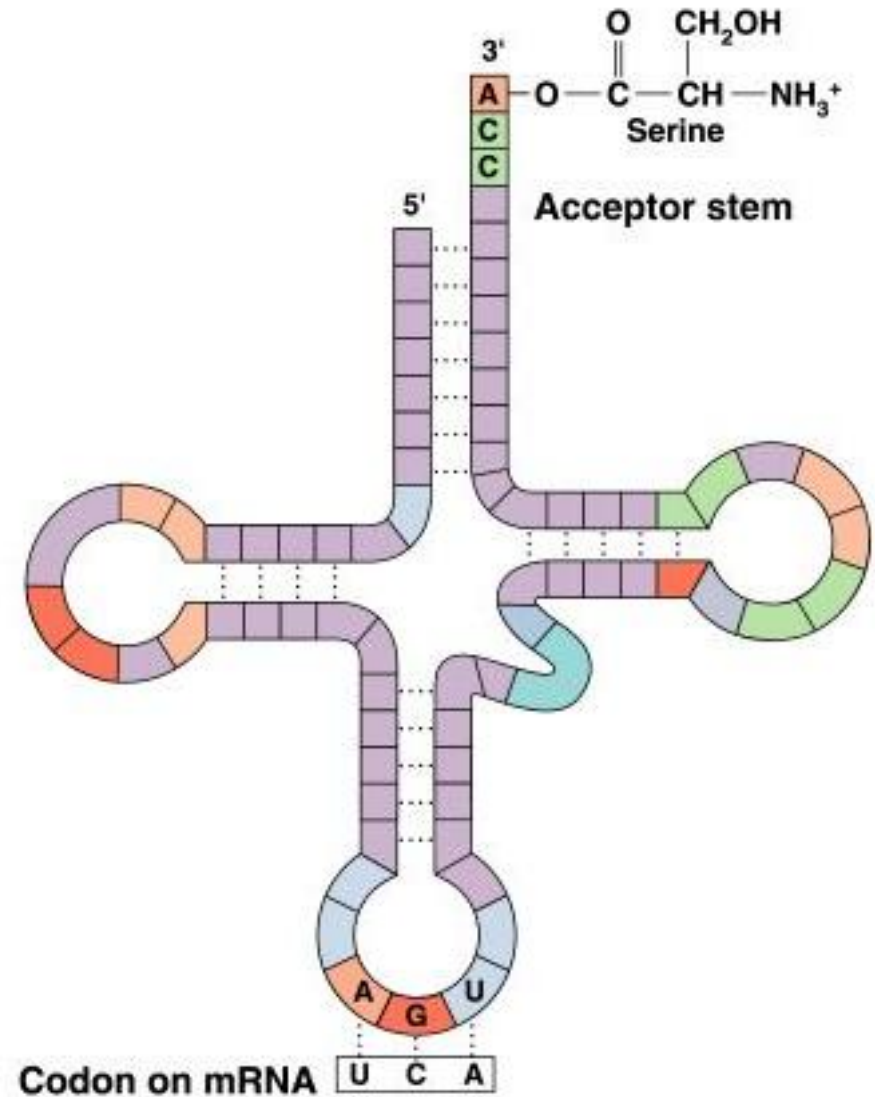
- According to the genetic code, the amino acids for these codons are:

