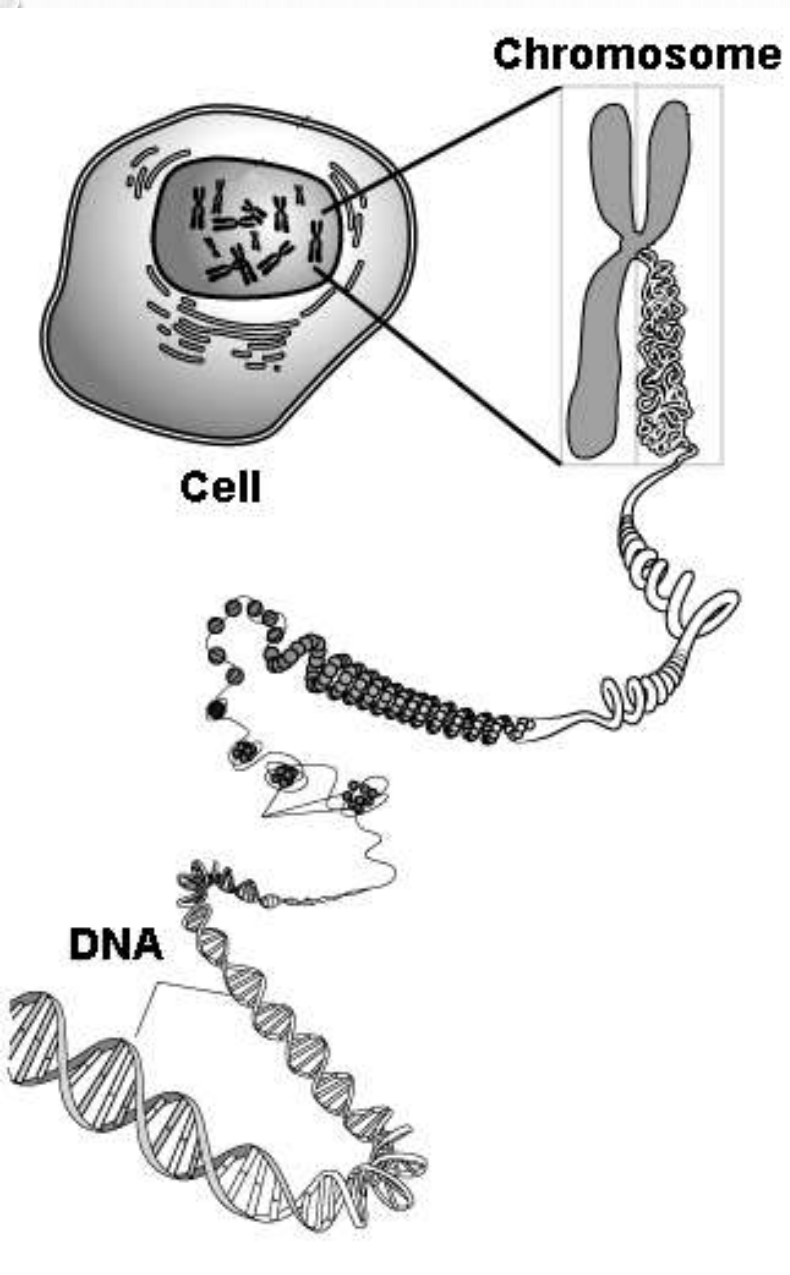


DEOXYRIBONUCLEIC ACID



Nucleus



Chromosomes



Genes



Segments of
DNA

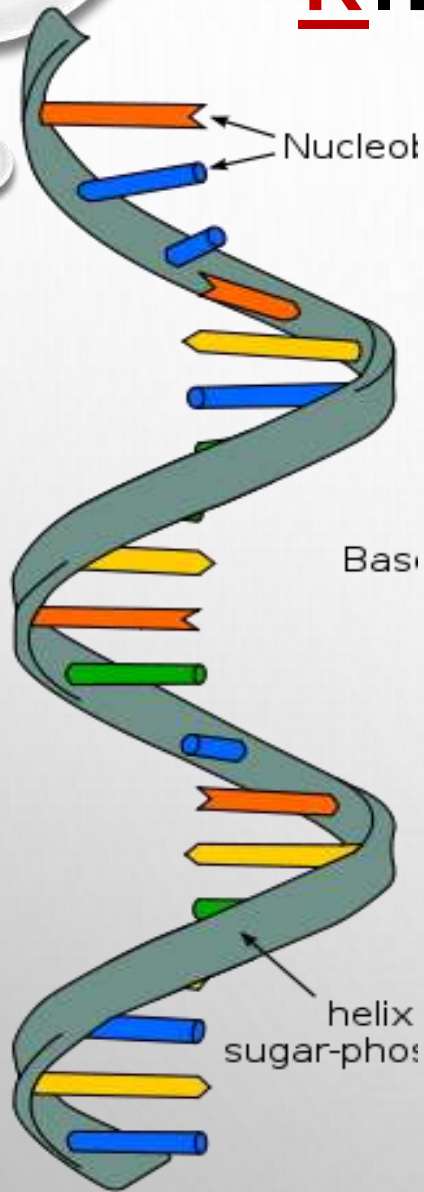
Portions of DNA are called genes.

DNA is tightly wound into chromosomes and located in the nucleus of cells.

DNA cannot leave the nucleus.

DNA is **DOUBLE STRANDED**(2 sides)

RIBONUCLEIC ACID



RNA

Ribonucleic acid

RNA is **SINGLE STRANDED** and does not have to stay in the nucleus!

RNA is not found in chromosomes because it does not carry the genetic code, however it can read the **DNA code and take the information out of the nucleus.**

RNA's main job is to build proteins!

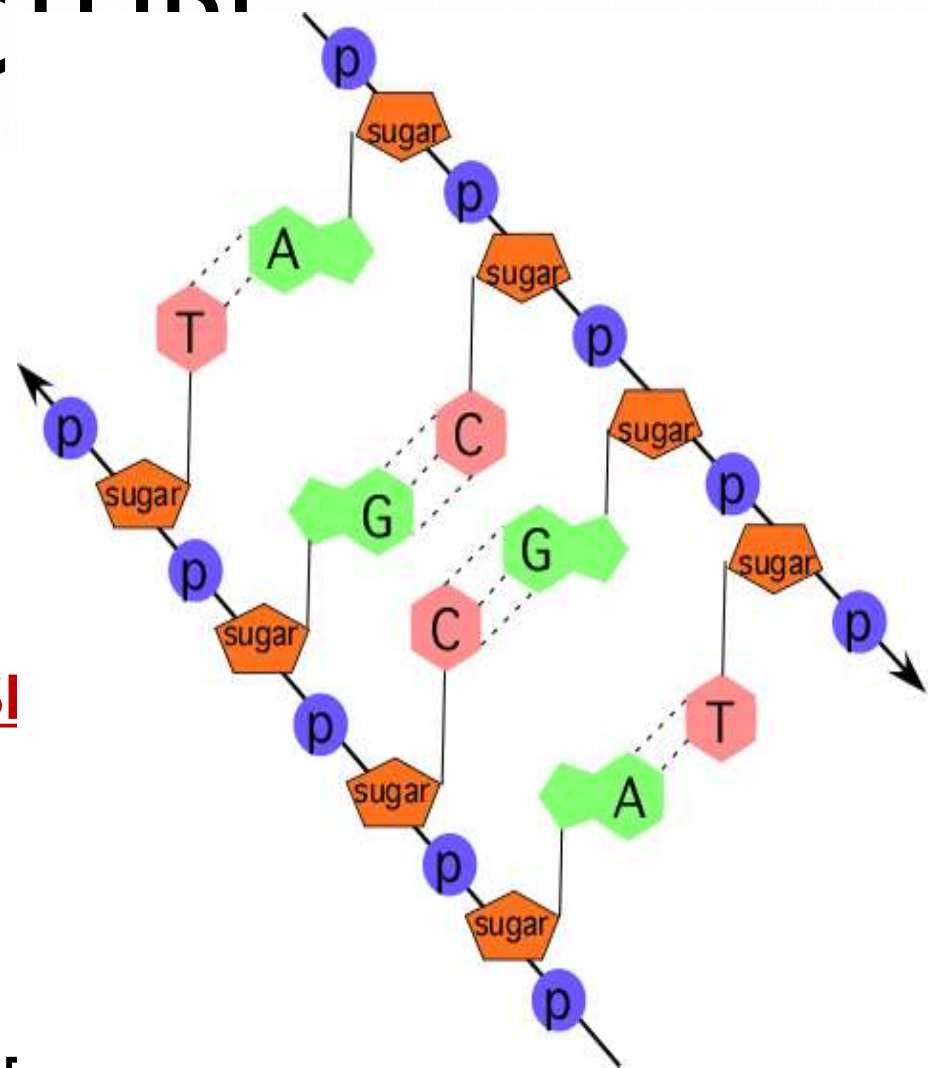
DNA STRUCTURE

○ THE BUILDING BLOCKS OF DNA ARE CALLED NUCLEOTIDES.

○ ONE NUCLEOTIDE IS MADE OF 3 IMPORTANT THINGS:

- 1. 5-CARBON SUGAR DEOXYRIBOSE
- 2. PHOSPHATE
- 3. NITROGEN BASE
- THERE ARE 4 NITROGEN BASES IN DNA: ADENINE, GUANINE, CYTOSINE, AND THYMINE THAT PAIR TOGETHER)

• A → T C → G



RNA STRUCTURE

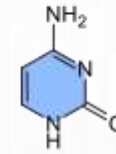
○ THE BUILDING BLOCKS OF RNA ARE NUCLEOTIDES, JUST LIKE DNA.

○ A NUCLEOTIDE IN RNA IS STILL MADE OF 3 IMPORTANT THINGS:

- 1. 6-CARBON SUGAR - RIBOSE (INSTEAD OF DEOXYRIBOSE)
- 2. PHOSPHATE
- 3. NITROGEN BASE
- THERE ARE 4 NITROGEN BASES IN RNA, A, G, C, AND U THAT PAIR TOGETHER)

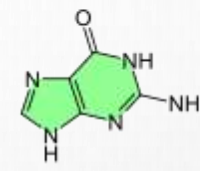
- $A \rightarrow \underline{U}$ $C \rightarrow G$

