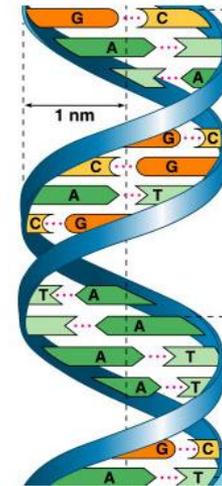


Chapter 10.



The Genetic Material ???



DNA History

- It took many years and the work of many different scientists to discover that **DNA** is the genetic material

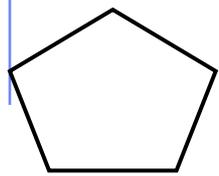
- ◆ Friedrich Meischer (1869)
- ◆ P.A. Levene (1920s)
- ◆ Frederick Griffith (1928)
- ◆ Avery, McCarty & MacLeod (1944)
- ◆ Ervin Chargaff (1947)
- ◆ Hershey & Chase (1952)
- ◆ Rosalind Franklin & Maurice Wilkins (1953)
- ◆ Watson & Crick (1953)

Friedrich Meischer (1869)

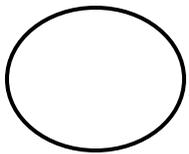
- As he studied salmon sperm for clues to understand heredity, he was the first to isolate a material called nuclein from the nucleus of the sperm.
- Chemical tests revealed that nuclein was part protein and part DNA
- Another clue that made him go “Hmmm” was when he later isolated nuclein from the nuclei of white blood cells in pus-soaked medical bandages.

P.A. Levene (1920s)

- Discovered that DNA is made from smaller **L-shaped** “building blocks” called nucleotides that have 3 chemical parts:



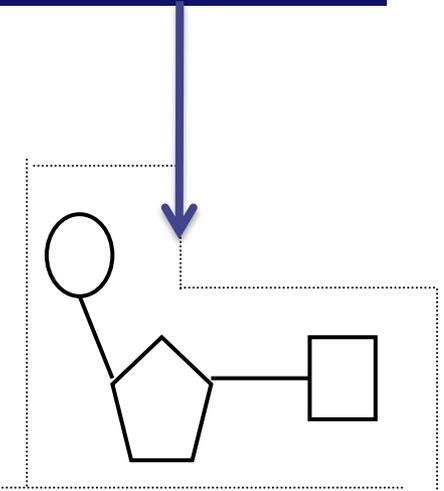
→ 5-C sugar = deoxyribose



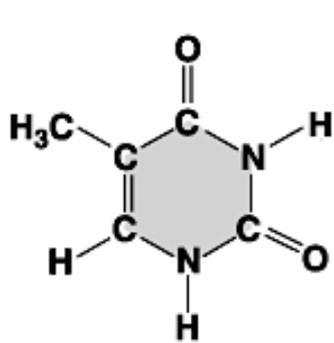
→ phosphate group (PO_4)



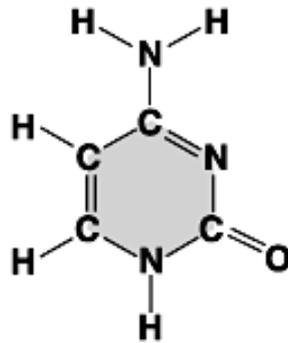
→ nitrogen base: 1 of 4



- DNA has four kinds of N-bases, A (Adenine), T (Thymine), C (Cytosine), and G (Guanine)

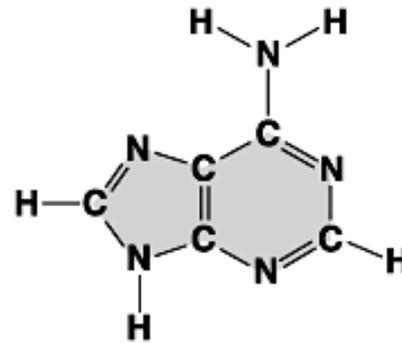


Thymine (T)

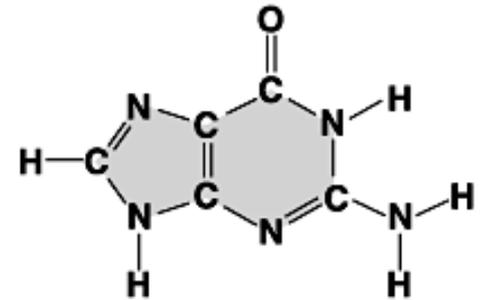


Cytosine (C)

Pyrimidines



Adenine (A)



Guanine (G)

Purines

1928

Fred Griffith (1928)

- While working on a cure for pneumonia with mice, he accidentally discovered the mystery of bacterial transformation



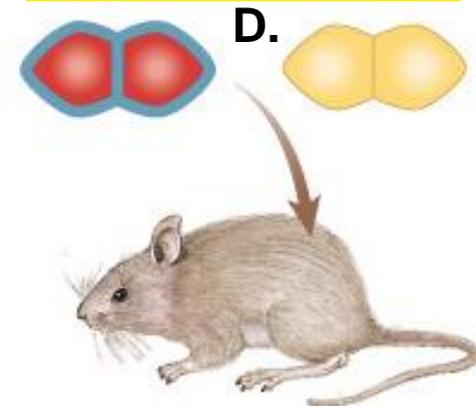
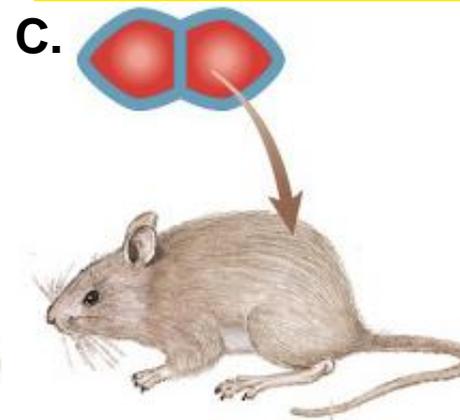
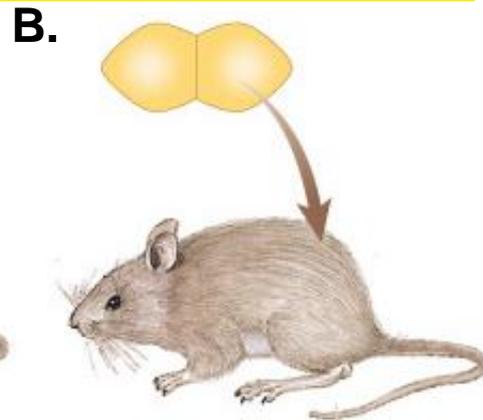
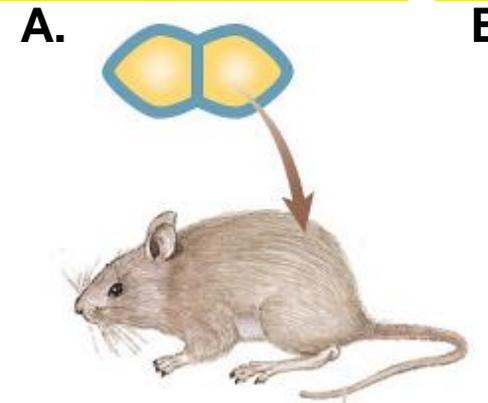
Griffith's 4-part experiments

live pathogenic
(smooth = S)
strain of *bacteria*

live non-pathogenic
(rough = R)
strain of *bacteria*

heat-killed
pathogenic (S)
bacteria

mix heat-killed
pathogenic (S) &
living (R) *bacteria*



Explanation? How could there be living **S** bacteria??

2 Hypotheses:

- 1) Resurrection

- ◆ Somehow the dead S bacteria came back to life ???

- 2) Transformation

- ◆ Some chemical factor in the dead S bacteria was able to communicate with and transform the non-harmful R bacteria into killer S bacteria

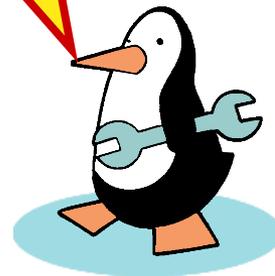
- Fred Griffith would spend many more years in his lab trying to identify this mysterious “transforming factor”

1944

Avery, McCarty & MacLeod (1944)

- ◆ purified both DNA & proteins from *Streptococcus pneumoniae* bacteria
 - which will transform non-pathogenic bacteria?
- ◆ injected protein into bacteria
 - no effect
- ◆ injected DNA into bacteria
 - transformed harmless (R) bacteria into virulent (S) bacteria
- ◆ DNA is the “Transforming Factor”

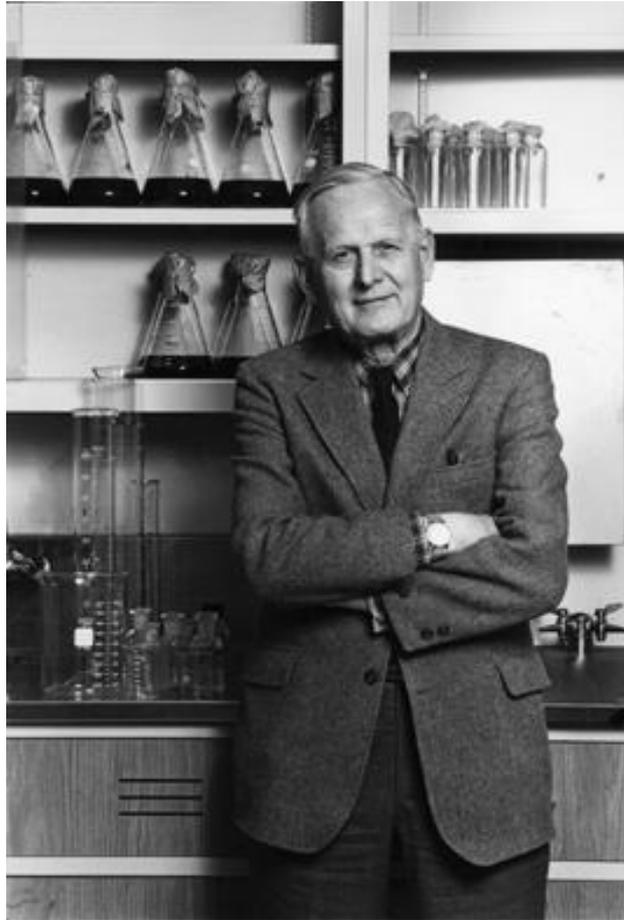
What's the conclusion?