CCU = Proline	AGC = Serine
GGA = Glycine	CUU = Leucine

- The mRNA section codes for the amino acid sequence of Pro—Ser—Gly—Leu

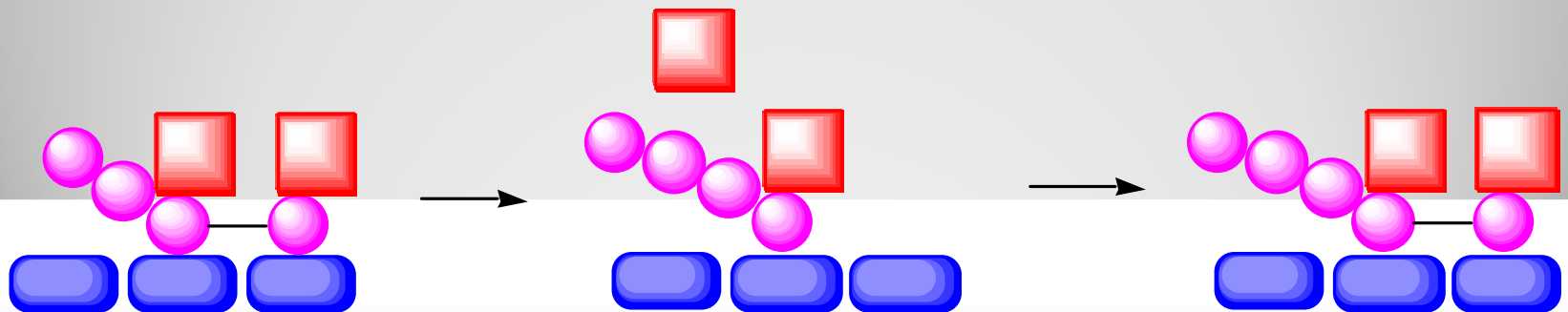
Translation and tRNA Activation

- Once the DNA has been transcribed to mRNA, the codons must be translated to the amino acid sequence of the protein
- The first step in **translation** is activation of the tRNA
- Each tRNA has a triplet called an **anticodon** that complements a codon on mRNA
- A *synthetase* uses ATP hydrolysis to attach an amino acid to a specific tRNA



Initiation and Translocation

- Initiation of protein synthesis occurs when a mRNA attaches to a ribosome
- On the mRNA, the **start codon (AUG)** binds to a tRNA with methionine
- The second codon attaches to a tRNA with the next amino acid
- A peptide bond forms between the adjacent amino acids at the first and second codons
- The first tRNA detaches from the ribosome and the ribosome shifts to the adjacent codon on the mRNA (this process is called **translocation**)
- A third codon can now attach where the second one was before translocation



Termination

- After a polypeptide with all the amino acids for a protein is synthesized, the ribosome reaches the the "**stop**" **codon**: UGA, UAA, or UAG
- There is no tRNA with an anticodon for the "stop" codons
- Therefore, protein synthesis ends (**termination**)
- The polypeptide is released from the ribosome and the protein can take on it's 3-D structure
(some proteins begin folding while still being synthesized, while others do not fold up until after being released from the ribosome)

The proteins build the cell structures

They also make enzymes

The DNA controls which enzymes are made
the enzymes determine what reactions take place

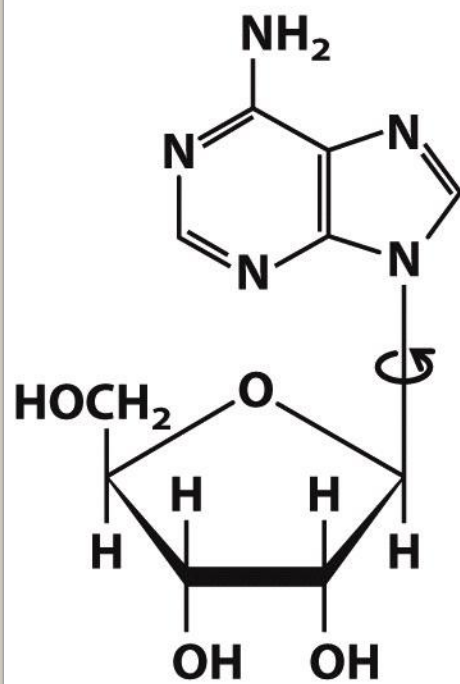
The structures and reactions in the cell determine
what sort of a cell it is and what its functions are