Cytosine



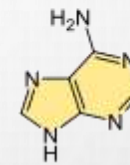
C

Guanine



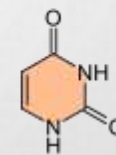
G

Adenine



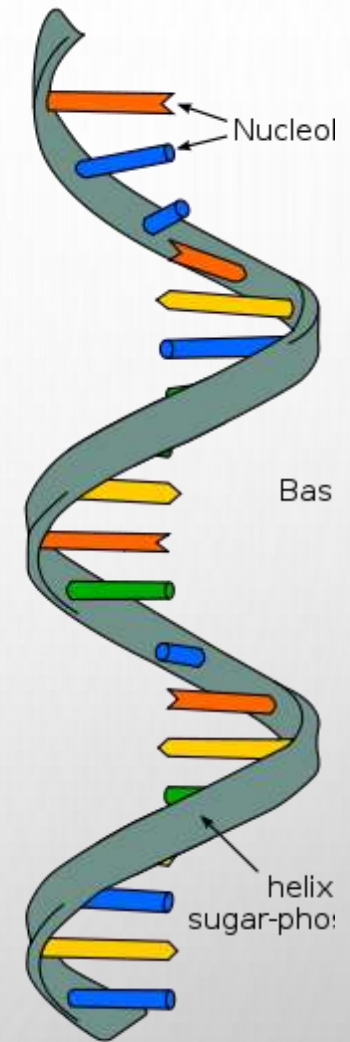
A

Uracil



U

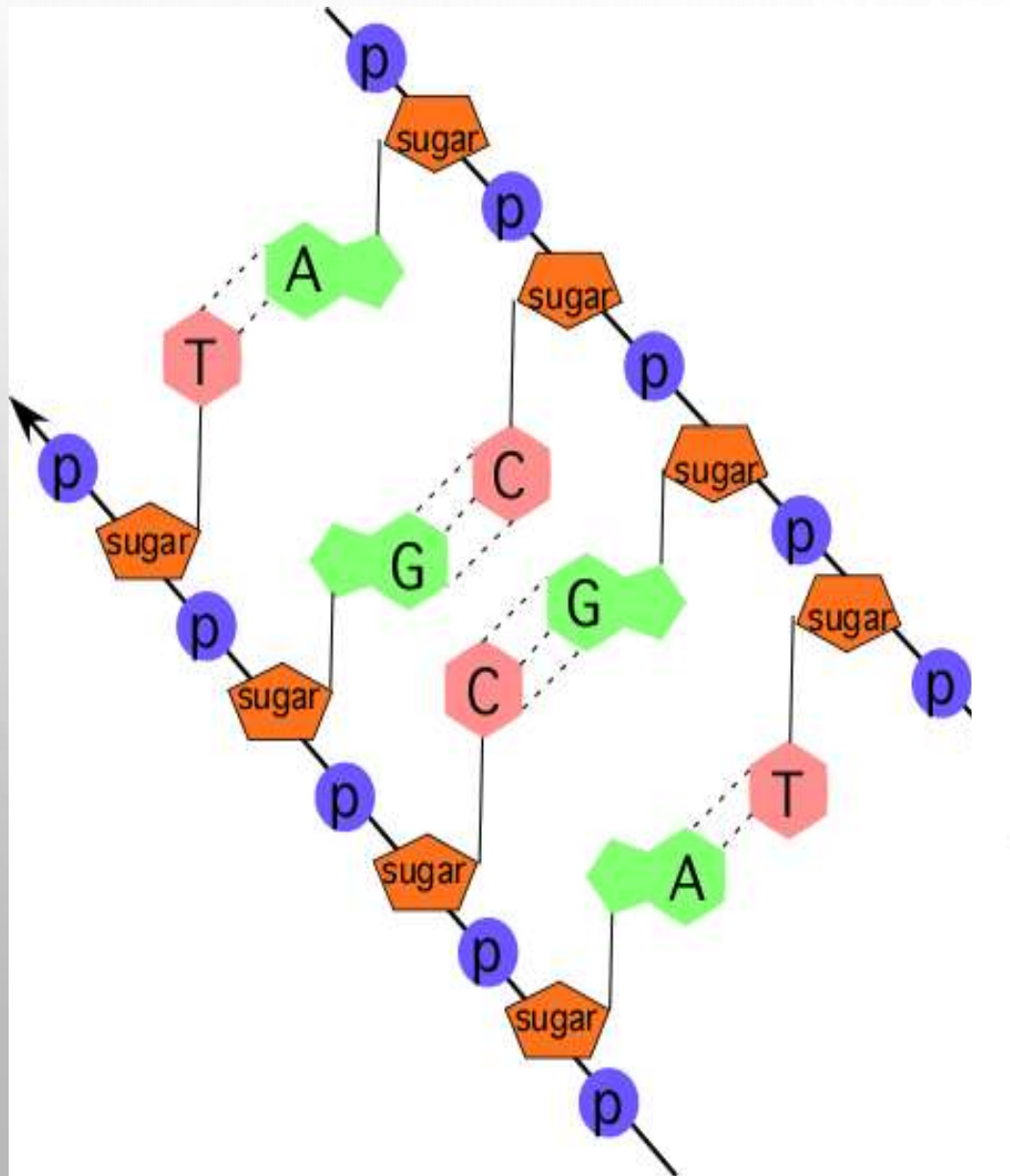
Nucleobases of RNA



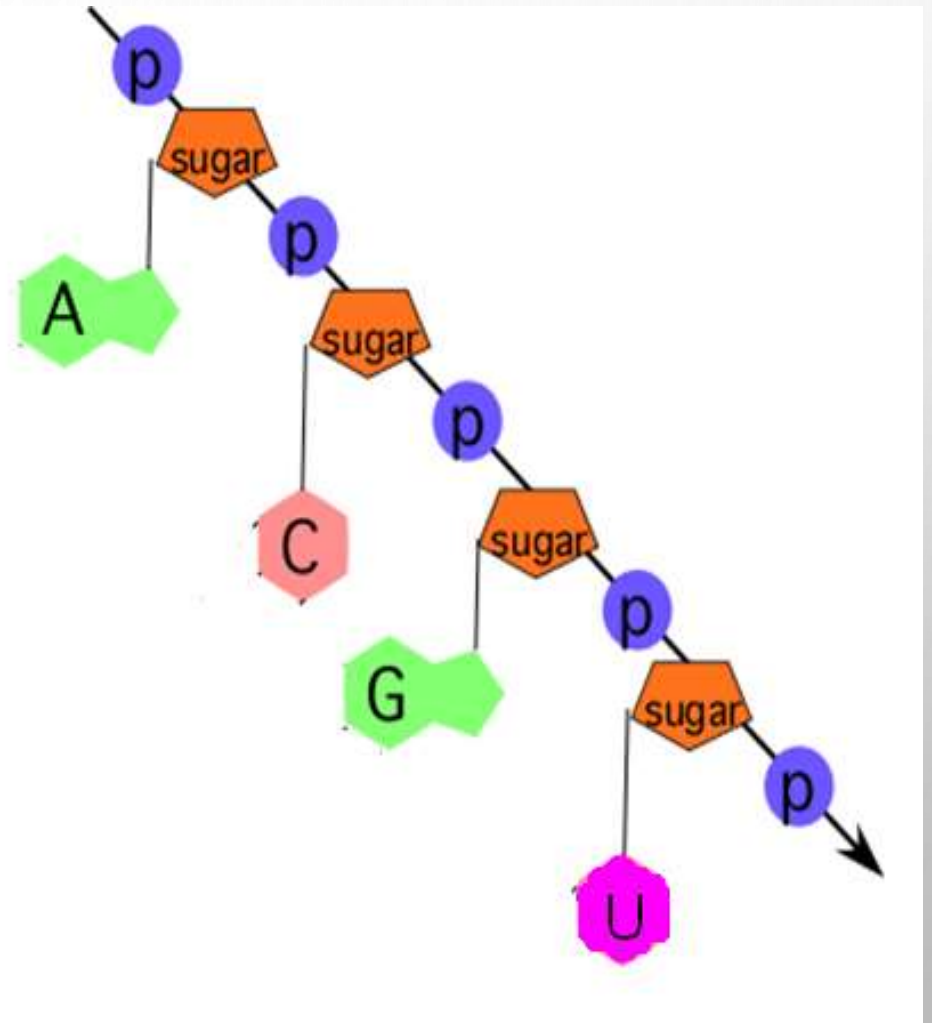
RNA

Ribonucleic acid

DNA




RNA

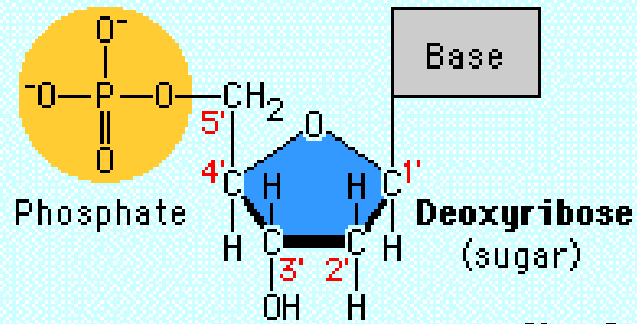




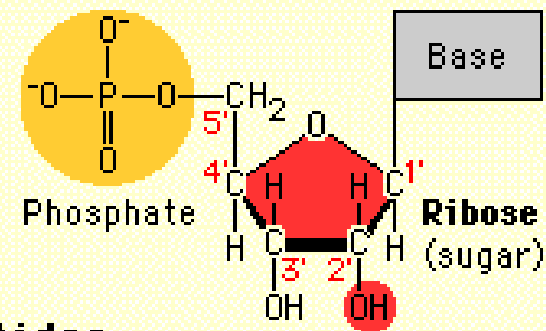
Both DNA and RNA:

- a. are single stranded
 - b. contain the same four nitrogenous bases
 - c. have the same five carbon sugars
 - d. contain phosphate groups
- 

DNA



RNA



Nucleotides

Pyrimidines



Purines



Bases

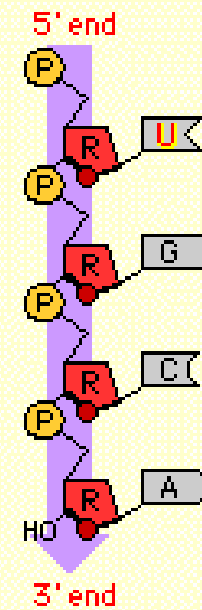
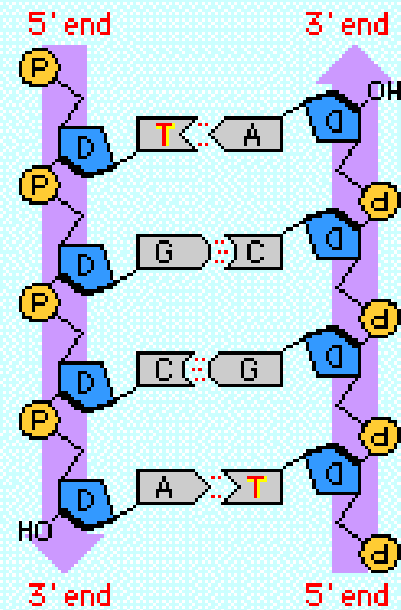
Pyrimidines