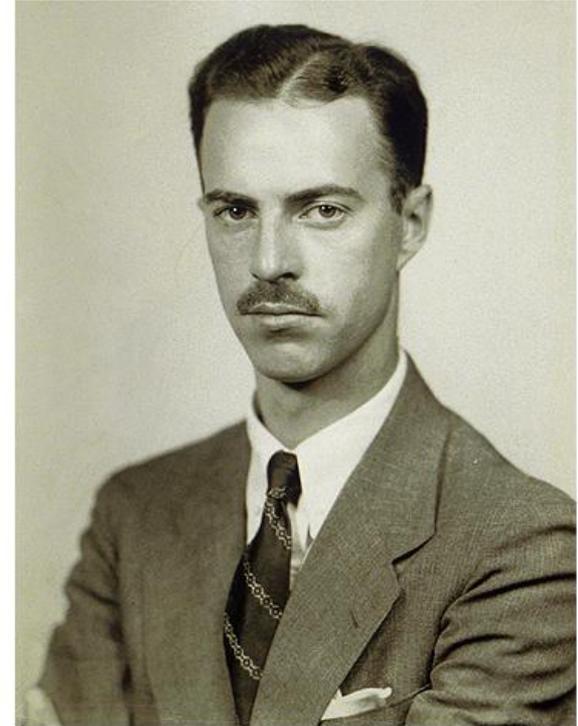
Avery, McCarty & MacLeod



Oswald Avery



Maclyn McCarty

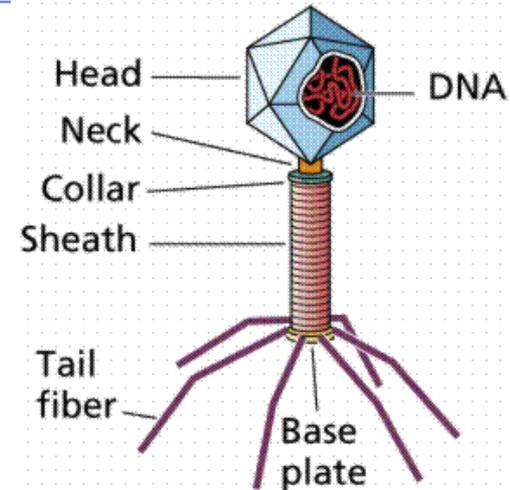


Colin MacLeod

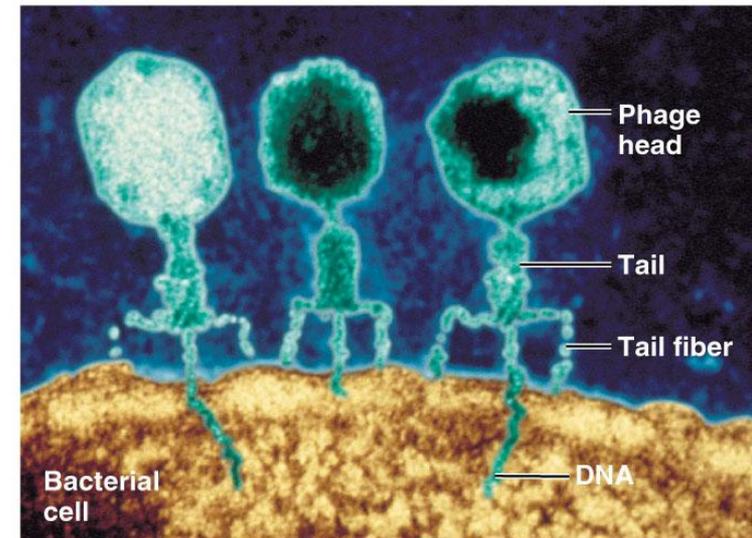
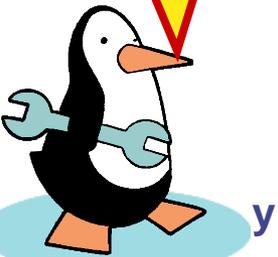
Confirmation of DNA = heredity molecule

■ Hershey & Chase (1952)

- ◆ classic “blender” experiment
- ◆ worked with **bacteriophage**
 - viruses that infect bacteria
- ◆ grew phage viruses in 2 media, radioactively labeled with either
 - ^{35}S in their proteins
 - ^{32}P in their DNA
- ◆ infected bacteria with labeled phages



Why use Sulfur vs. Phosphorus?



Hershey & Chase



Hershey & Chase

Protein coat labeled with ^{35}S

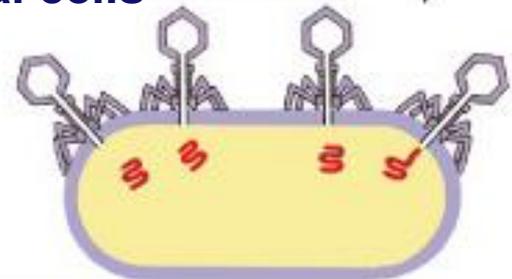
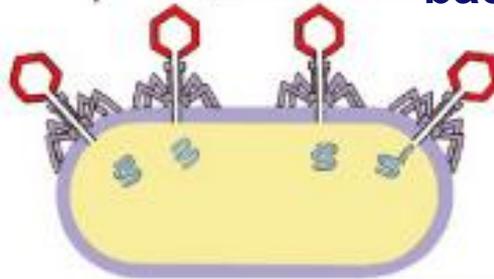


T2 bacteriophages are labeled with radioactive isotopes
S vs. P

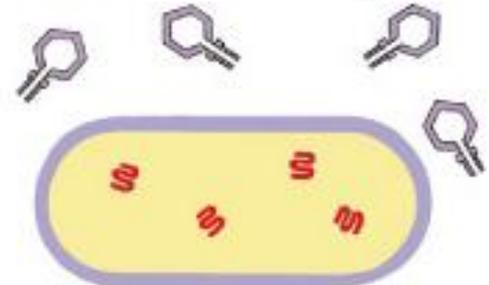
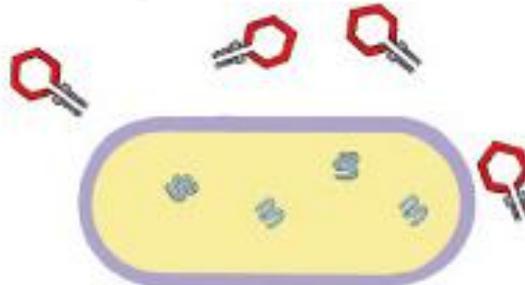
DNA labeled with ^{32}P



bacteriophages infect bacterial cells



bacterial cells are agitated to remove viral protein coats



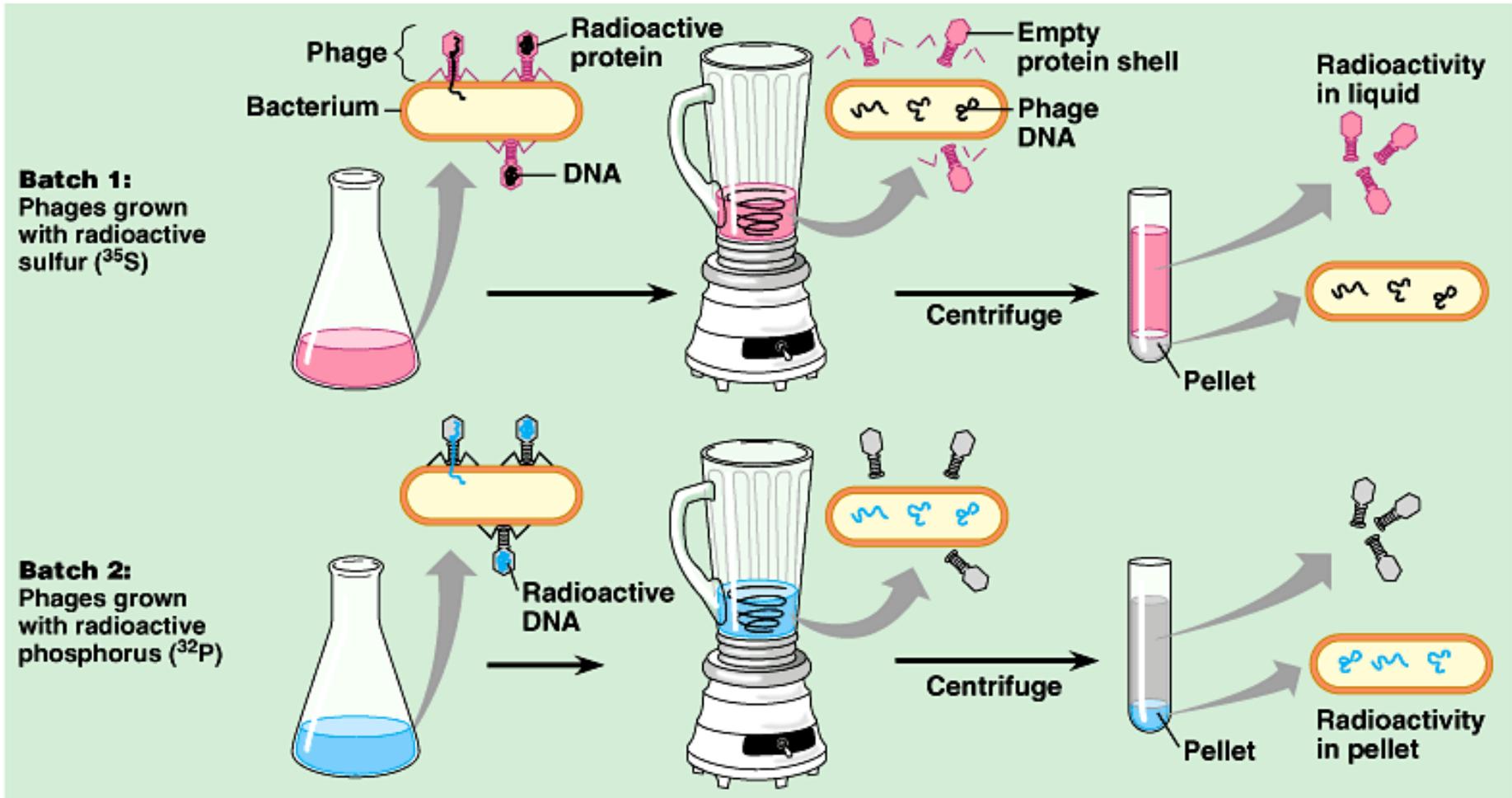
^{35}S radioactivity found in the medium

^{32}P radioactivity found in the bacterial cells

Which radioactive marker is found inside the cell?

Which molecule carries viral genetic info?

- 1 Mix radioactively labeled phages with bacteria. The phages infect the bacterial cells.
- 2 Agitate in a blender to separate phages outside the bacteria from the cells and their contents.
- 3 Centrifuge the mixture so bacteria form a pellet at the bottom of the test tube.
- 4 Measure the radioactivity in the pellet and the liquid.



Blender experiment conclusions

- Radioactive phage & bacteria in blender
 - ◆ ^{35}S phage = indicates protein
 - radioactive proteins stayed in upper liquid
 - therefore protein did NOT enter bacteria
 - ◆ ^{32}P phage = indicates DNA
 - radioactive DNA found in lower pellet
 - therefore DNA did enter bacteria
 - ◆ **Confirmed DNA is “transforming factor”**



Chargaff's data:

TABLE 11-2

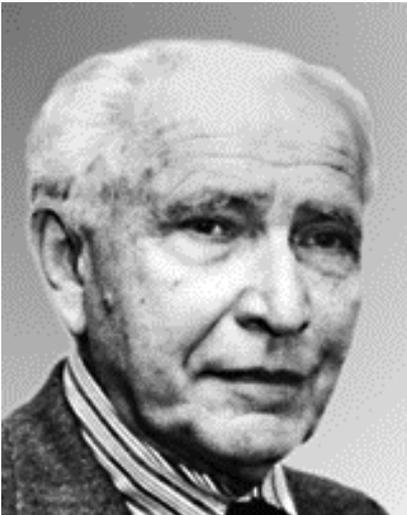
Base Compositions in DNA from Selected Organisms

Source of DNA	Percentage of DNA Bases				Ratios	
	A	T	G	C	A/T	G/C
<i>E. coli</i>	26.1	23.9	24.9	25.1	1.09	0.99
Yeast	31.3	32.9	18.7	17.1	0.95	1.09
Sea urchin sperm	32.5	31.8	17.5	18.2	1.02	0.96
Herring sperm	27.8	27.5	22.2	22.6	1.01	0.98
Human liver	30.3	30.3	19.5	19.9	1.00	0.98
Corn (<i>zea mays</i>)	25.6	25.3	24.5	24.6	1.01	1.00

1947

Chargaff (1947)

- DNA composition: “Chargaff’s rules”
 - ◆ varies from species to species
 - ◆ all 4 bases not in equal quantity
 - ◆ bases present in characteristic ratio
- humans:



Erwin Chargaff

A = 30.9%

T = 29.4%

G = 19.9%

C = 19.8%

What do
you notice?!