So DNA exerts its control through the enzymes

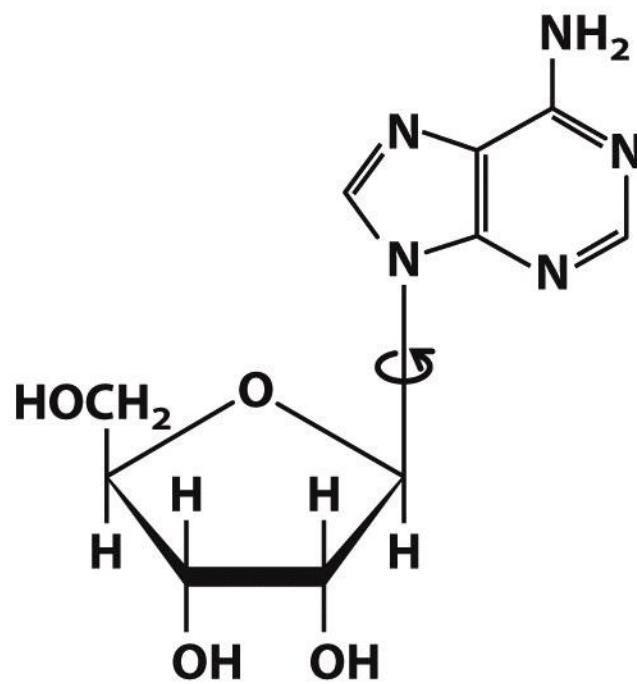
A sequence of triplets in the DNA molecule code for a complete protein

Such a sequence forms a **gene**

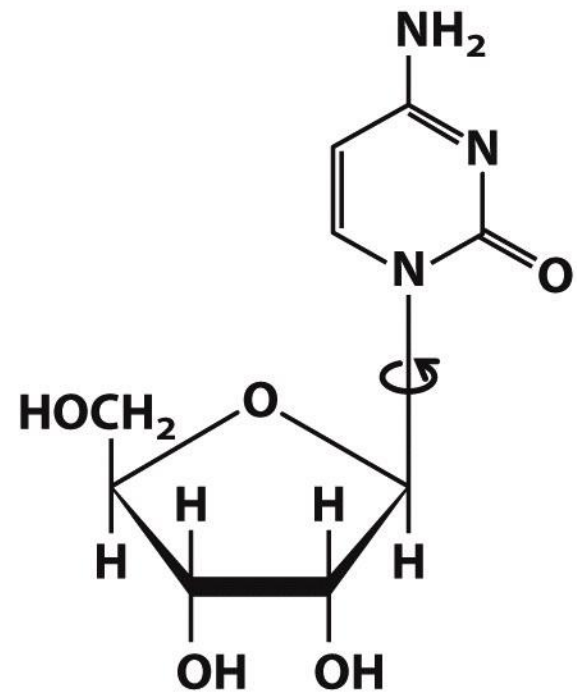
There may be a thousand or more base pairs in one gene