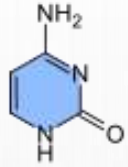
Purines



Polynucleotides

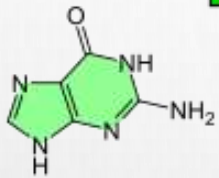


Cytosine



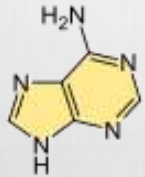
C

Guanine



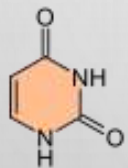
G

Adenine



A

Uracil



U

Nucleobases
of RNA

Nucleobases

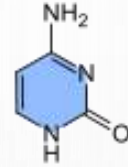
Base pair

helix of
sugar-phosphates

RNA

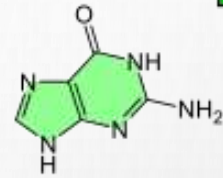
Ribonucleic acid

Cytosine



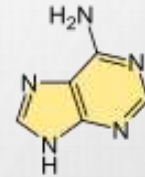
C

Guanine



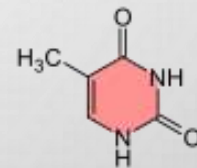
G

Adenine



A

Thymine



T

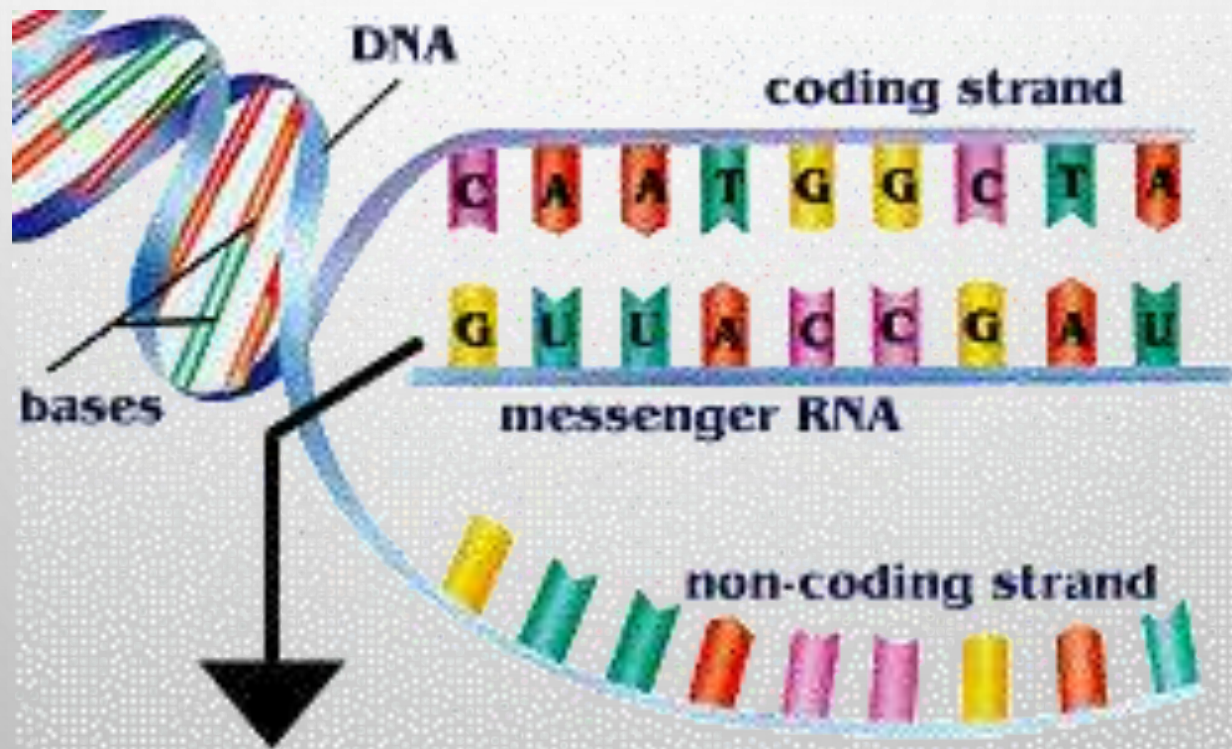
Nucleobases
of DNA

DNA

Deoxyribonucleic acid

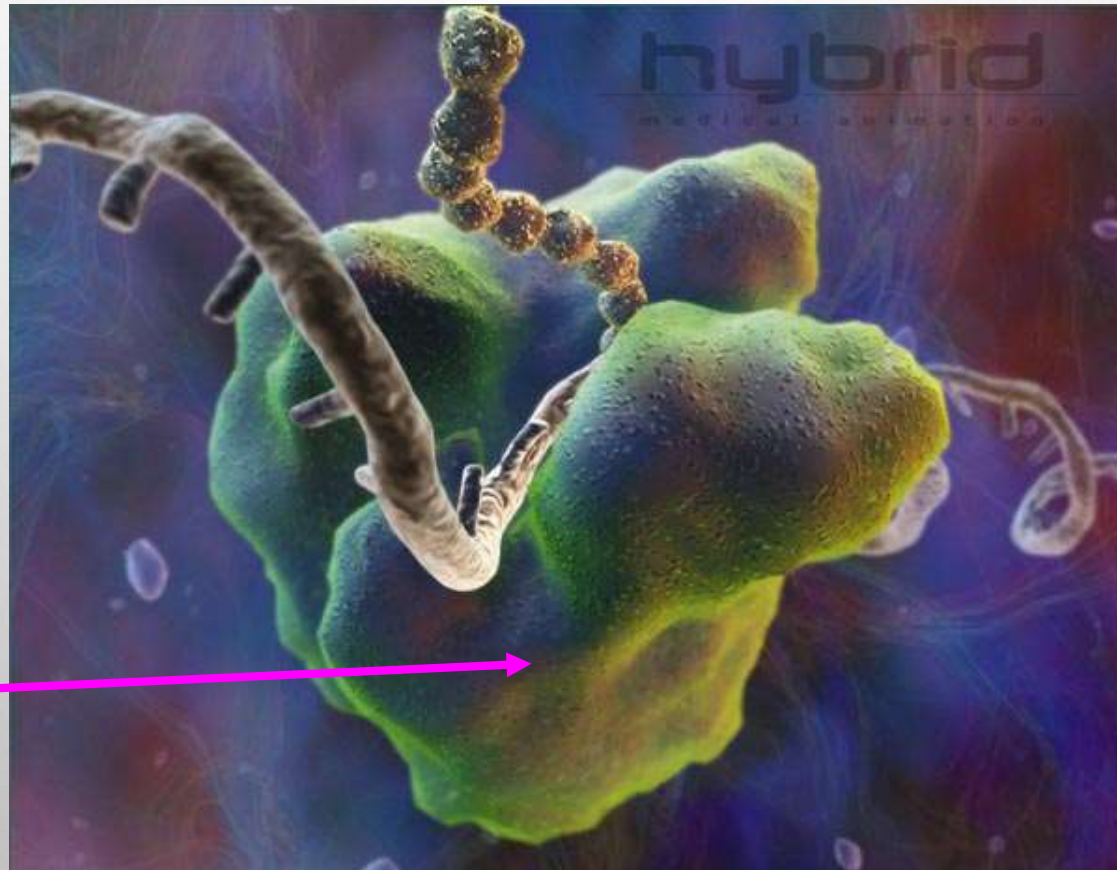
THREE MAIN TYPES OF RNA

- 1. MESSENGER RNA (MRNA) - CARRIES COPIES OF INSTRUCTIONS FOR THE ASSEMBLY OF AMINO ACIDS INTO PROTEINS FROM DNA TO THE REST OF THE CELL (SERVE AS “MESSENGER”)



THREE MAIN TYPES OF RNA

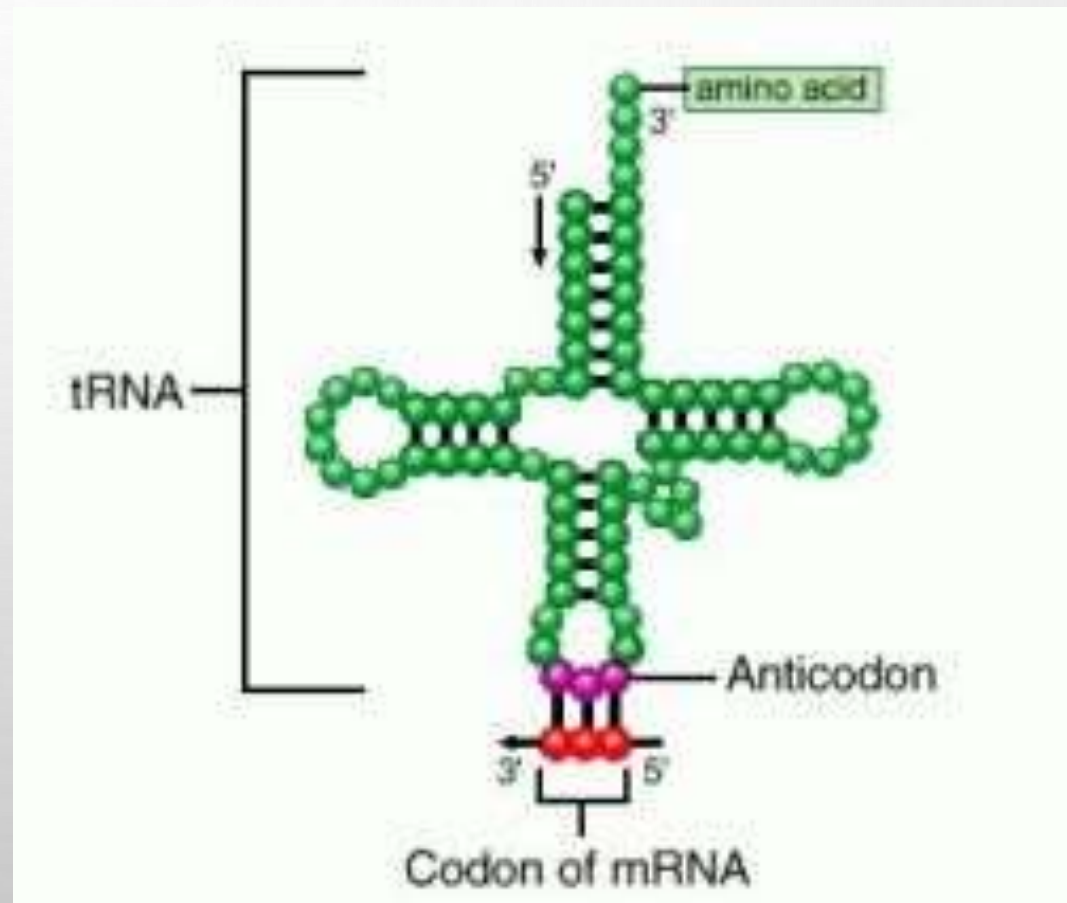
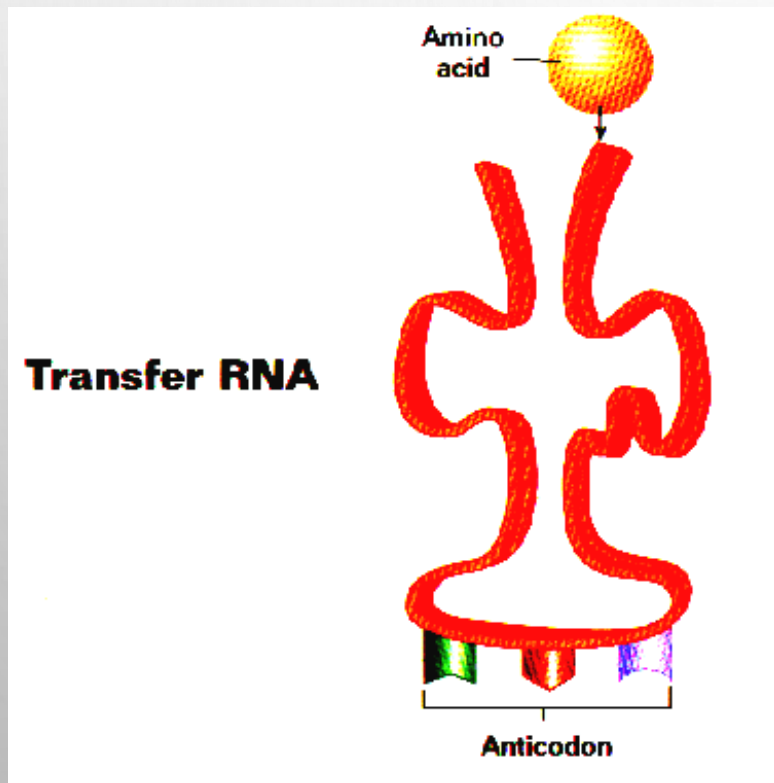
2. RIBOSOMAL RNA (RRNA) – MAKES UP THE MAJOR PART OF RIBOSOMES, WHICH IS WHERE PROTEINS ARE MADE.



**Ribosomal
RNA**

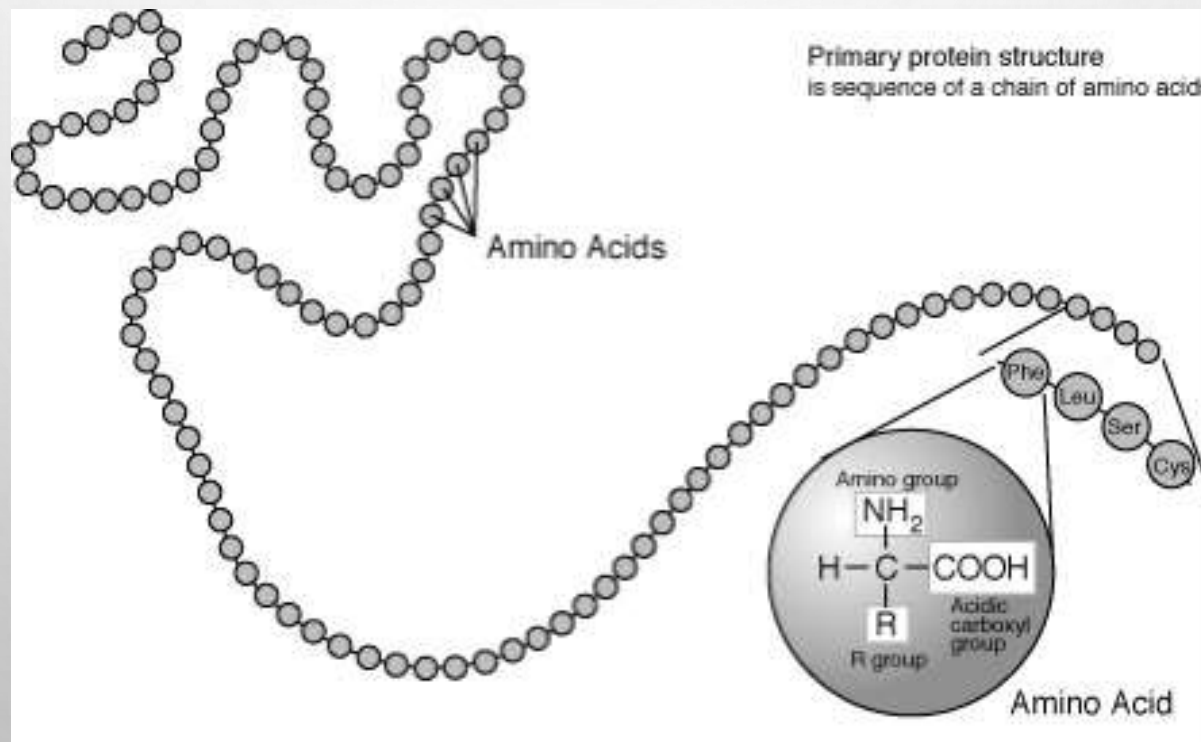
THREE MAIN TYPES OF RNA

3. TRANSFER RNA (TRNA) - TRANSFERS AMINO ACIDS TO RIBOSOMES DURING PROTEIN SYNTHESIS



PROTEINS

- PROTEINS ARE MADE UP OF A CHAIN OF AMINO ACIDS.
- PROTEINS ARE ENZYMES, WHICH CATALYZE AND REGULATE CHEMICAL REACTIONS.