Chargaff's big clue: Base Pair RATIOS in DNA

■ Purines

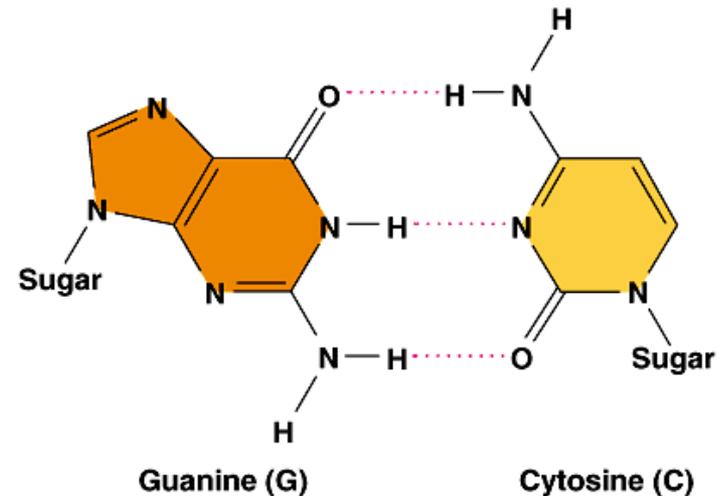
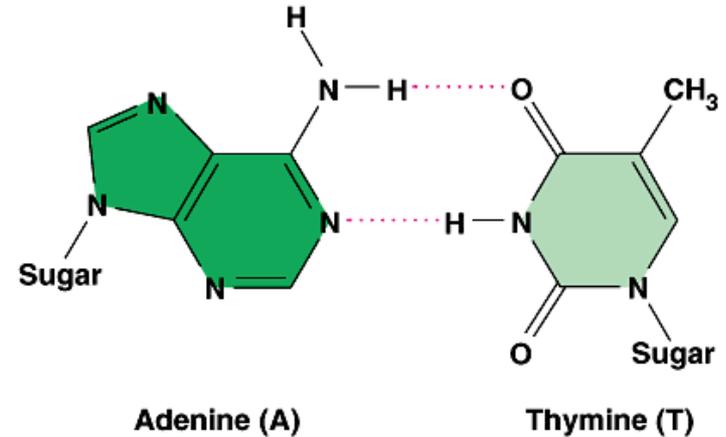
- ◆ adenine (A)
- ◆ guanine (G)

■ Pyrimidines

- ◆ thymine (T)
- ◆ cytosine (C)

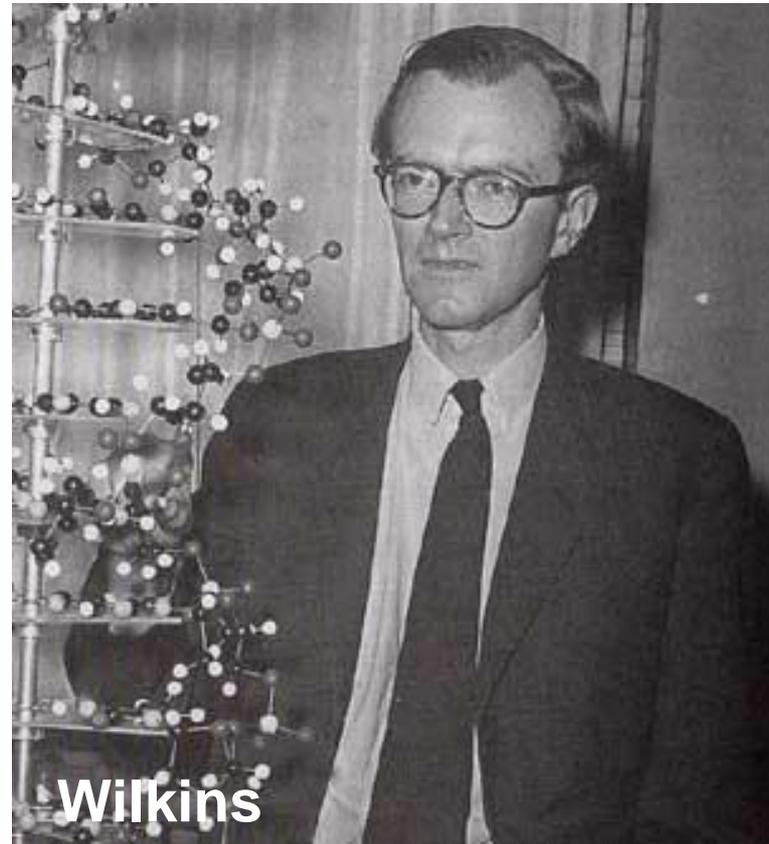
■ Pairing

- ◆ A : T
- ◆ C : G

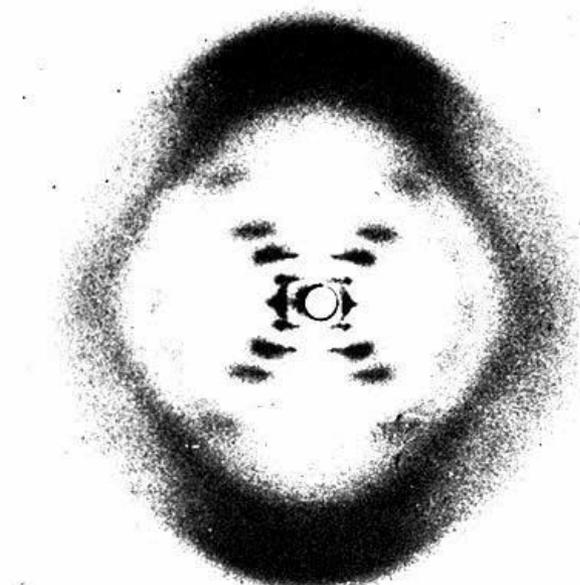
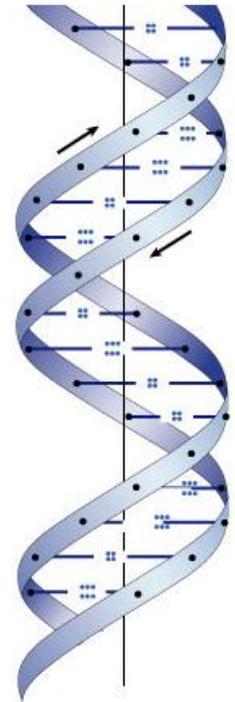
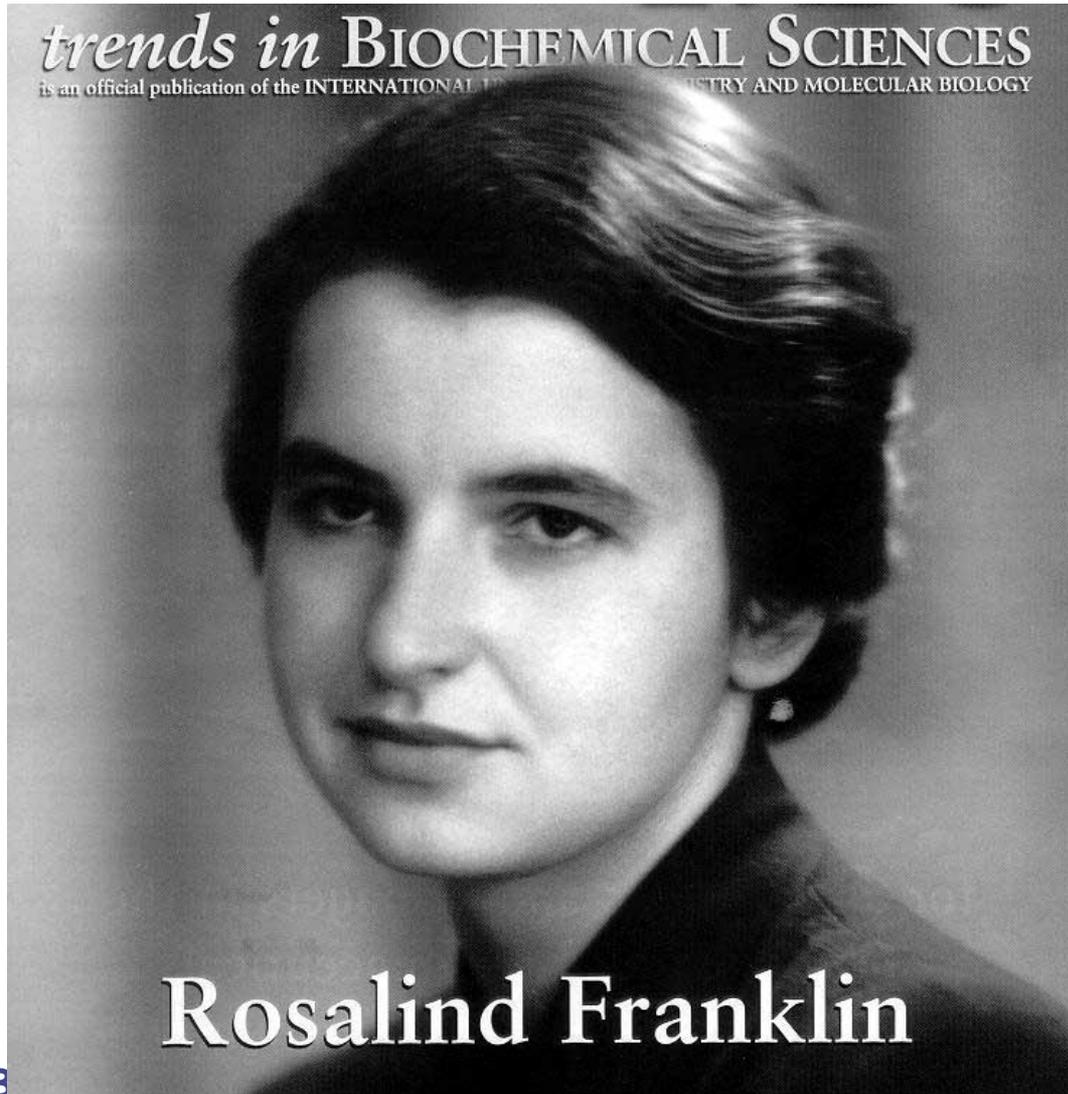


Another clue: Franklin & Wilkins (1953)