***syn*-Adenosine**



***anti*-Adenosine**



***anti*-Cytidine**

Figure 8-16b

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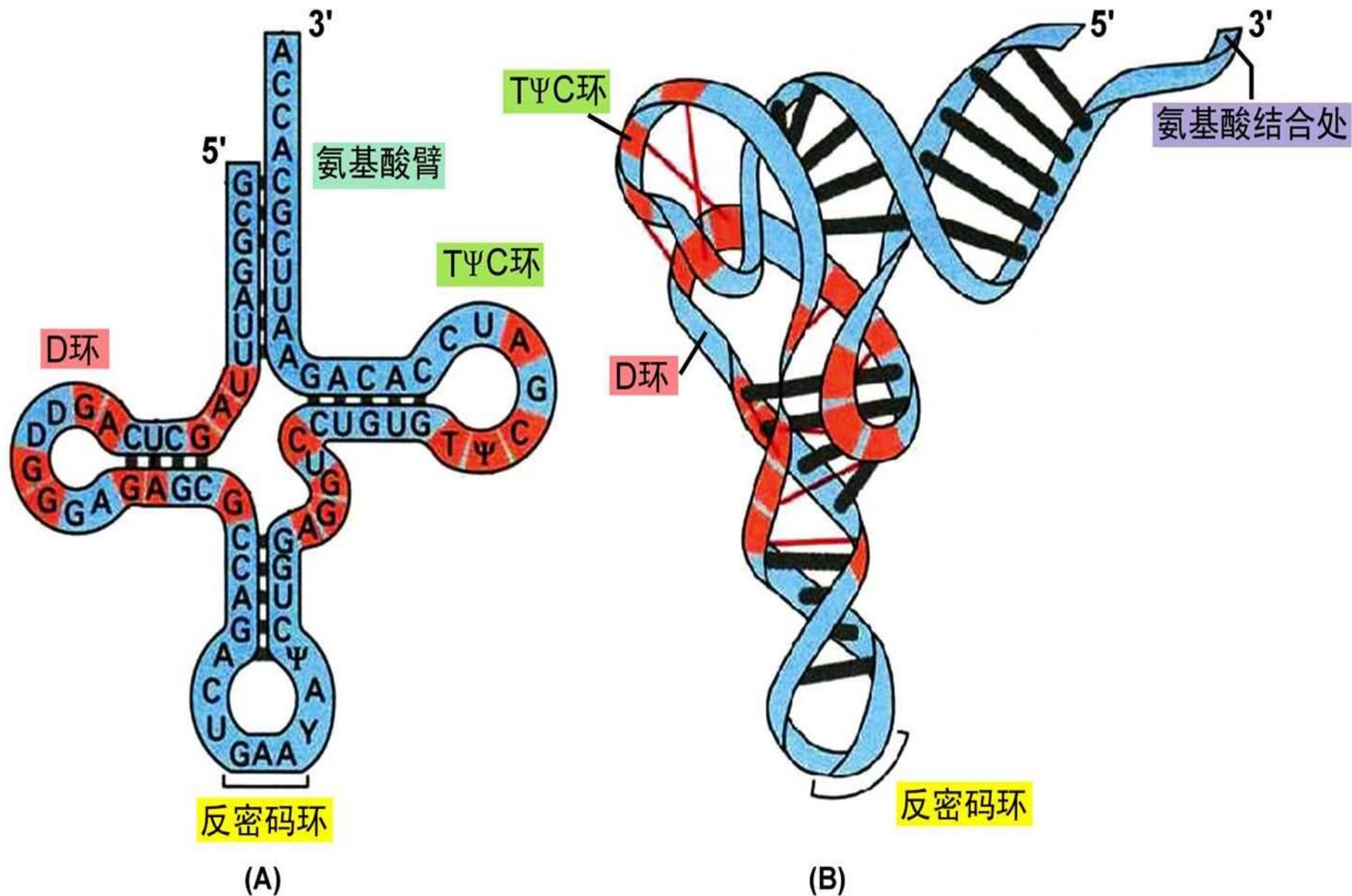
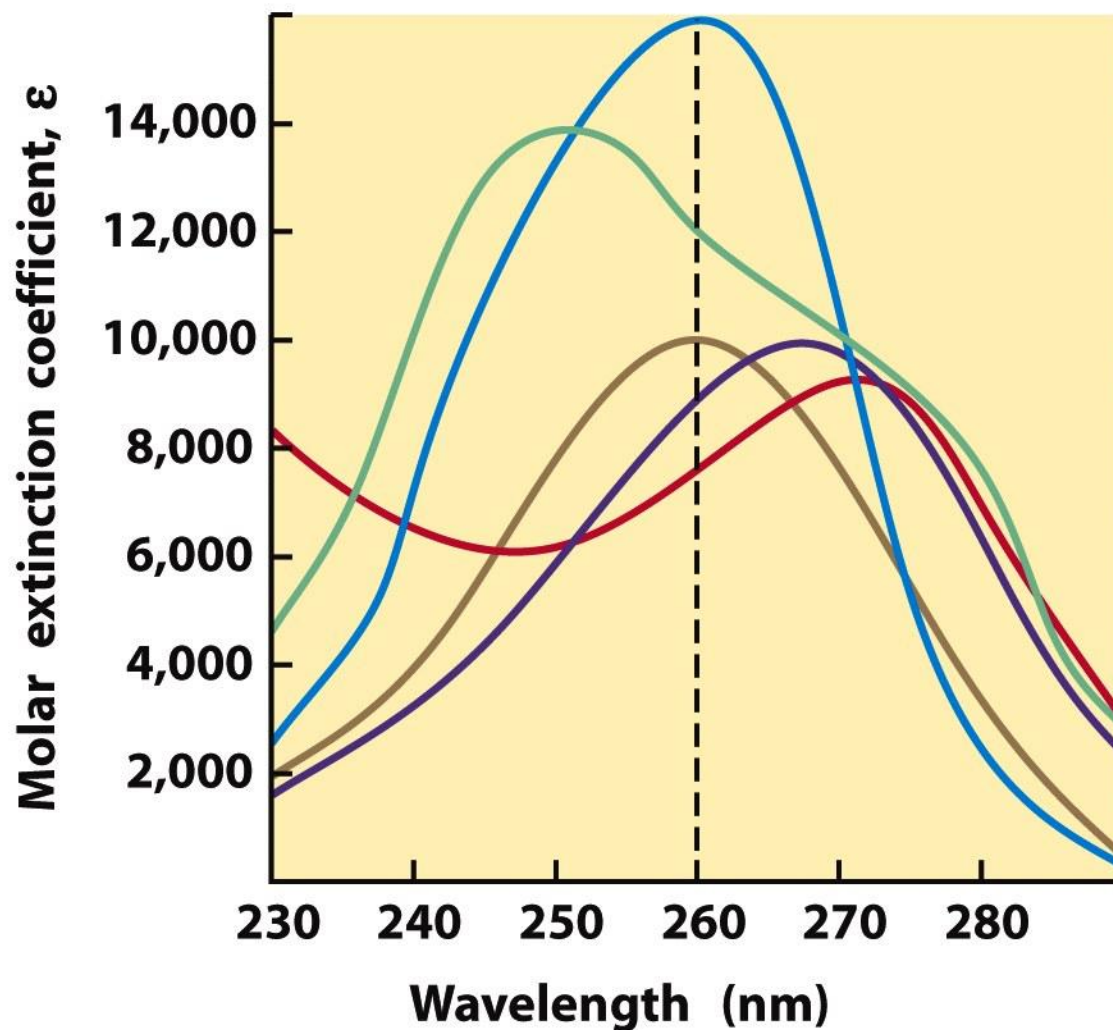