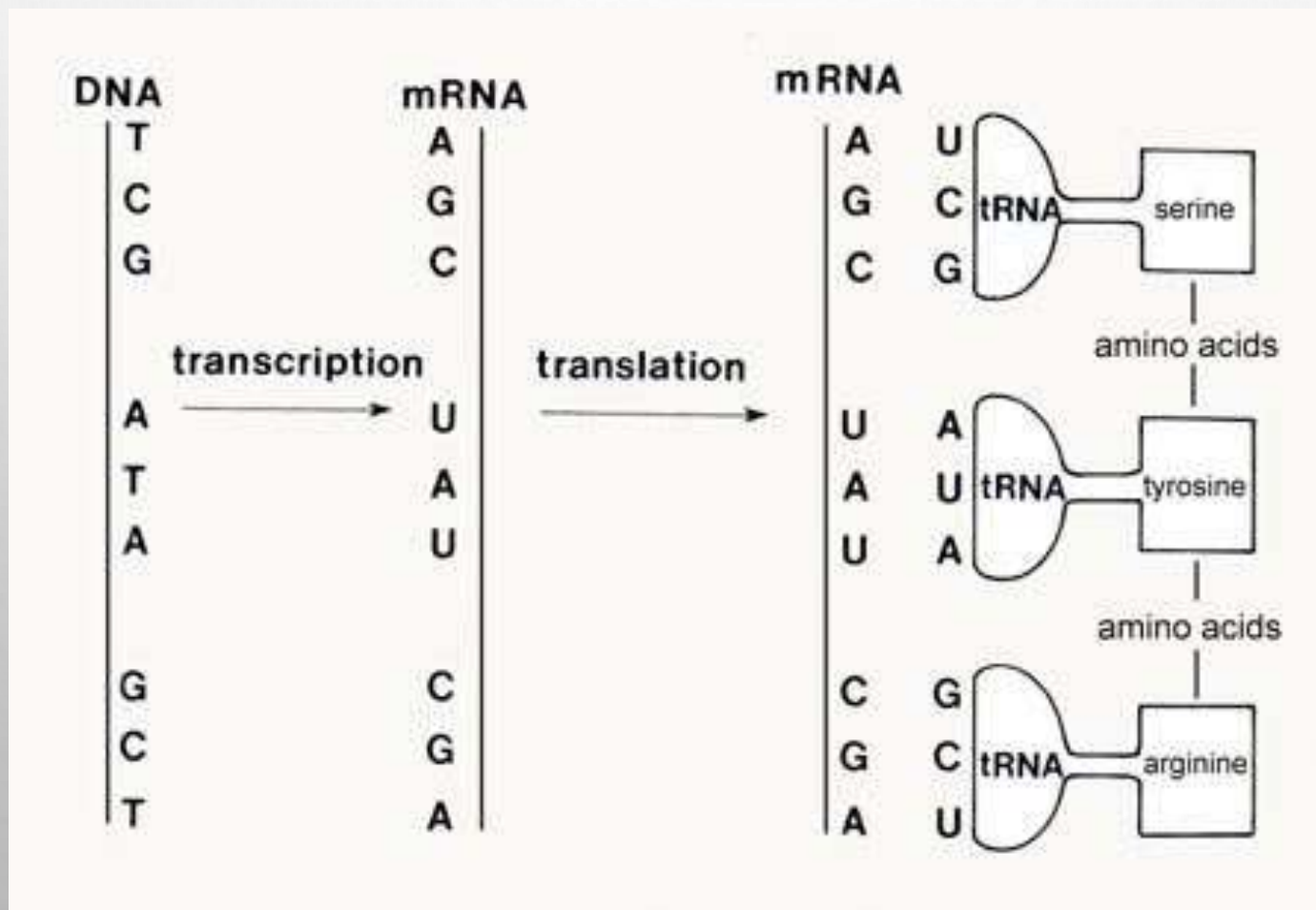
2 STEPS TO MAKE A PROTEIN

1. TRANSCRIPTION

- DNA → RNA

2. TRANSLATION

- RNA → PROTEIN (CHAIN OF AMINO ACIDS)



- WHEN TRANSCRIPTION NEEDS TO TAKE PLACE, DNA MUST PROVIDE THE CODE IN ORDER TO CREATE AN MRNA STRAND.

- MRNA WILL BE ABLE TO LEAVE THE NUCLEUS AND NOW IT HAS THE CODE TRANSCRIBED INSIDE IT'S BASE PAIRS!

- PRACTICE:

- DNA STRAND: TTA ACG GGT CTA
- MATCHING DNA STRAND: AAT TGC CCA GAT
- MRNA: UUA ACG GGU CUA

DNA REPLICATION:

- Copying genetic information for transmission to the next generation
- Occurs in S phase of cell cycle
- Process of DNA duplicating itself
- Begins with the unwinding of the double helix to expose the bases in each strand of DNA
- Each unpaired nucleotide will attract a complementary nucleotide from the medium
 - will form base pairing via hydrogen bonding.
- Enzymes link the aligned nucleotides by phosphodiester bonds to form a continuous strand.

DNA REPLICATION:

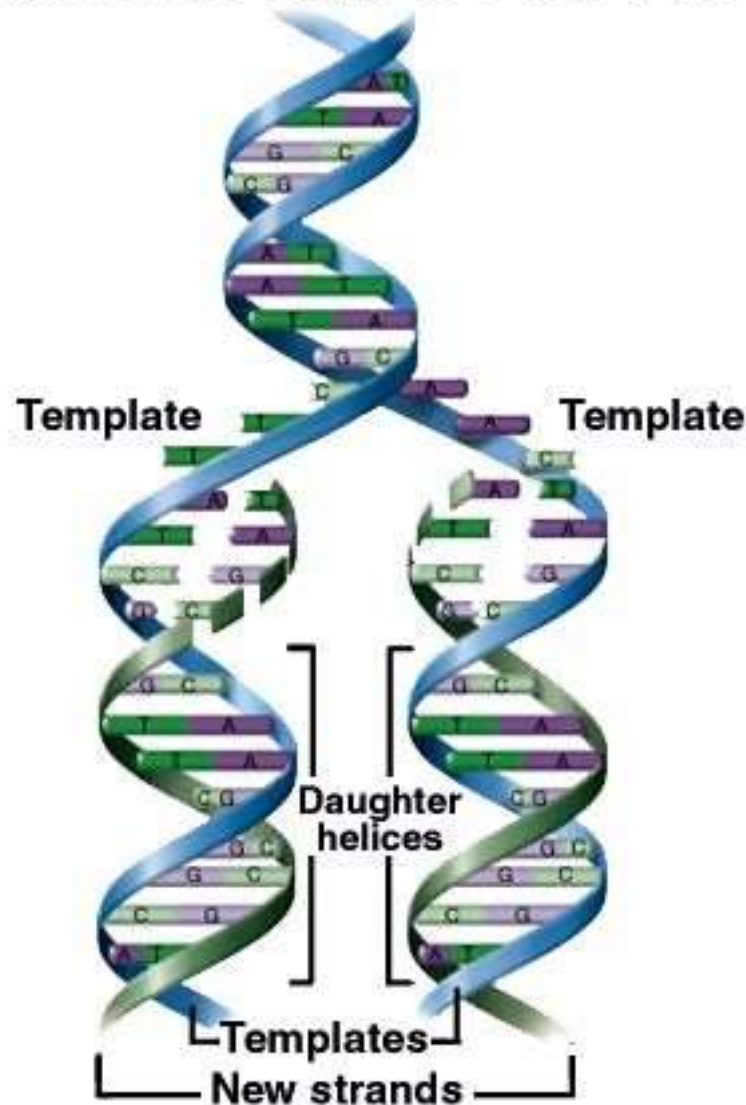
- First question asked was whether duplication was semiconservative or conservative
 - Meselson and Stahl expt
 - Semiconservative -
 - one strand from parent in each new strand
 - Conservative-
 - both strands from parent and other is all new strands

DNA REPLICATION:

- Complementary base pairing produces semiconservative replication
 - Double helix unwinds
 - Each strand acts as template
 - Complementary base pairing ensures that T signals addition of A on new strand, and G signals addition of C
 - Two daughter helices produced after replication

DNA replication: an overview

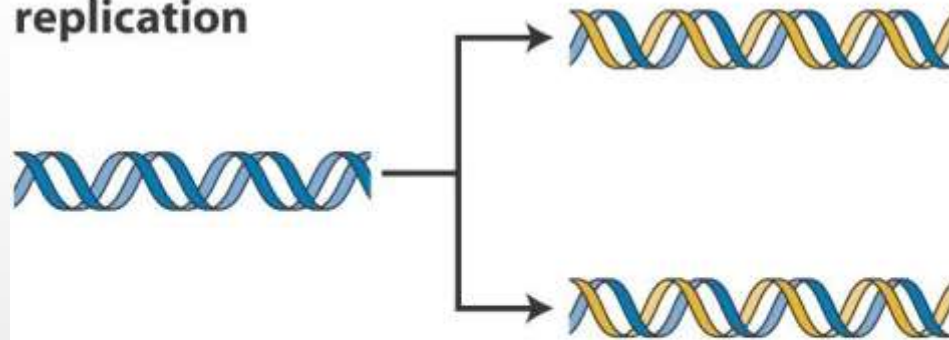
1. Original double helix
2. Strands separate
3. Complementary bases align opposite templates
4. Enzymes link sugar-phosphate elements of aligned nucleotides into a continuous new strand



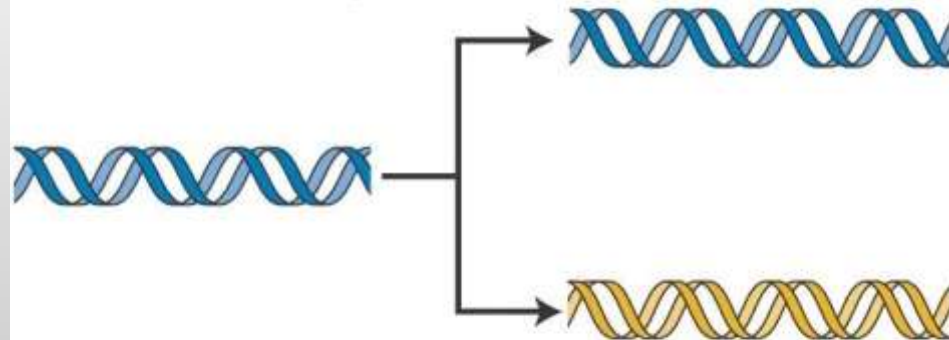
EXPERIMENTAL PROOF OF SEMICONSERVATIVE REPLICATION – THREE POSSIBLE MODELS

- Semiconservative replication –
 - Watson and Crick model
- Conservative replication:
 - The parental double helix remains intact;
 - both strands of the daughter double helix are newly synthesized
- Dispersive replication:
 - At completion, both strands of both double helices contain both original and newly synthesized material.