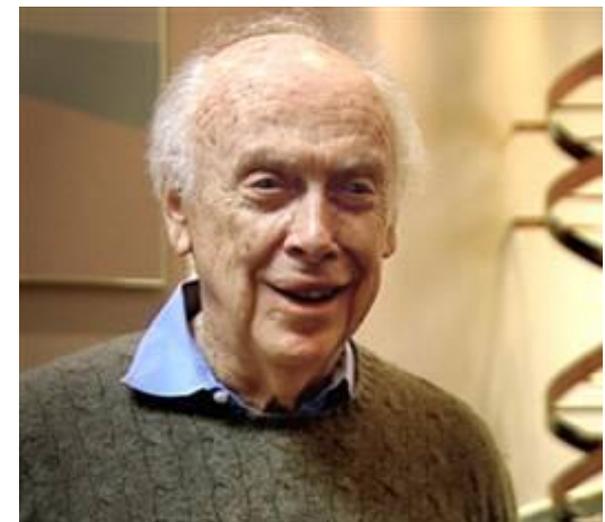
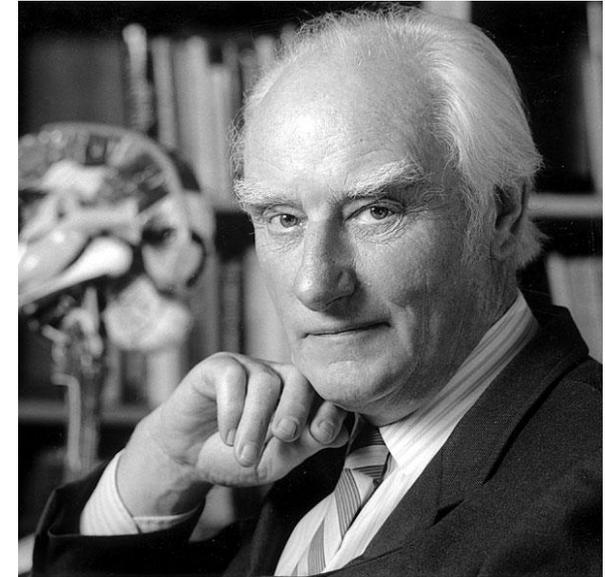
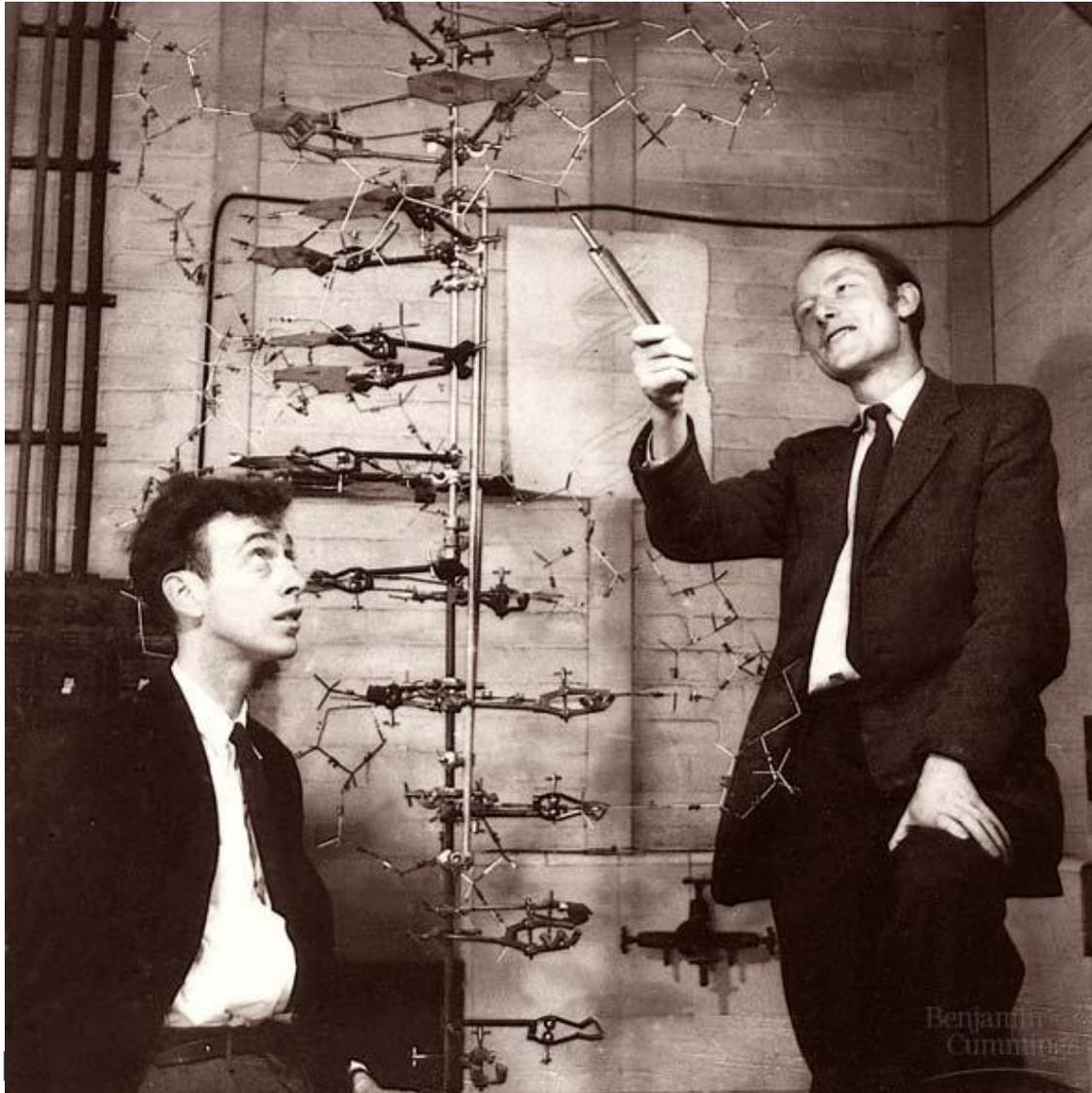
- ◆ X-ray pictures of DNA suggest that it consists of 2 chains twisted in a helix



Rosalind Franklin (1920-1958)

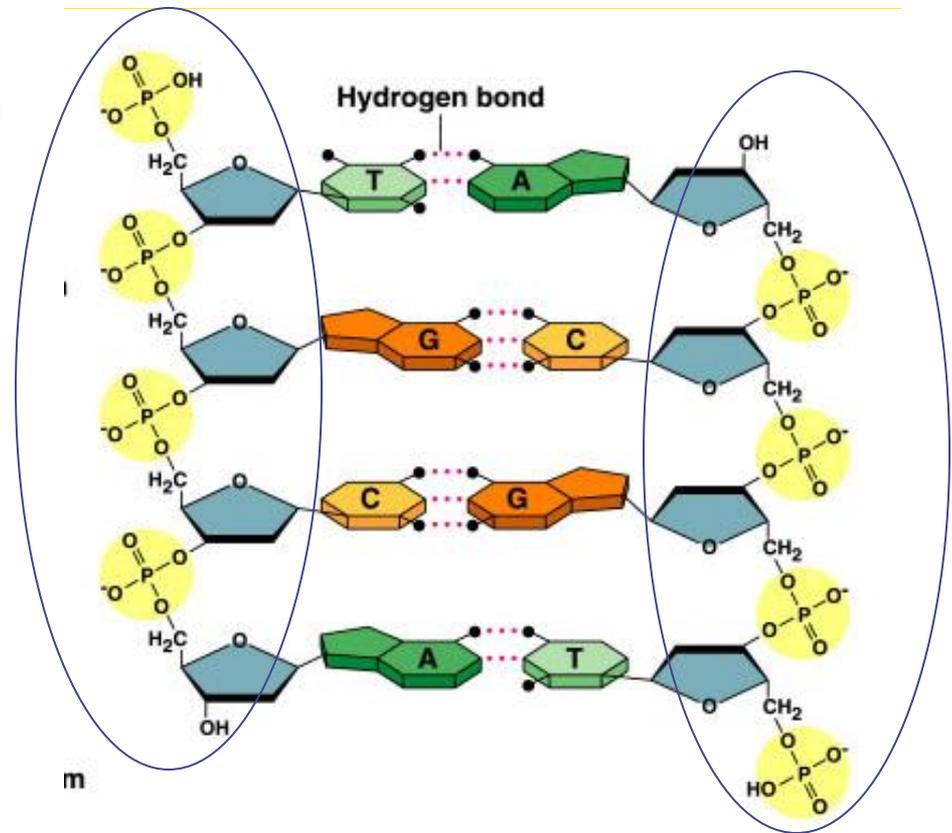
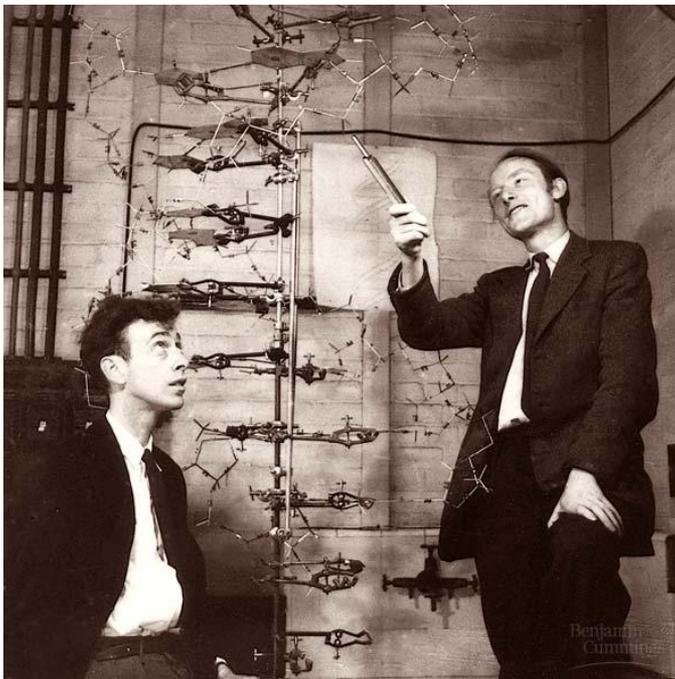


The “Fathers” of DNA Structure

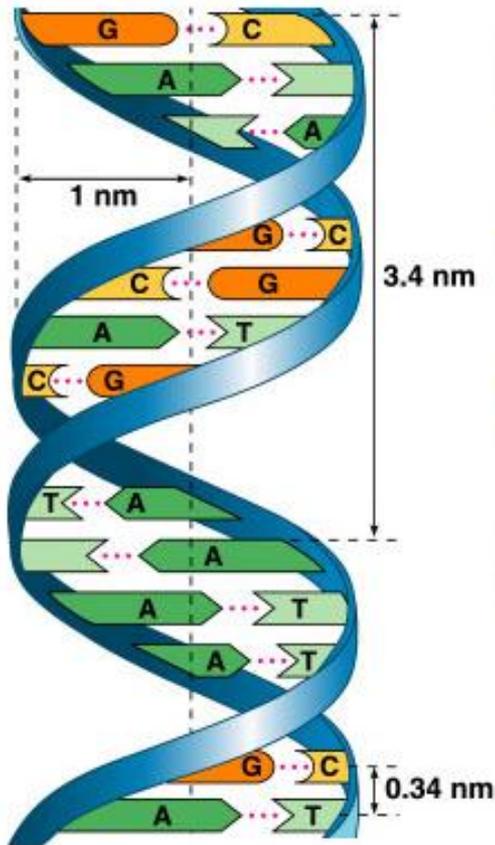


Watson and Crick (1953)