图3-16 tRNA的二级(A)及三级结构(B)

Conformation around N-glycosidic Bond

- Relatively **free rotation** can occur around the N-glycosidic bond in free nucleotides
- The torsion angle about the N-glycosidic bond (N-C1') is denoted by the symbol χ
- The sequence of atoms chosen to define this angle is O4'-C1'-N9-C4 for purine, and O4'-C1'-N1-C2 for pyrimidine derivatives
- Angle near 0° corresponds to ***syn conformation***
- Angle near 180° corresponds to ***anti conformation***
- Anti conformation is found in normal B-DNA



Molar extinction coefficient at 260 nm, ϵ_{260} ($M^{-1}cm^{-1}$)

AMP	15,400
GMP	11,700
UMP	9,900
dTMP	9,200
CMP	7,500

Figure 8-10

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Replication of Genetic Code

- Strand separation occurs first
- Each strand serves as a template for the synthesis of a new strand
- Synthesis is catalyzed by enzymes known as DNA polymerases
- Newly made DNA molecule has one daughter strand and one parent strand.

“It has not escaped our notice that the specific pairing we have postulated immediately suggests a possible copying mechanism for the genetic material”

Watson and Crick, in their Nature paper, 1953

Message RNA

Code Carrier for the Sequence of Proteins

- Is synthesized using DNA template
- Contains ribose instead of deoxyribose
- Contains uracil instead of thymine
- One mRNA may code for more than one protein



(a) Monocistronic



(b) Polycistronic

Figure 8-21

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Factors Affecting DNA Denaturation

- The midpoint of melting (T_m) depends on base composition
 - high CG increases T_m
- T_m depends on DNA length
 - Longer DNA has higher T_m
 - Important for short DNA
- T_m depends on pH and ionic strength
 - High salt increases T_m