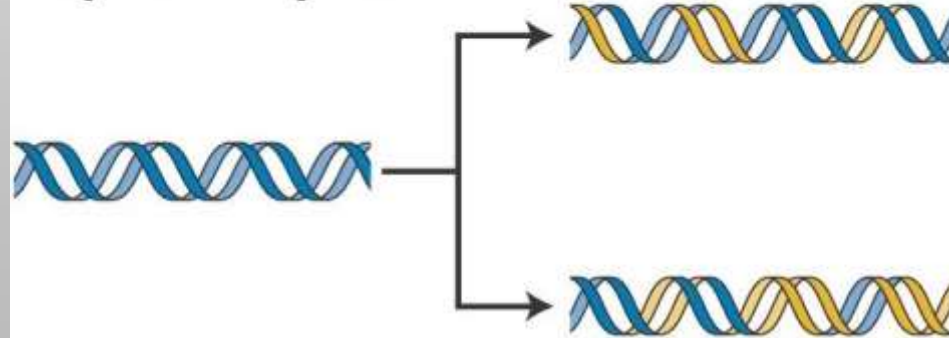
Semiconservative replication



Conservative replication



Dispersive replication



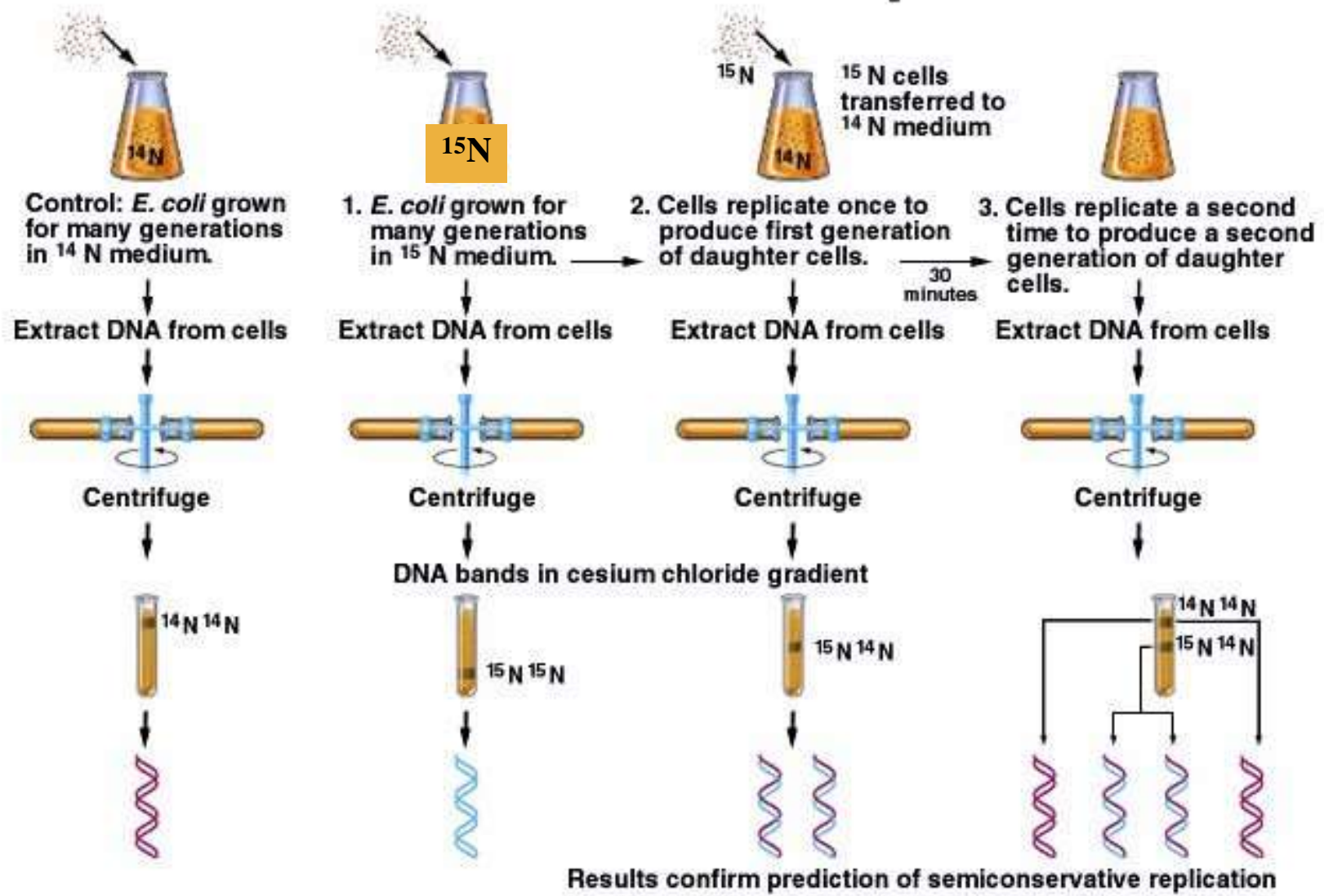
MESELSON-STAHN EXPERIMENTS CONFIRM SEMICONSERVATIVE REPLICATION

- Experiment allowed differentiation of parental and newly formed DNA.
- Bacteria were grown in media containing either normal isotope of nitrogen (^{14}N) or the heavy isotope (^{15}N).
- DNA banded after **equilibrium density gradient centrifugation** at a position which matched the density of the DNA:
 - heavy DNA was at a higher density than normal DNA.

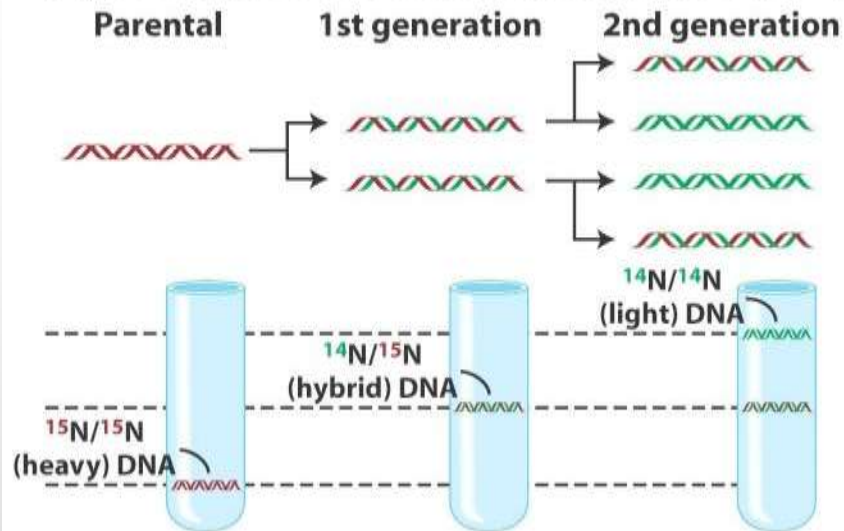
MESELSON-STAHN EXPERIMENTS CONFIRM SEMICONSERVATIVE REPLICATION

- When bacteria grown in ^{15}N were transferred to normal ^{14}N containing medium,
 - the newly synthesized DNA strand had the ^{14}N while the parental strand had ^{15}N .
- They checked the composition of the resulting DNA molecules by density gradient centrifugation,
 - found an intermediate band,
 - indicating a hybrid molecule
 - containing both ^{14}N and ^{15}N DNA.

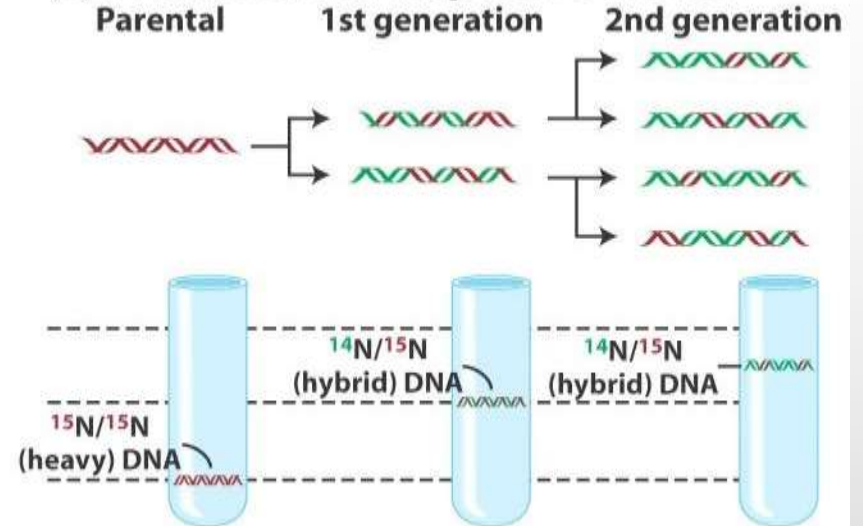
The Meselson-Stahl experiment



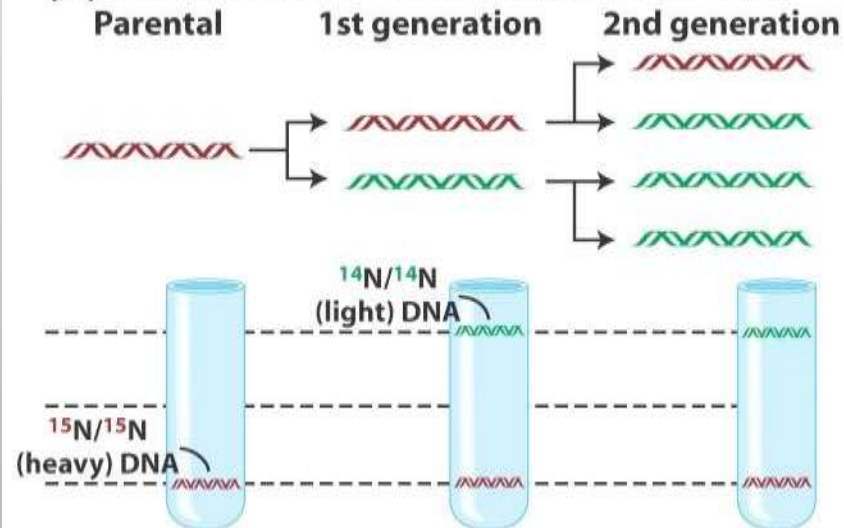
(a) Predictions of semiconservative model