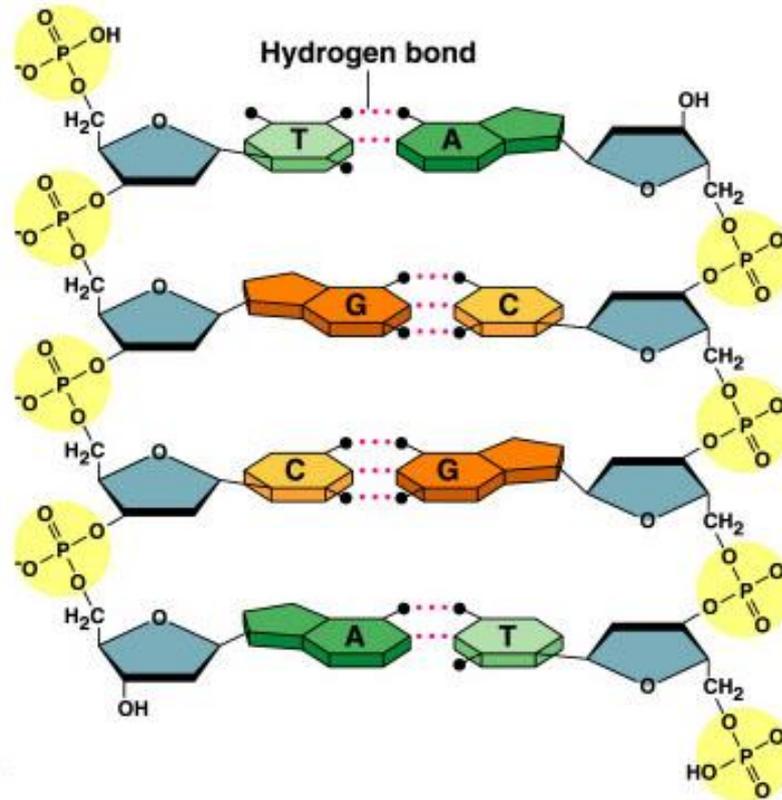
- Win the “race” to build the first accurate model of DNA
- Double Helix Model = 2 DNA chains that spiral like a “twisted rope ladder”
- **ladder sides** formed by alternating 5-C sugars & phosphates = **backbone**
- **ladder rungs** formed by nitrogen base pairs held together by **hydrogen** bonds
 - A – T (**2 H bonds**)
 - C – G (**3 H bonds**)



Double helix structure of DNA



(a) Key features of DNA structure



(b) Partial chemical structure



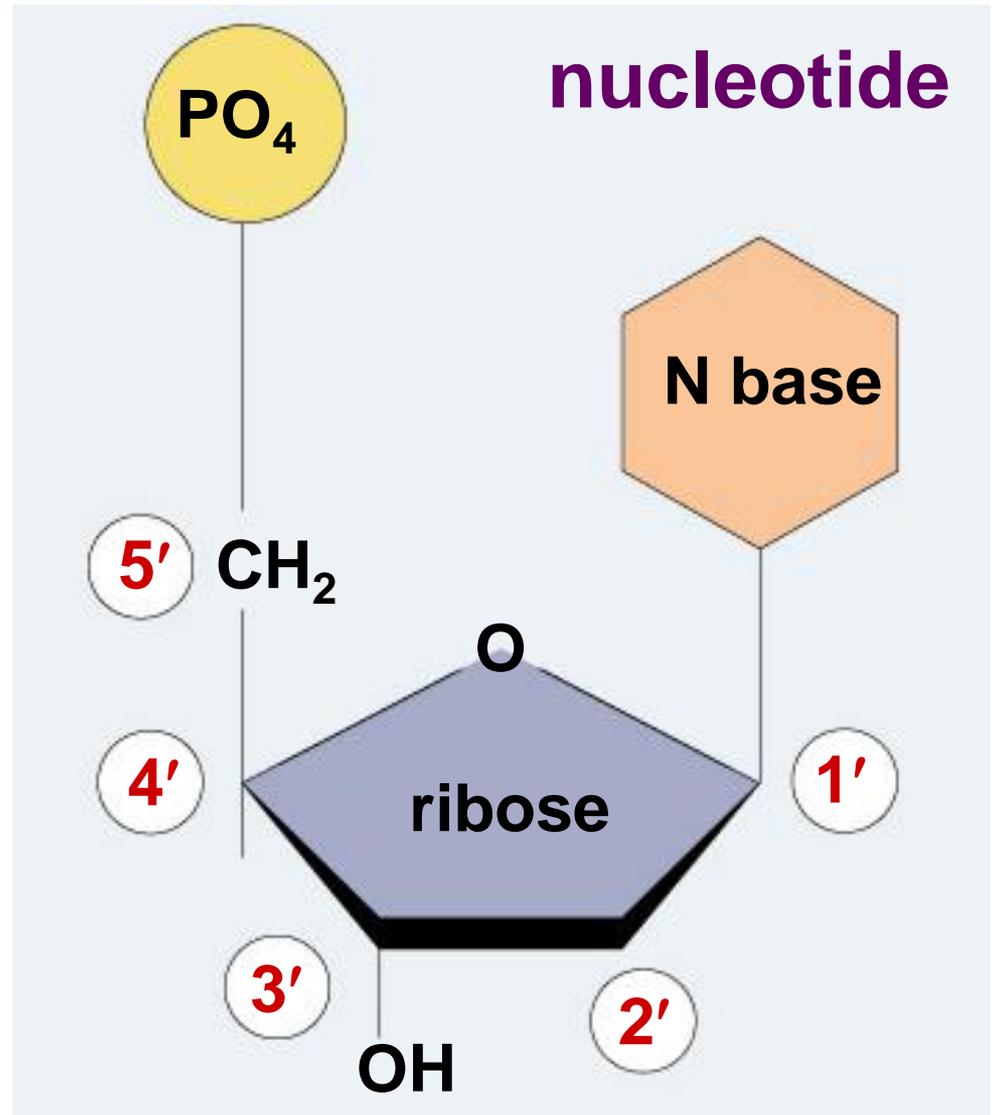
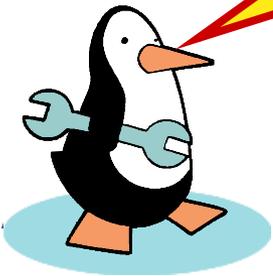
(c) Space-filling model

the structure of DNA suggested a mechanism for **how DNA is copied** by the cell

Directionality of DNA

- You need to number the carbons!
 - ◆ it matters!

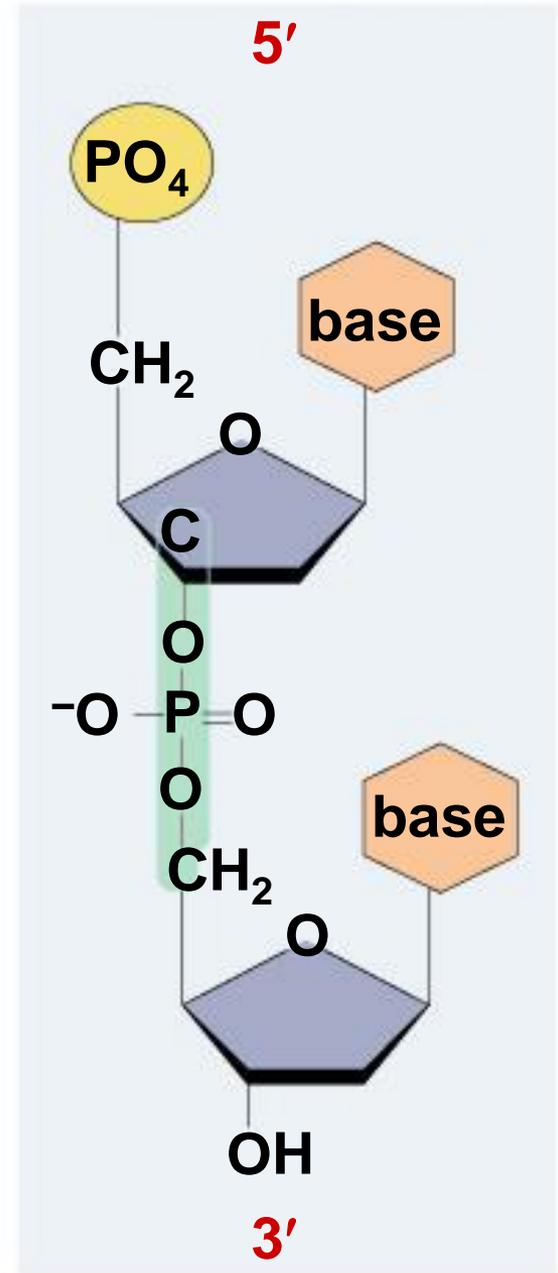
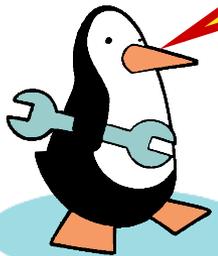
This will be IMPORTANT!!



The DNA backbone

- Putting the DNA backbone together
 - ◆ refer to the 3' and 5' ends of the DNA

I mean it...
This will be
IMPORTANT!!



- Each strand of the double helix is oriented in the opposite direction = Antiparallel