(c) Predictions of dispersive model



(b) Predictions of conservative model



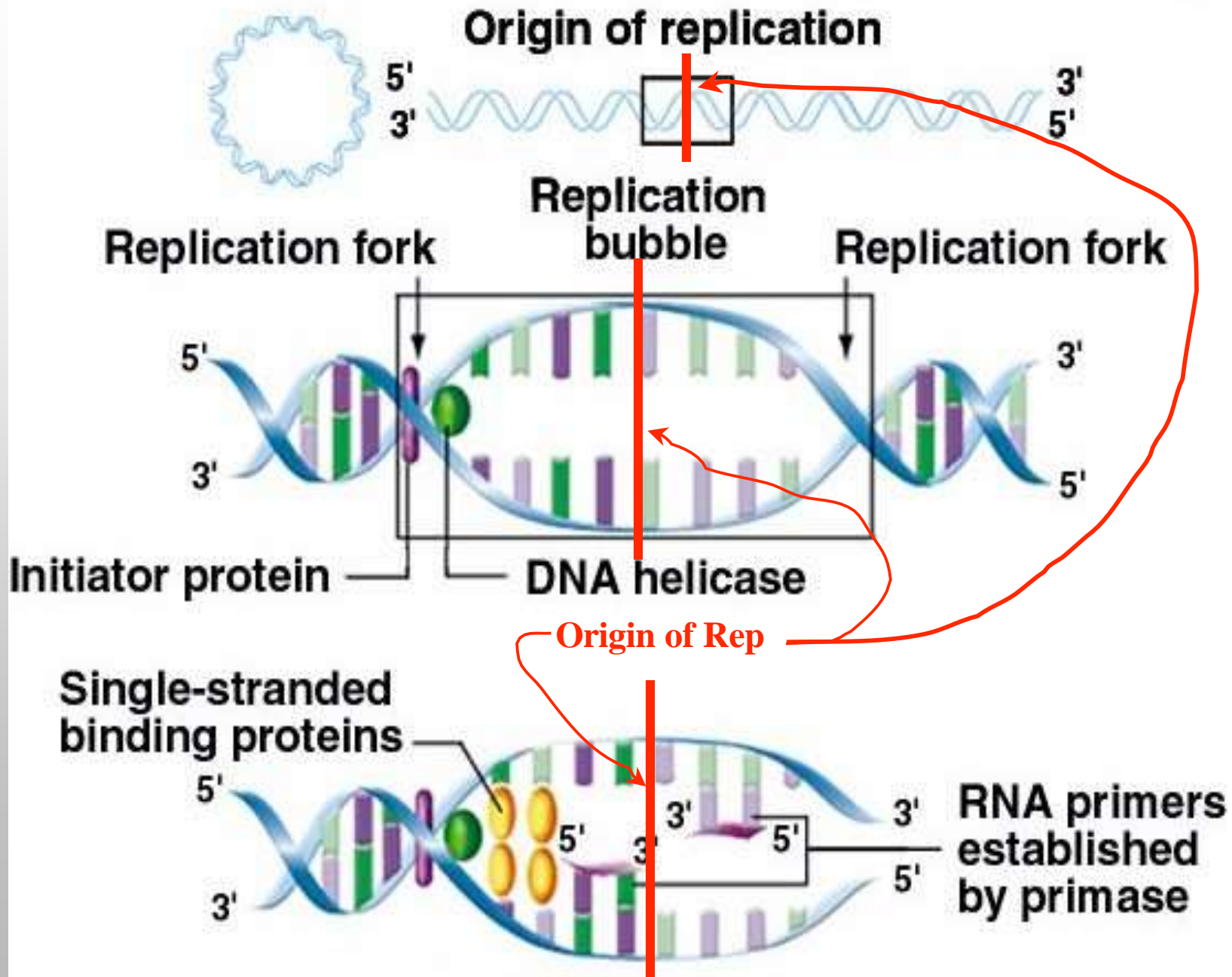
THE MECHANISM OF DNA REPLICATION

- Tightly controlled process,
 - occurs at specific times during the cell cycle.
- Requires:
 - a set of **proteins** and **enzymes**,
 - and requires energy in the form of **ATP**.
- Two basic steps:
 - **Initiation**
 - **Elongation**.
- Two basic components:
 - **template**
 - **primer**.

THE MECHANISM OF DNA REPLICATION (PROKARYOTIC)

- DNA polymerase
 - the enzyme that extends the primer;
 - Pol III –
 - produces new stands of complementary DNA
 - Pol I –
 - fills in gaps between newly synthesized Okazaki segments
- additional enzymes/proteins
 - i) DNA helicase –
 - unwinds double helix
 - ii) Single-stranded binding proteins –
 - keep helix open
 - iii) Primase –
 - creates RNA primers to initiate synthesis
 - iv) Ligase –
 - welds together Okazaki fragments

Mechanism of DNA replication, 1

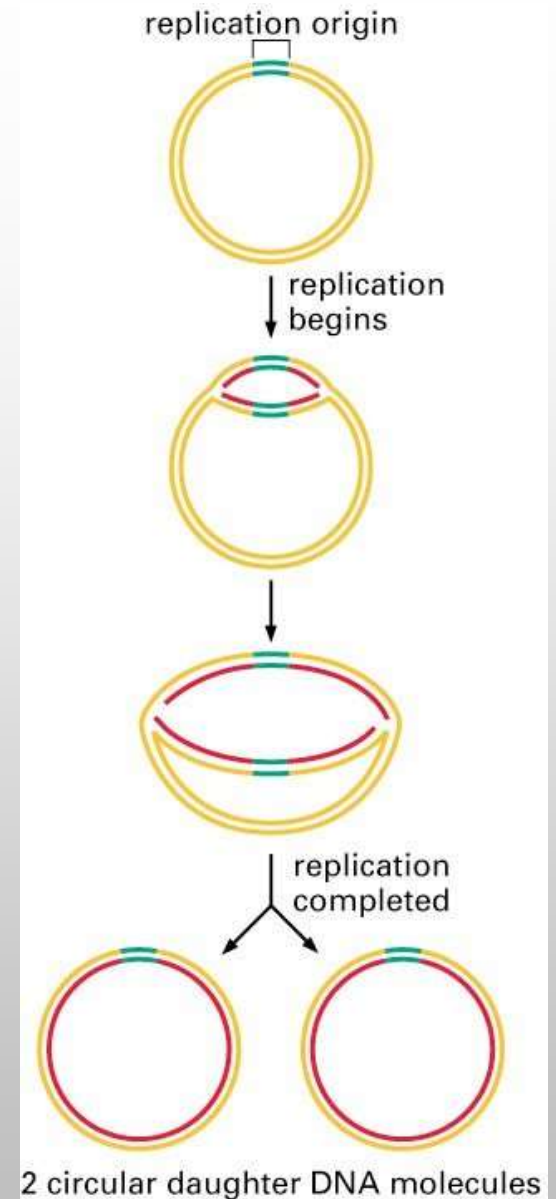


ORIGINS OF REPLICATION

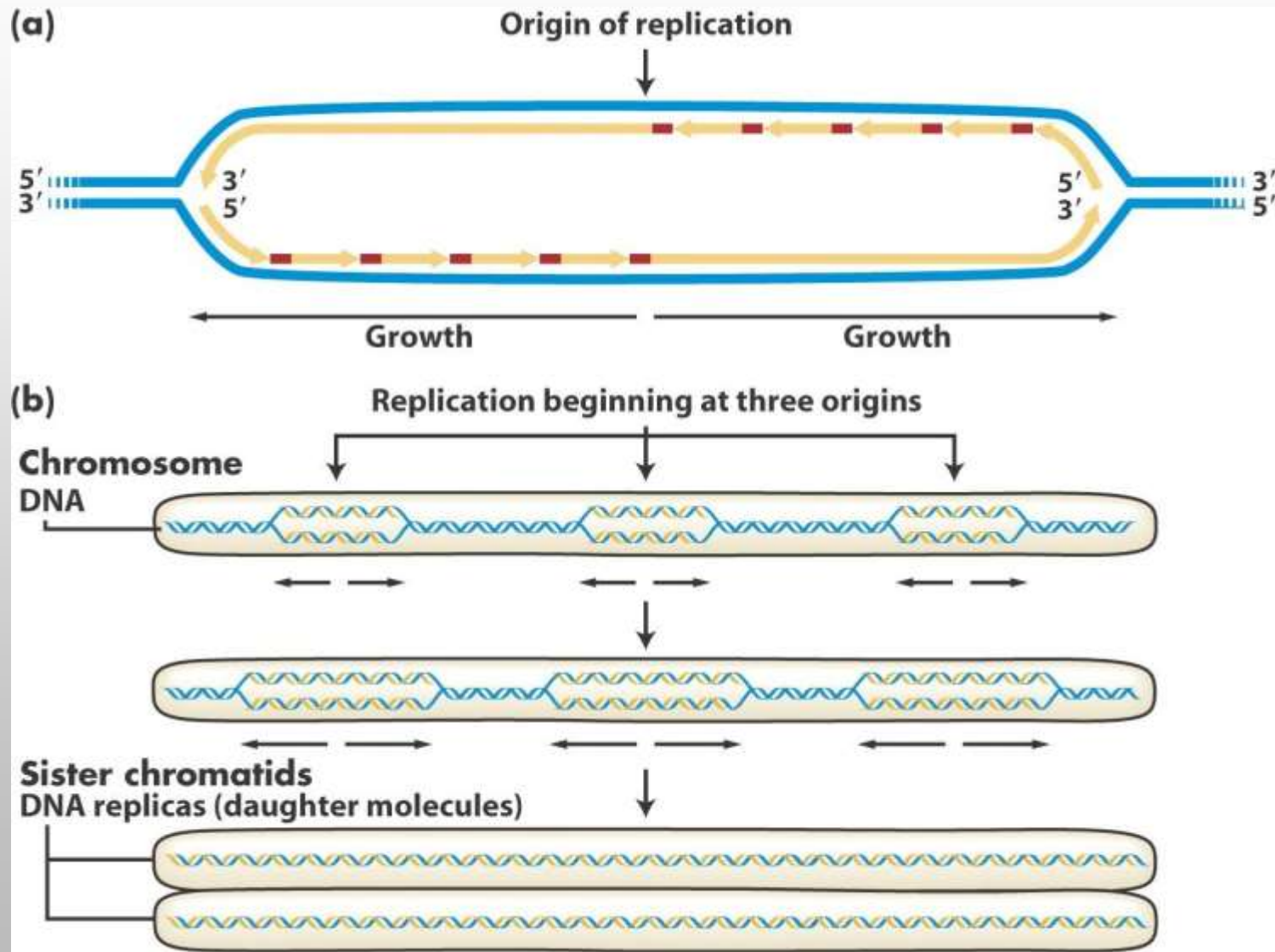
- Replication proceeds in both directions (bidirectionally) from a single origin of replication on the prokaryotic circular chromosome
- Replication proceeds in both directions (bidirectionally) from hundreds or thousands of origins of replication on each of the linear eukaryotic chromosomes.

ORIGINS OF REPLICATION

- Bacteria have 1 origin of replication per one chromosome
- They only have one chromosome = 1 origin!



EUKARYOTIC ORIGINS OF REPLICATION



- Replication Initiation:

- Primase and the RNA Primer

- Replication Elongation:

- DNA polIII

- Must have 3' to add to

- Replication is Finished:

- DNA polI removes primer

- Fills gap using 3'ends

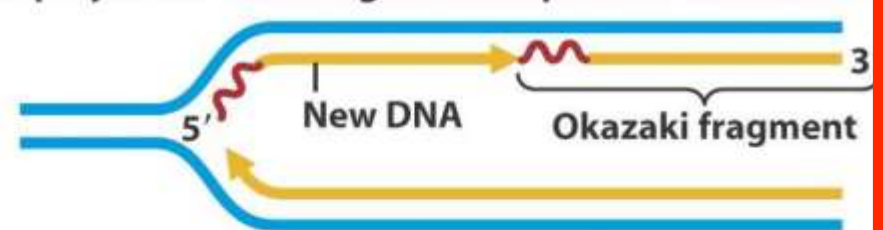
- DNA ligase connects frags

- Uses 5' ends!