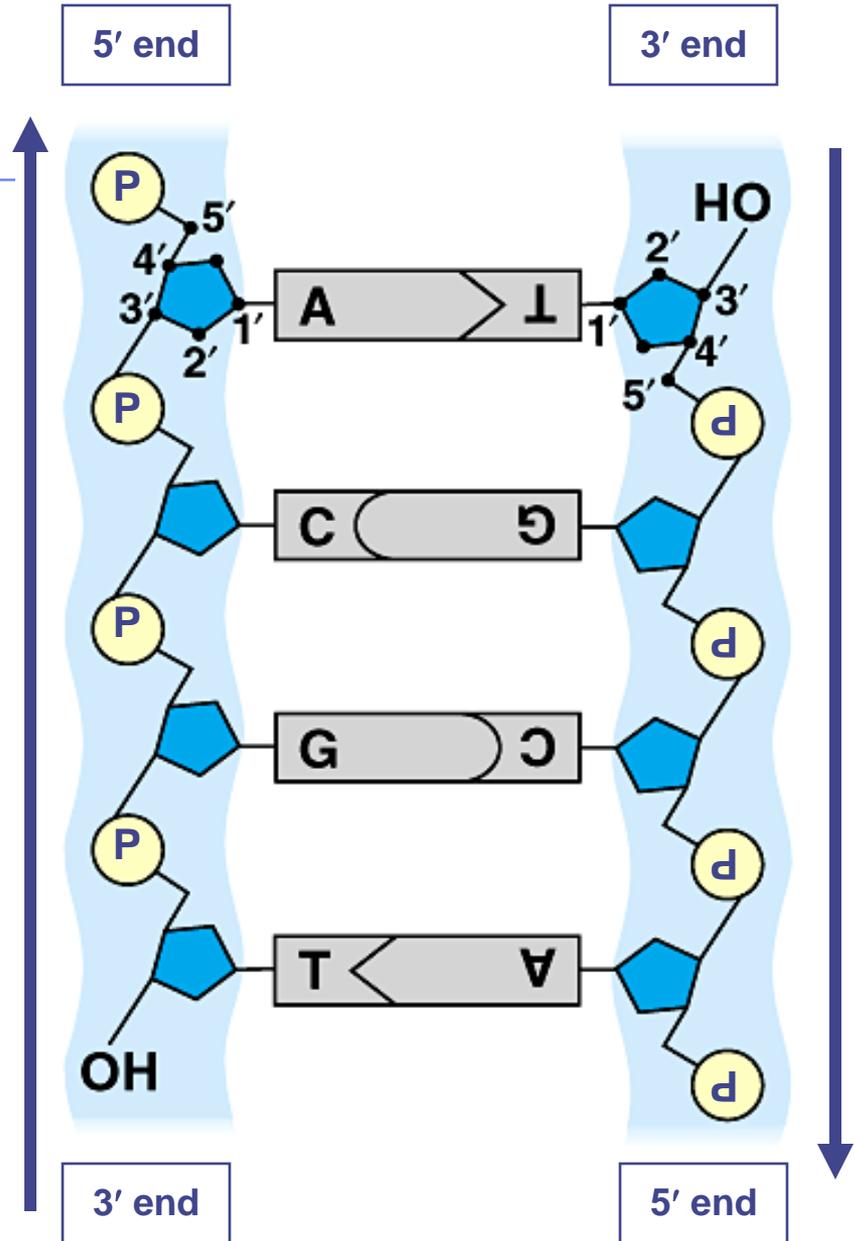
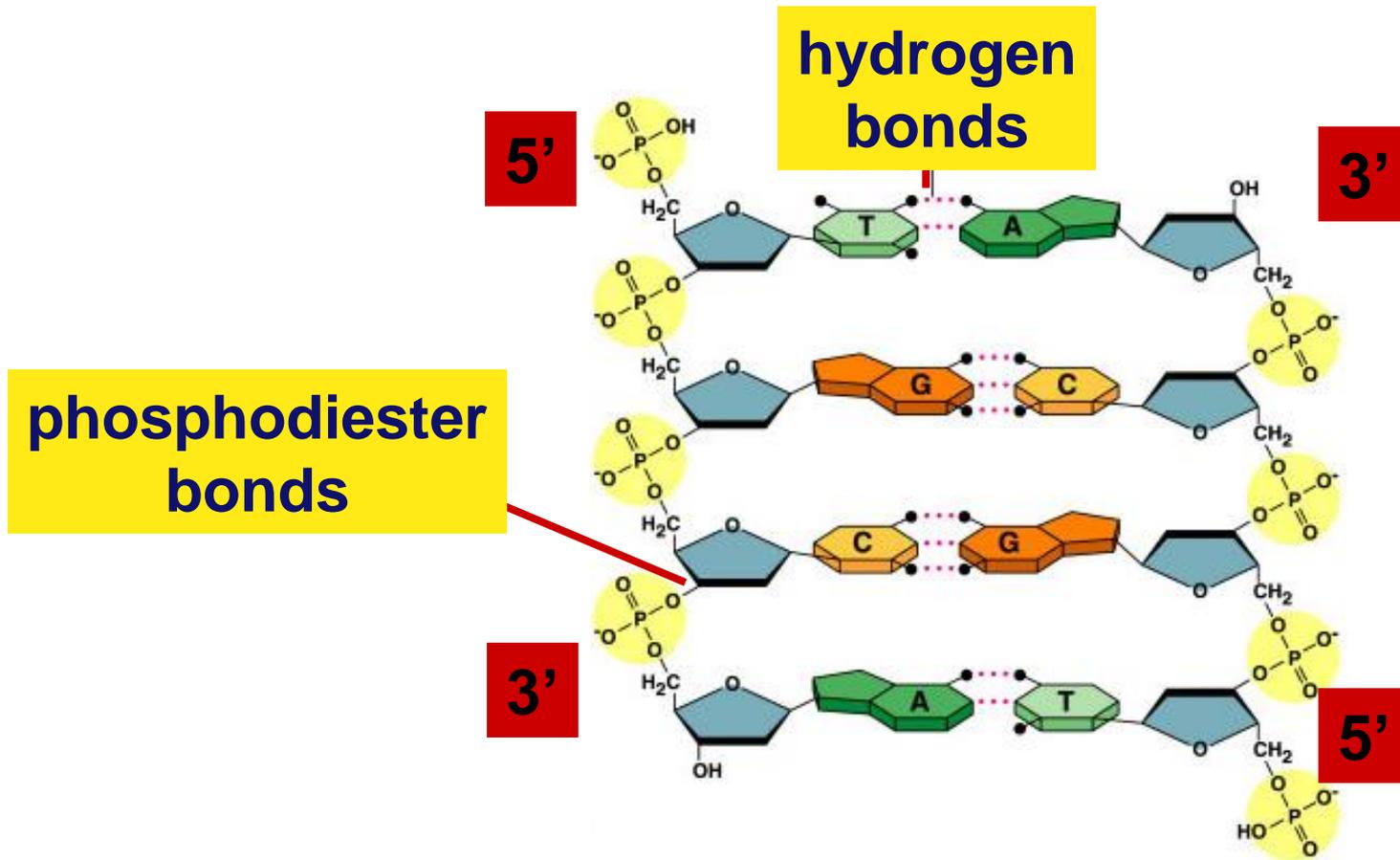


Figure 10.5B

Bonding in DNA



....strong or weak bonds?

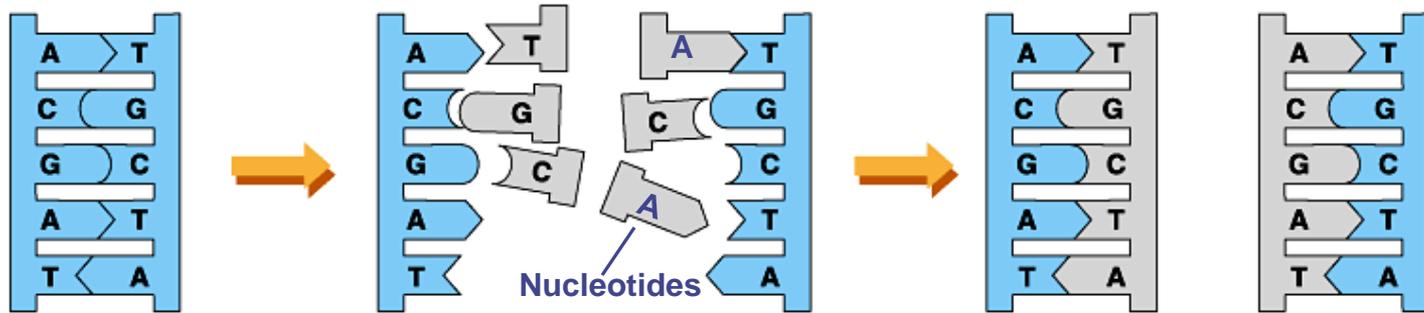
How do the bonds fit the mechanism for copying DNA?

DNA REPLICATION

- **WHAT ?** → a process that produces two DNA molecules identical to the original.
- **WHY?** → so when cells divide they can give each daughter cell its own identical copy of the DNA blueprint
- **WHEN?** → right BEFORE cell division as new body cells are made:
 - During growth
 - Repairing injury
 - Replacing old cells that wear out
 - Producing eggs or sperm (gametes)

DNA REPLICATION: HOW?

- 1) DNA Helicase enzymes bind to the DNA at different locations and then unwind and unzip the 2 strands
- 2) DNA Polymerase enzymes use each old strand as a template to assemble the new strand
- 3) DNA Ligase enzymes link together DNA fragments until long, continuous strands are formed