1. Primase synthesizes short RNA oligonucleotides (primer) copied from DNA.



2. DNA polymerase III elongates RNA primers with new DNA.



3. DNA polymerase I removes RNA at 5' end of neighboring fragment and fills gap.

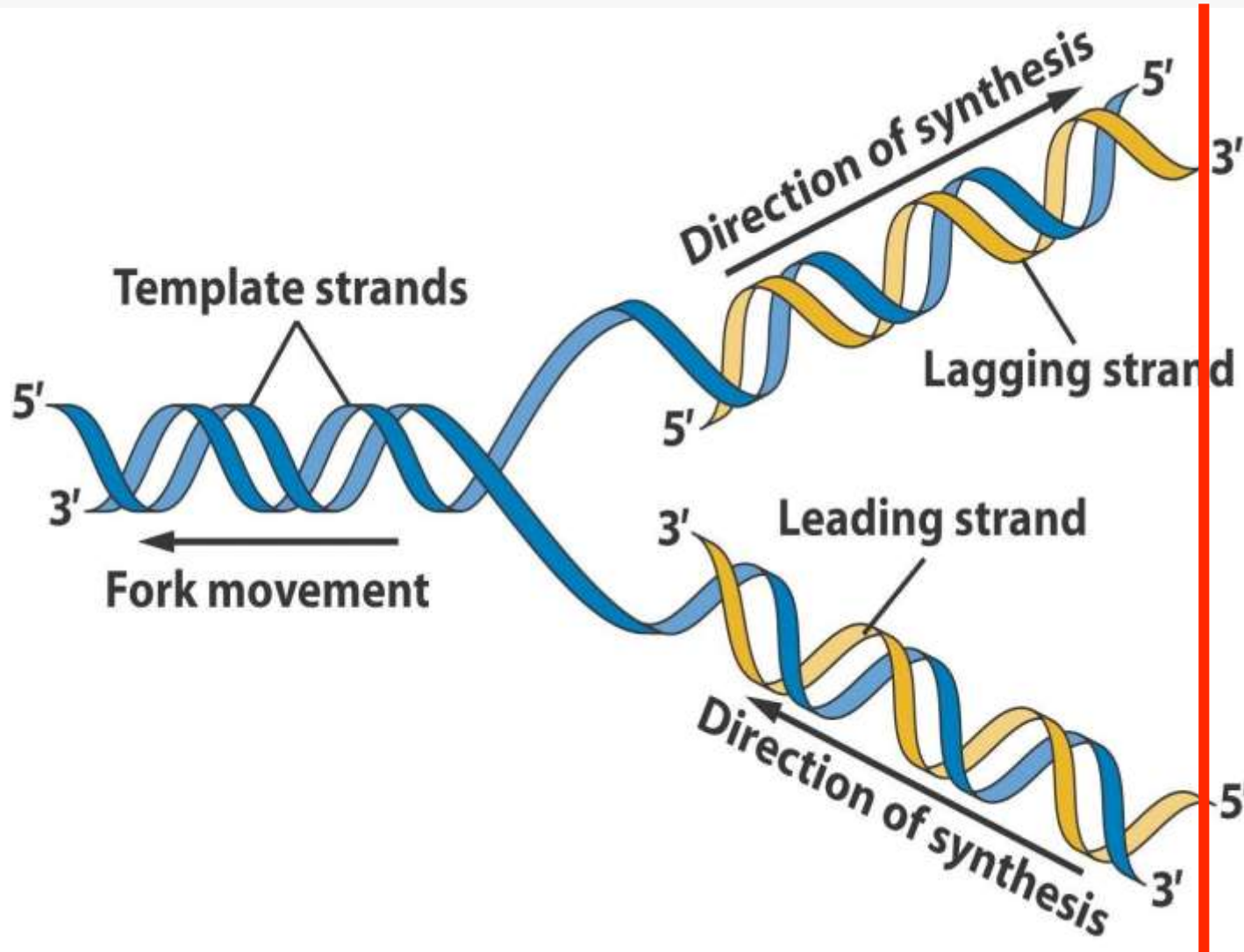


4. DNA ligase connects adjacent fragments.



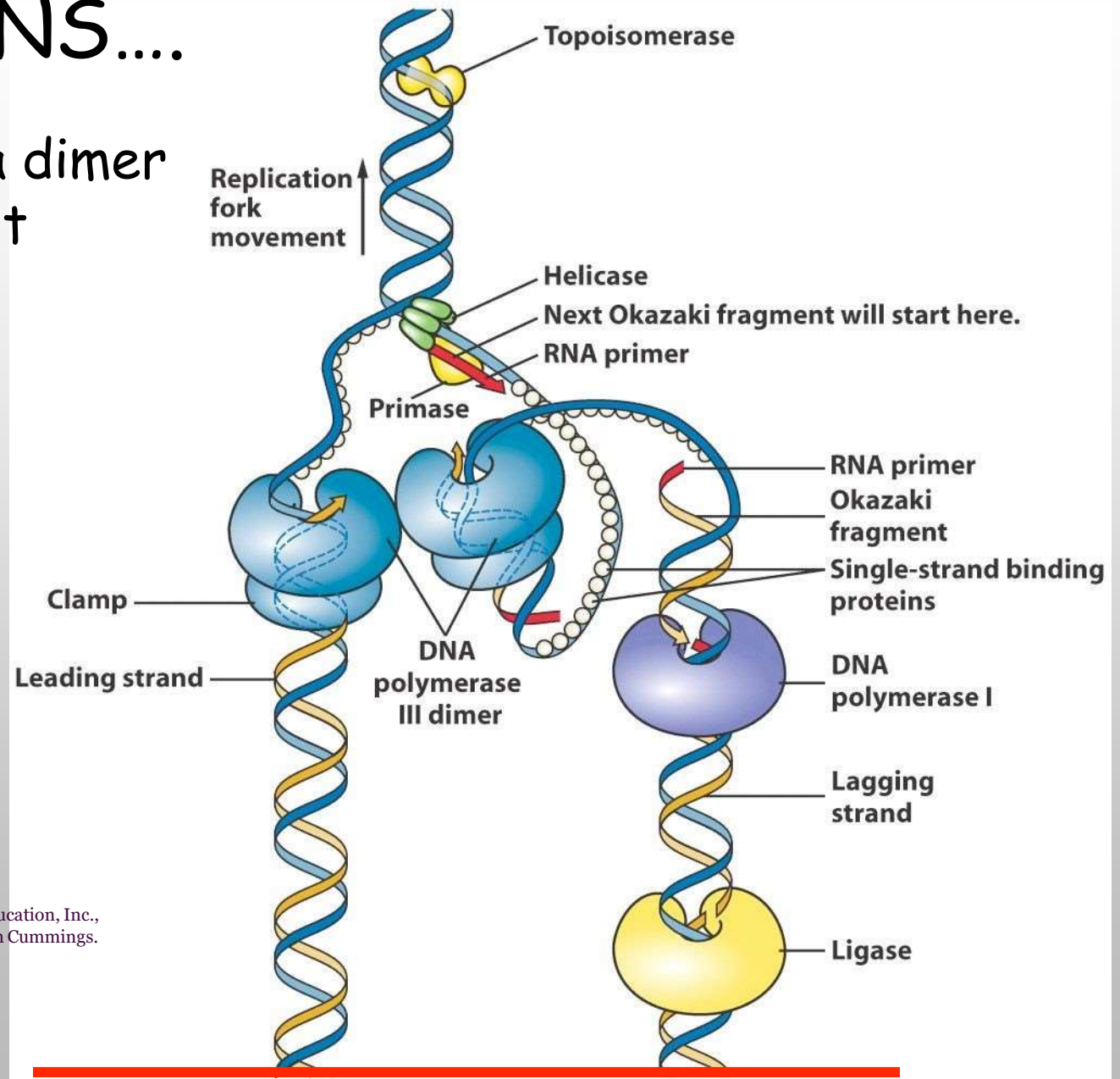
REPLICATION FORK

Origin of Rep



WHAT REALLY HAPPENS....

DNA pol works as a dimer
Lagging strand must
loop around to
accommodate
dimerization



Peter J. Russell, *iGenetics*: Copyright © Pearson Education, Inc.,
publishing as Benjamin Cummings.

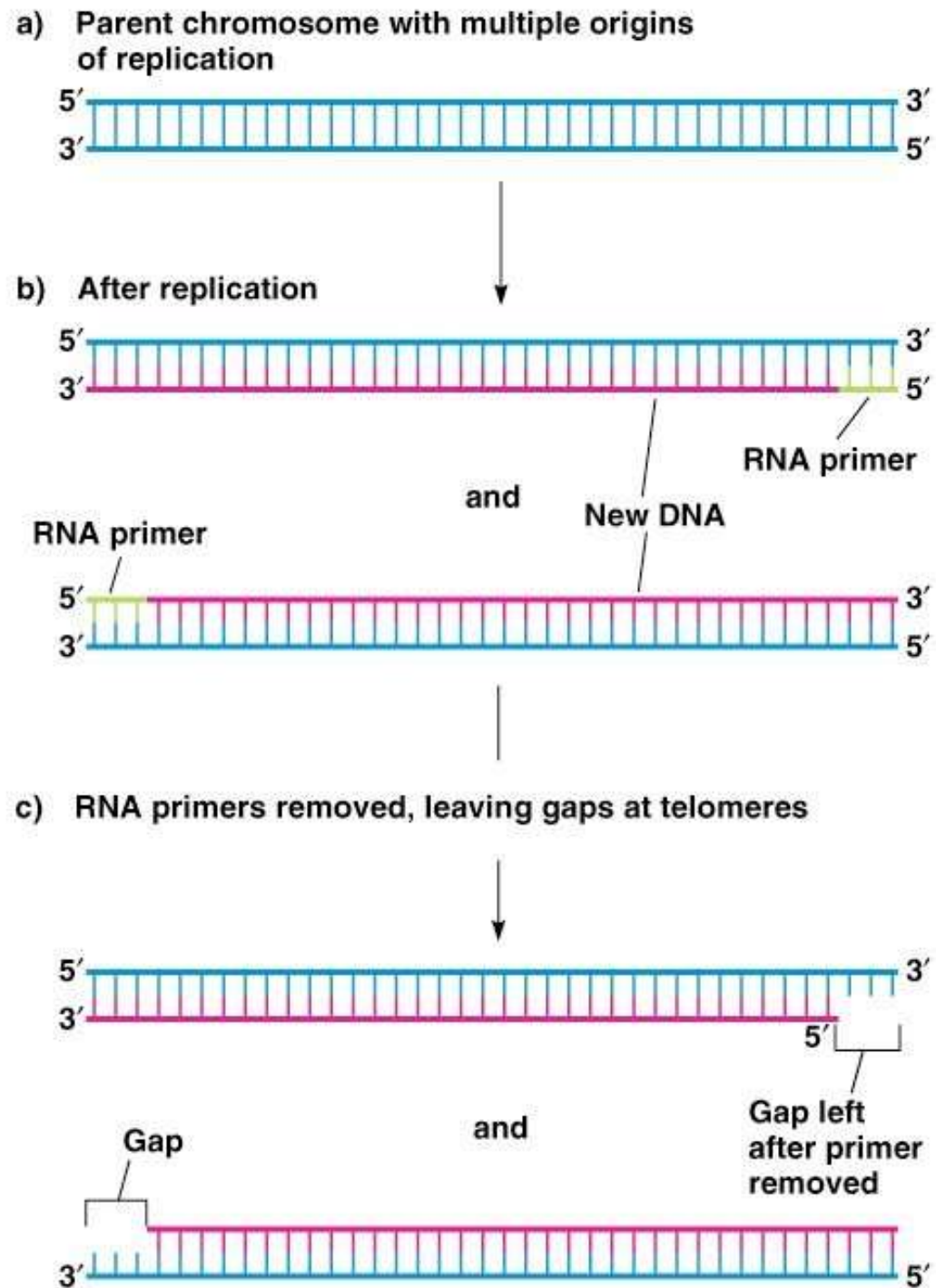
Origin of Rep

REPLICATION TERMINATION

- THE ENDS OF CHROMOSOMES (TELOMERES) CANNOT BE REPLICATED ON THE LAGGING STRAND BECAUSE THERE IS NO PRIMER AVAILABLE.
- **TELOMERASES**
 - ENZYMES THAT CONTAIN RNA PRIMERS WHICH EXTEND THE ENDS OF CHROMOSOMES (NOT NORMALLY EXPRESSED IN SIGNIFICANT LEVELS)
 - TELOMERES FORM A SORT OF SINGLE STRANDED CAP AROUND THE CHROMOSOME ENDS TO PROTECT THEM FROM BEING DEGRADED
 - CHROMOSOME ENDS ARE PROGRESSIVELY SHORTENED WITH EACH ROUND OF REPLICATION.
 - “OLD” CELLS WITH SHORTENED TELOMERES UNDERGO APOPTOSIS
 - No death signal
 - PROTECTIVE FOR NORMAL CELLS
 - KILL THE OLD AND POSSIBLY MUTATED