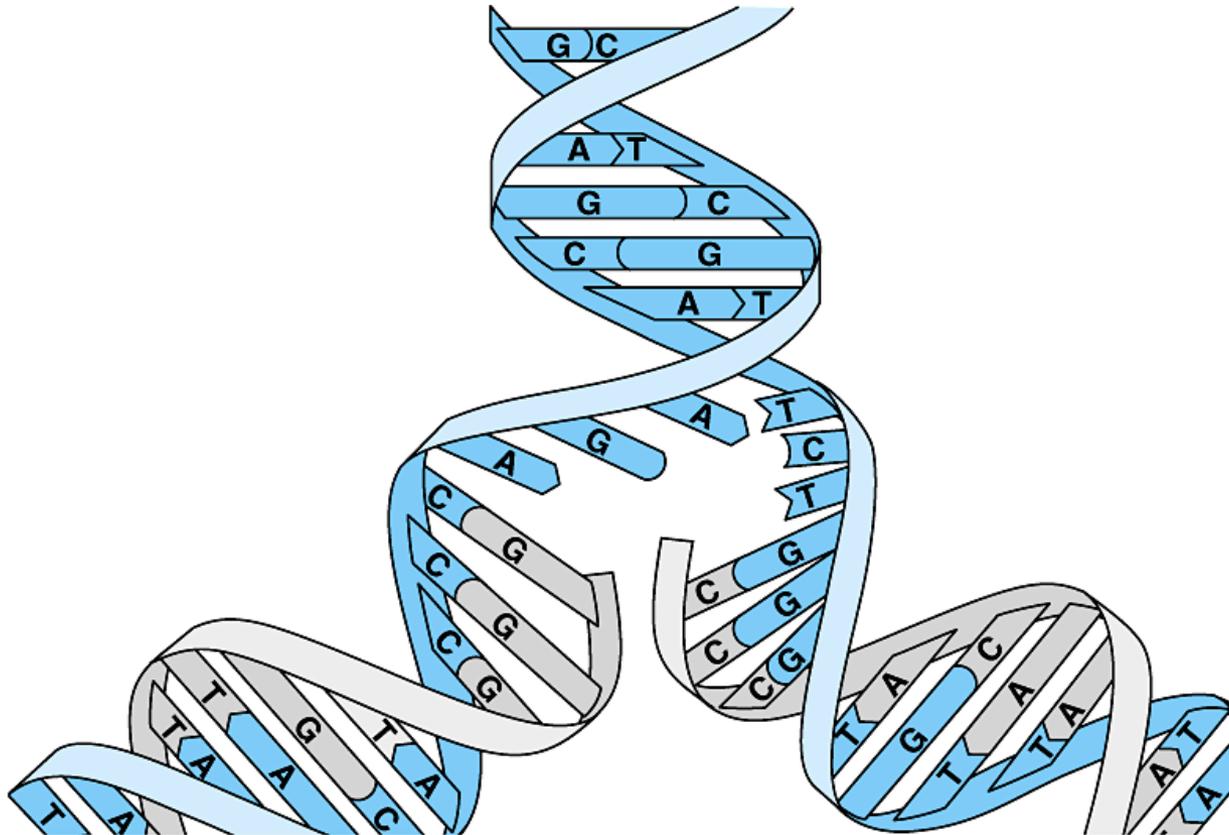


Parental molecule
of DNA

Both parental strands serve
as templates

Two identical daughter
molecules of DNA

■ Untwisting and replication of DNA

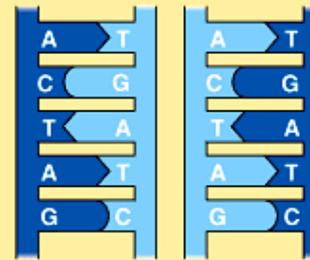
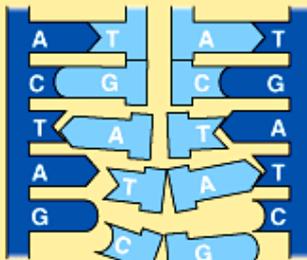
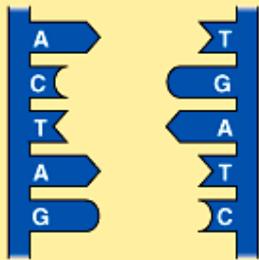
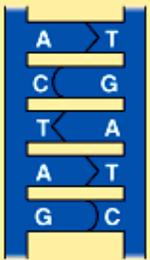
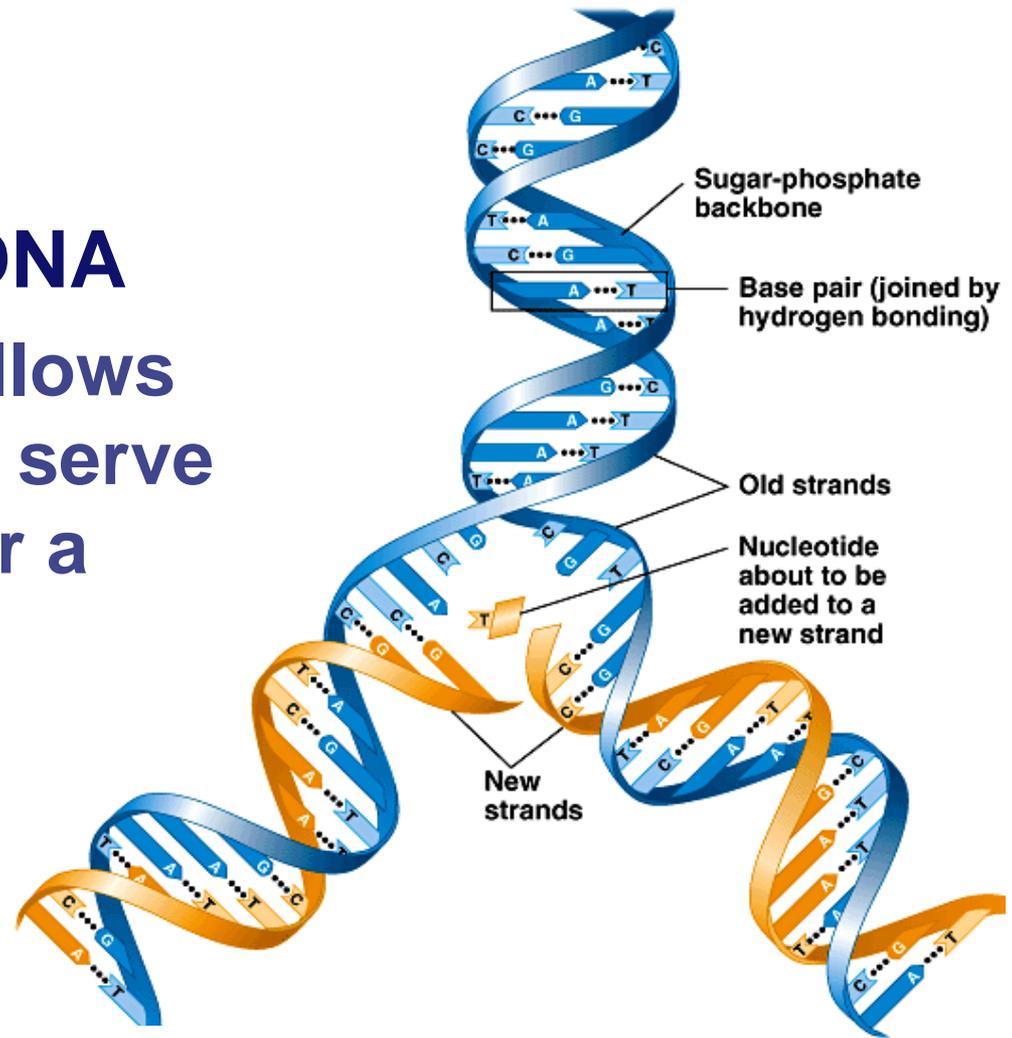


- DNA replication is a semi-conservative process = 1 strand is old and 1 strand is new

Figure 10.4B

Copying DNA

- **Replication of DNA**
 - ◆ base pairing allows each strand to serve as a pattern for a new strand



10.5 DNA replication: A closer look

- DNA replication begins at many different sites, so... the whole molecule can be copied much more quickly

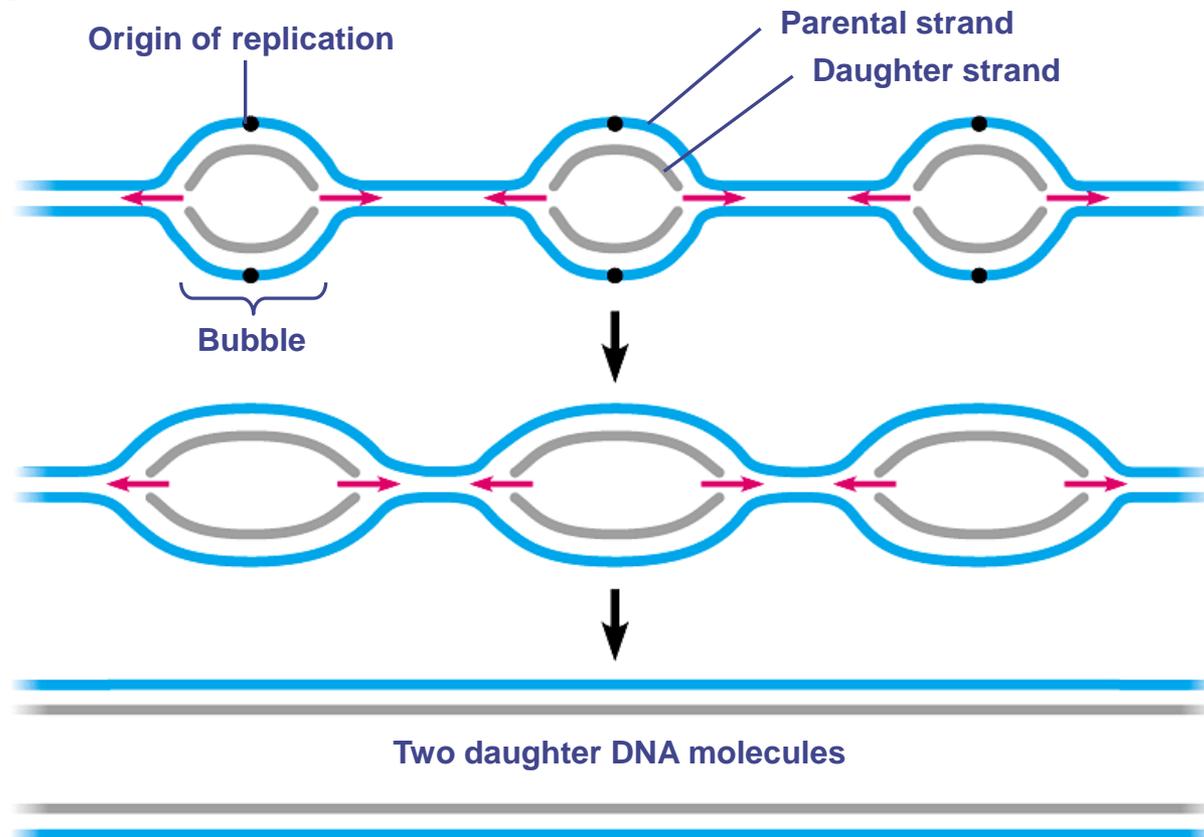


Figure 10.5A

- Because DNA is Antiparallel and Polymerase only builds new DNA in ONE direction ($5' \rightarrow 3'$), this affects how DNA daughter strands are synthesized:
 - 1 side continuously
 - 1 side in fragments
- The fragments on one side are linked together by Ligase enzymes.