FIG.
11.14

The problem of
replicating
completely a linear
chromosome in
eukaryotes

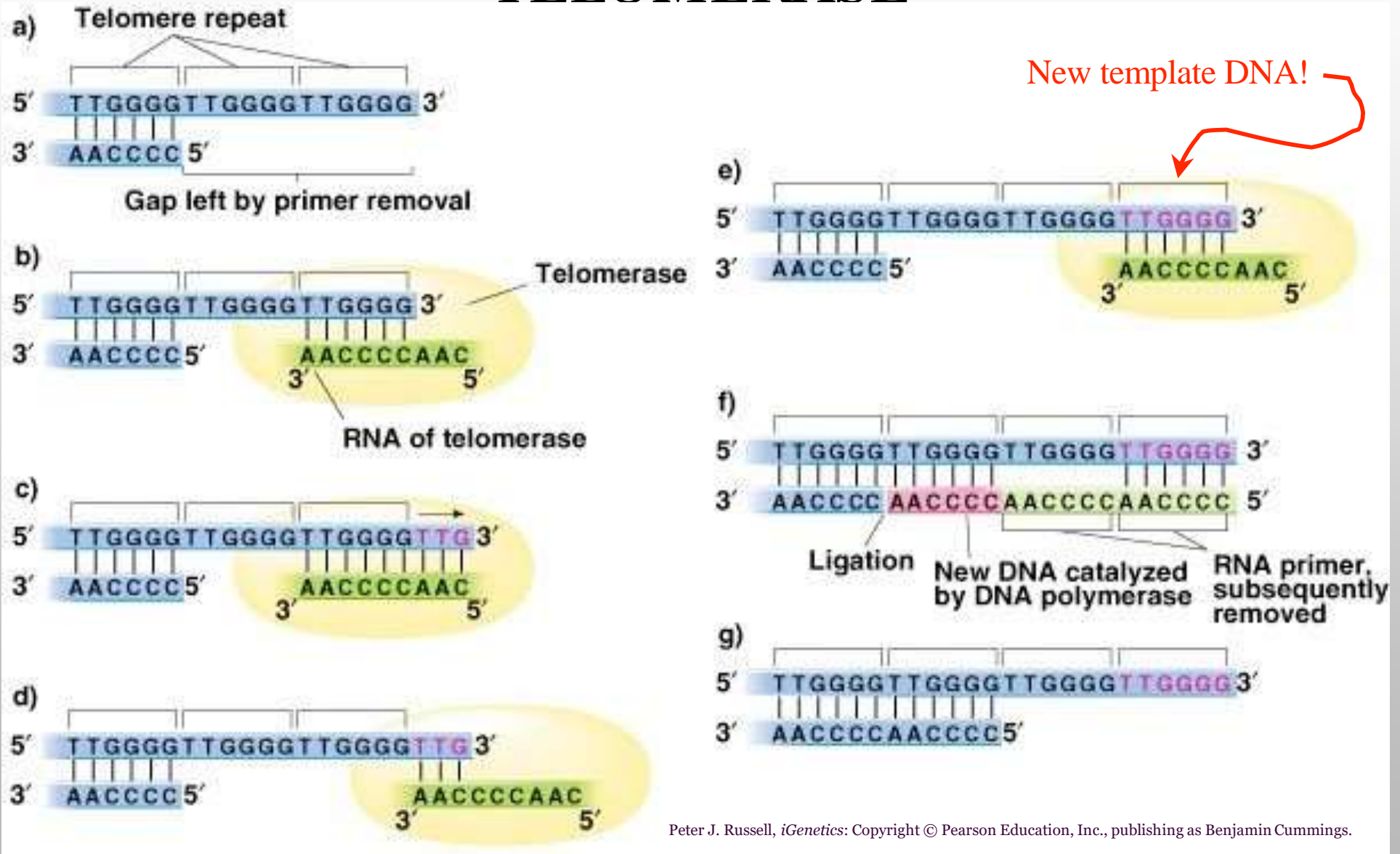


REPLICATING THE ENDS OF CHROMOSOMES

- telomerase adds an RNA primer complementary to telomere sequences
 - chromosomal replication proceeds by adding to the 3' end of the primer
- Fills the gap left behind by replication
- Telomerase enzyme can also add DNA basepairs to the TEMPLATE DNA
 - complementary to the RNA primer basepairs
 - Using an RNA template to make DNA, telomerase functions as a reverse transcriptase called TERT (telomerase reverse transcriptase).
 - This goes against the Central Dogma....
 - Evolutionarily thought to be derived from a Retrovirus

FIG. 3.19

SYNTHESIS OF TELOMERIC DNA BY TELOMERASE



REPLICATION AT THE CHROMOSOMAL LEVEL

- Replication is bidirectional.
- For circular DNA (and linear chromosomes)
 - the unwinding at the replication forks causes **supercoiling**.
- **DNA topoisomerases**
 - enzymes that help relax the DNA by nicking the strands
 - releasing the twists
 - then rejoining the DNA ends.
 - Example is DNA gyrase

THE BIDIRECTIONAL REPLICATION OF A CIRCULAR CHROMOSOME (PROKARYOTIC)

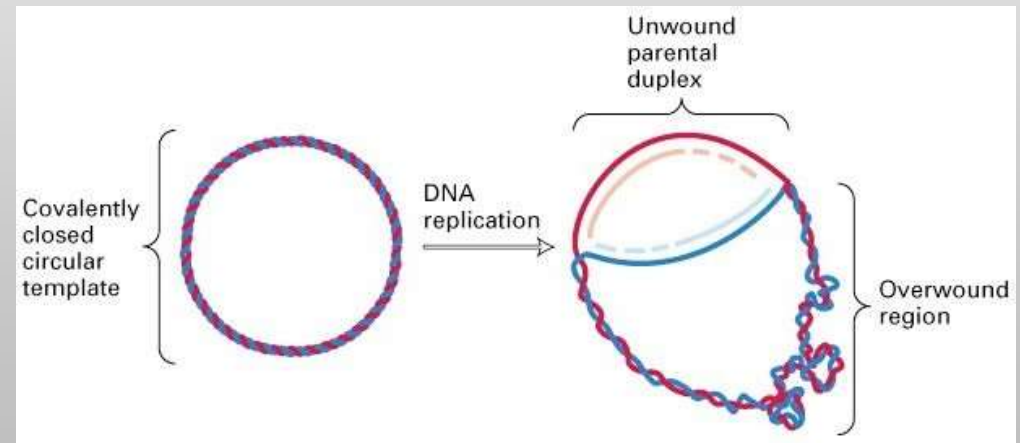
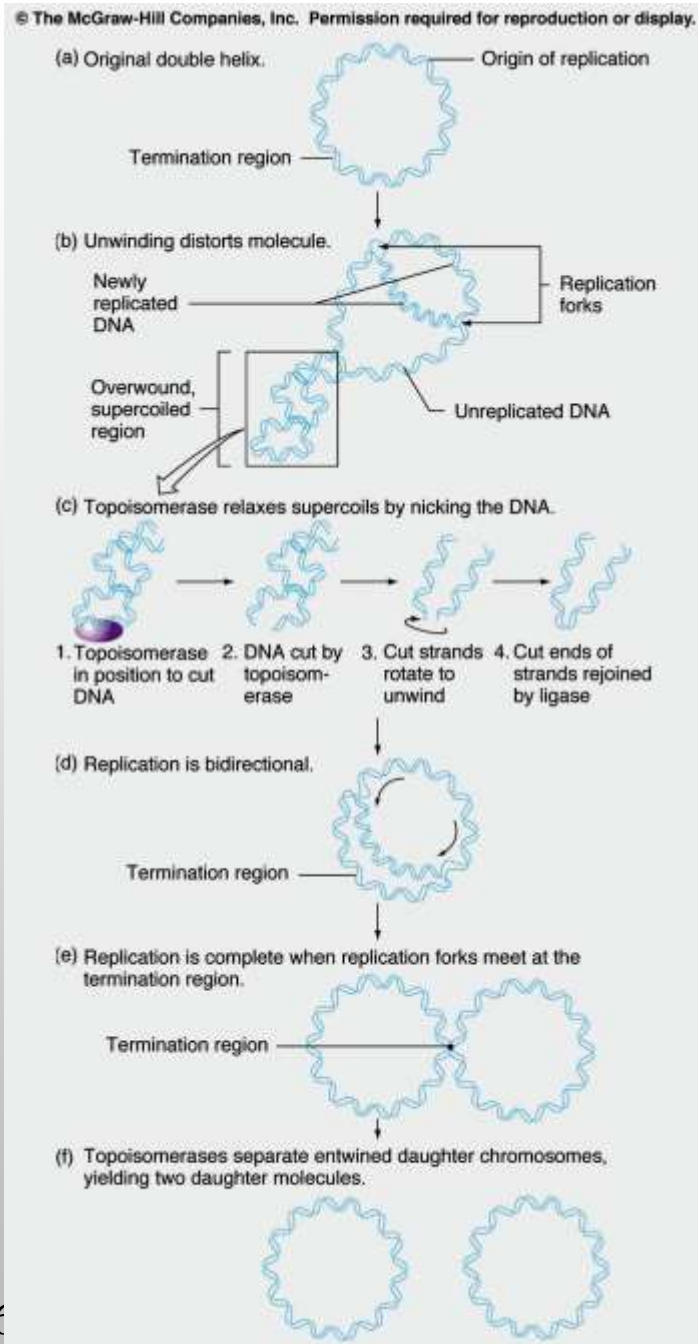
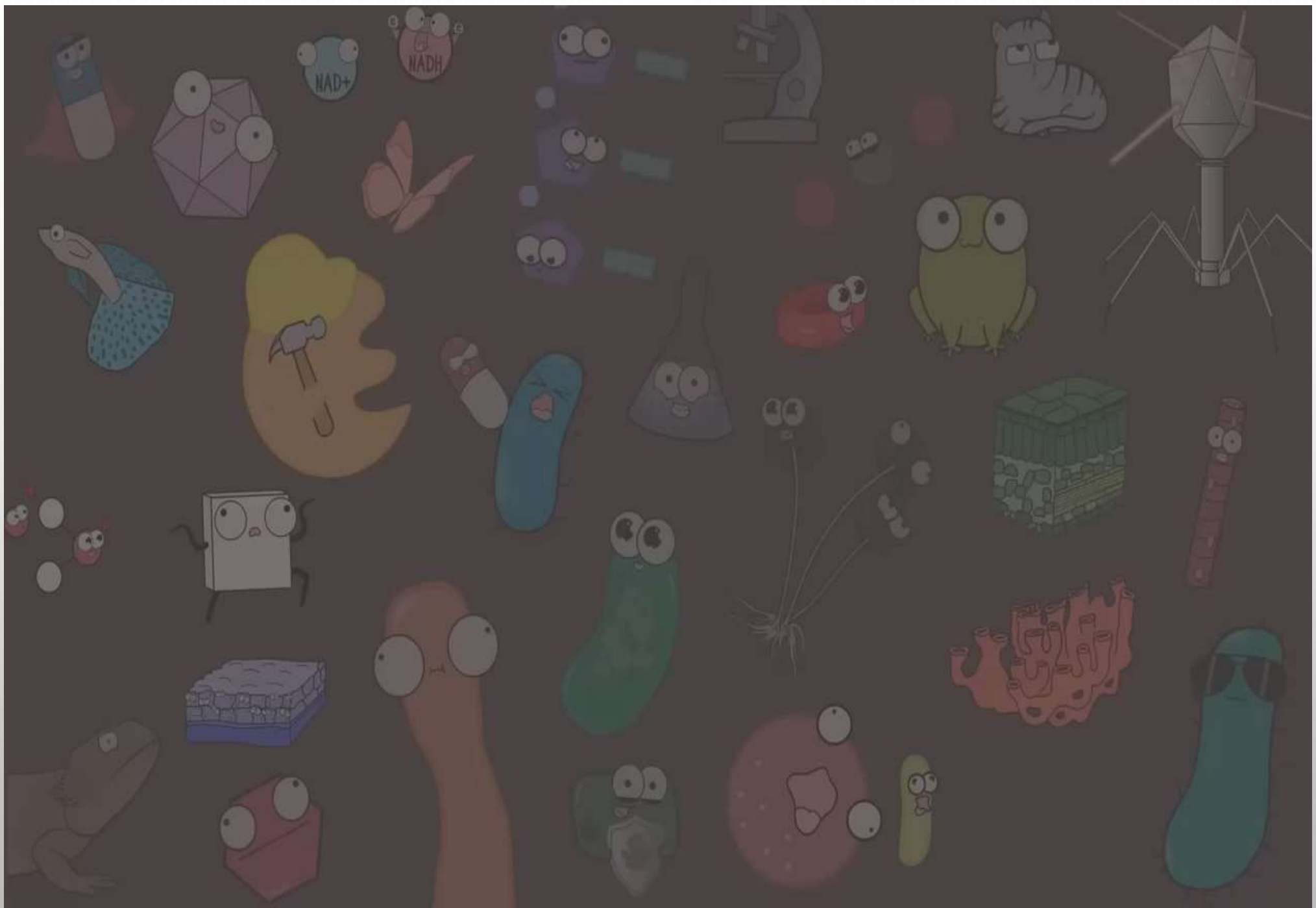


Fig. 6



ASSEMBLING NEWLY REPLICATED DNA INTO NUCLEOSOMES

- When eukaryotic DNA is replicated, it complexes with histones.
 - This requires synthesis of histone proteins and assembly of new nucleosomes.
- Transcription of histone genes is initiated near the end of G1 phase, and translation of histone proteins occurs throughout S phase.
- Assembly of newly replicated DNA into nucleosomes is shown in Figure 11.16.

THE ASSEMBLY OF NUCLEOSOMES AFTER REPLICATION