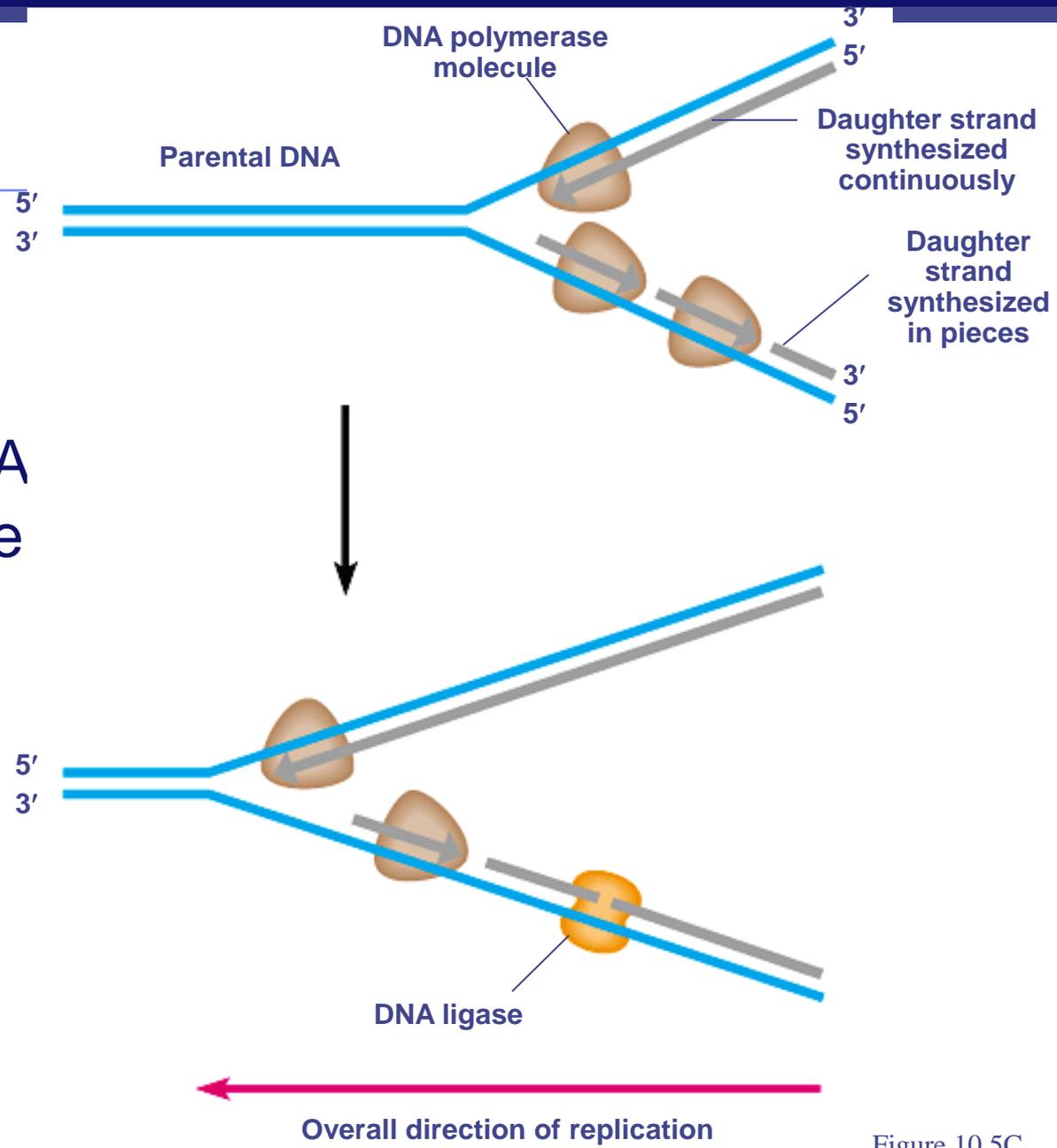
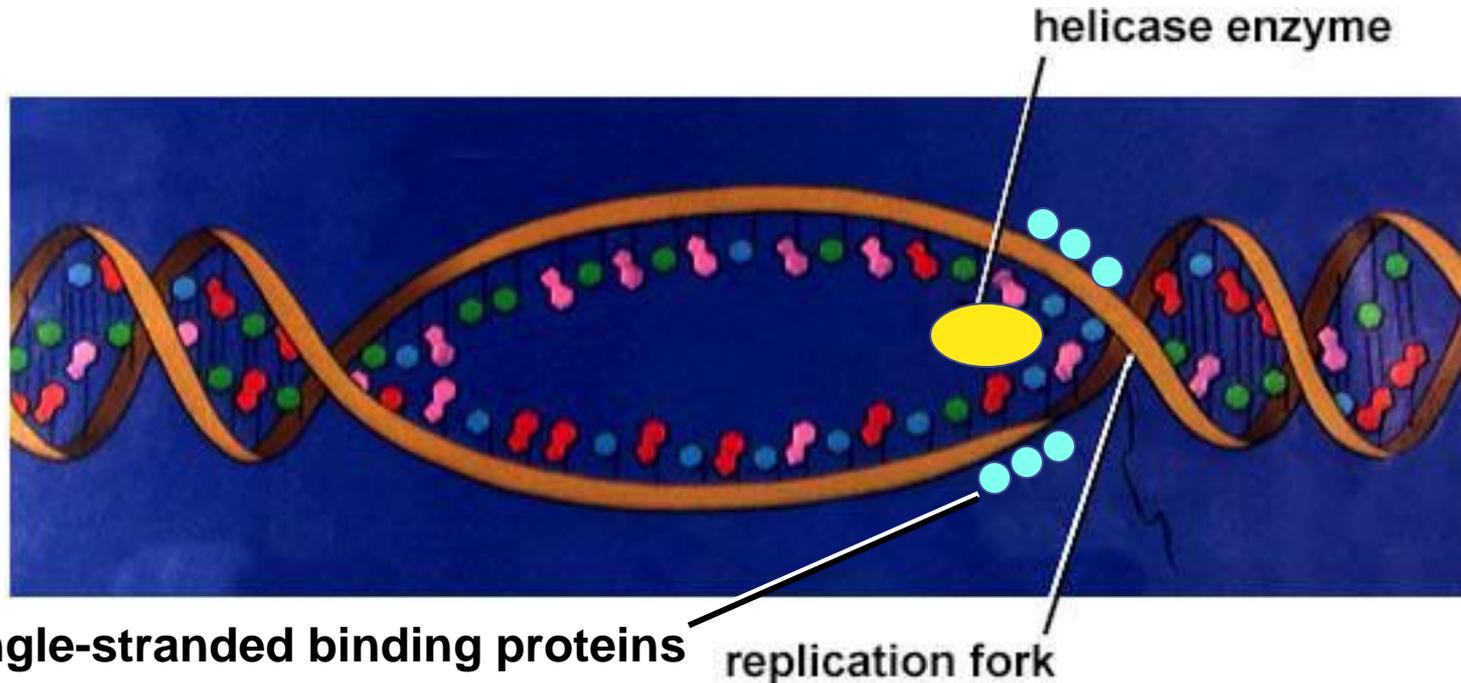


Figure 10.5C

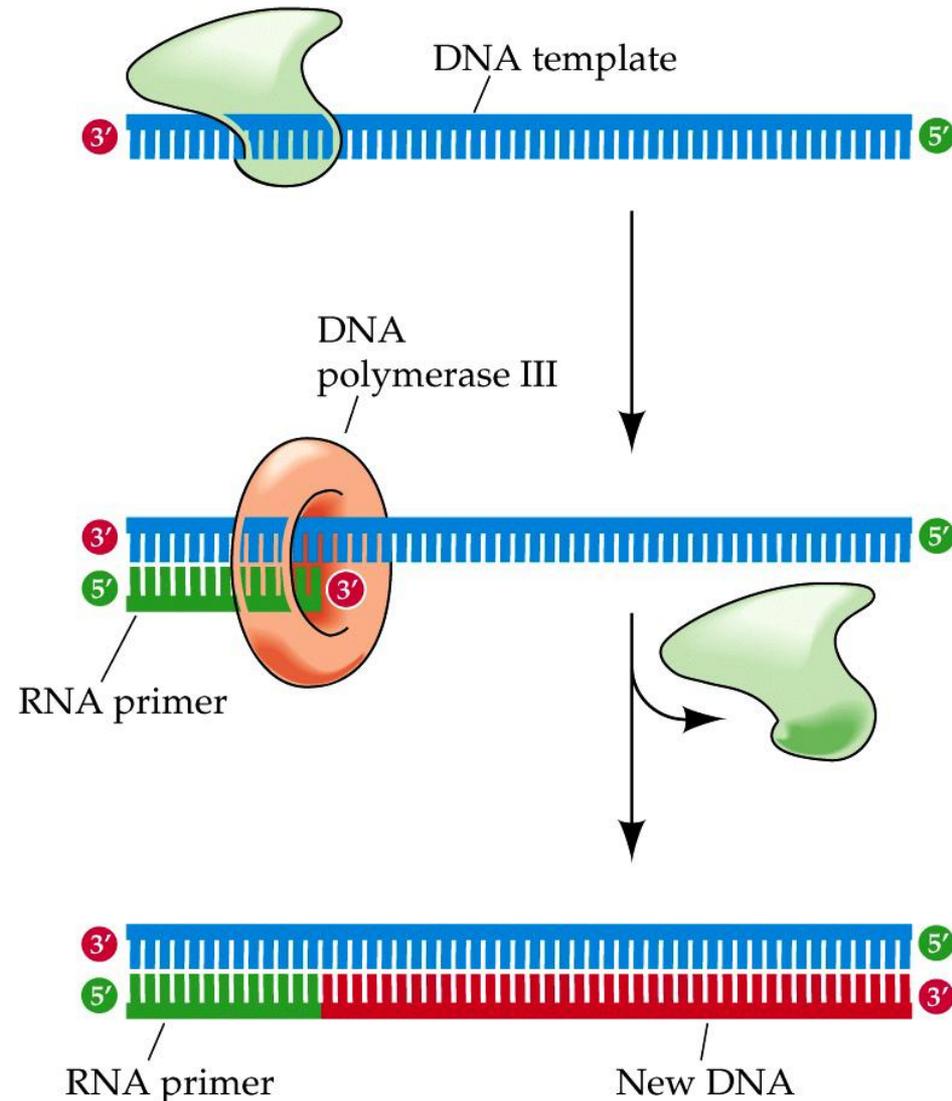
Replication: 1st step

- Helicase unwinds & unzips the 2 strands
 - ◆ stabilized by single-stranded binding proteins



Step 2: Priming DNA synthesis

- DNA polymerase I can only attach a new nucleotide on a **FREE 3' end**
 - ◆ cannot start building a new DNA strand without a free 3' end
 - ◆ short **RNA primer** with a free 3' end is built first by **primase**

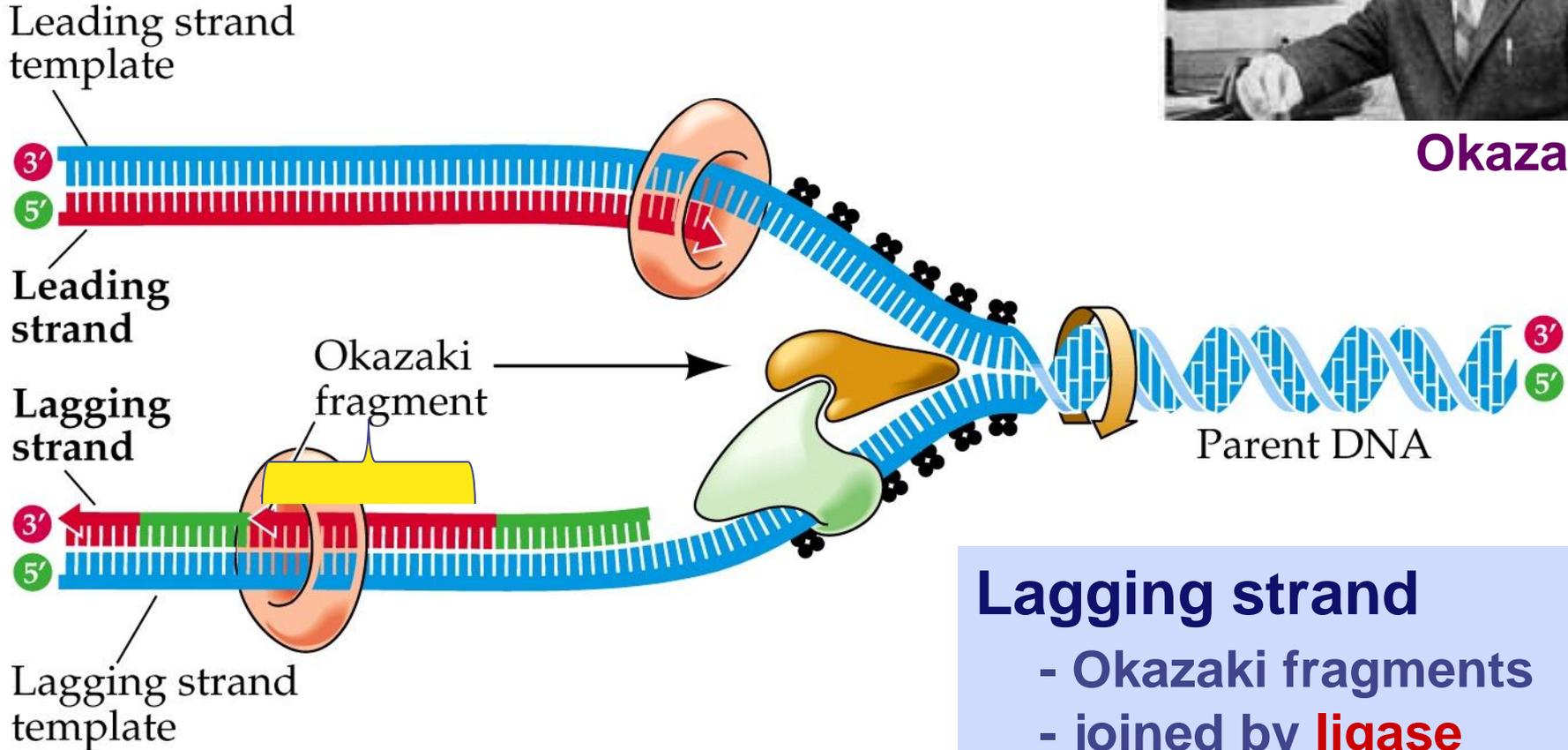


Step 3: Polymerase I builds Leading & Lagging strands

Leading strand
- continuous synthesis

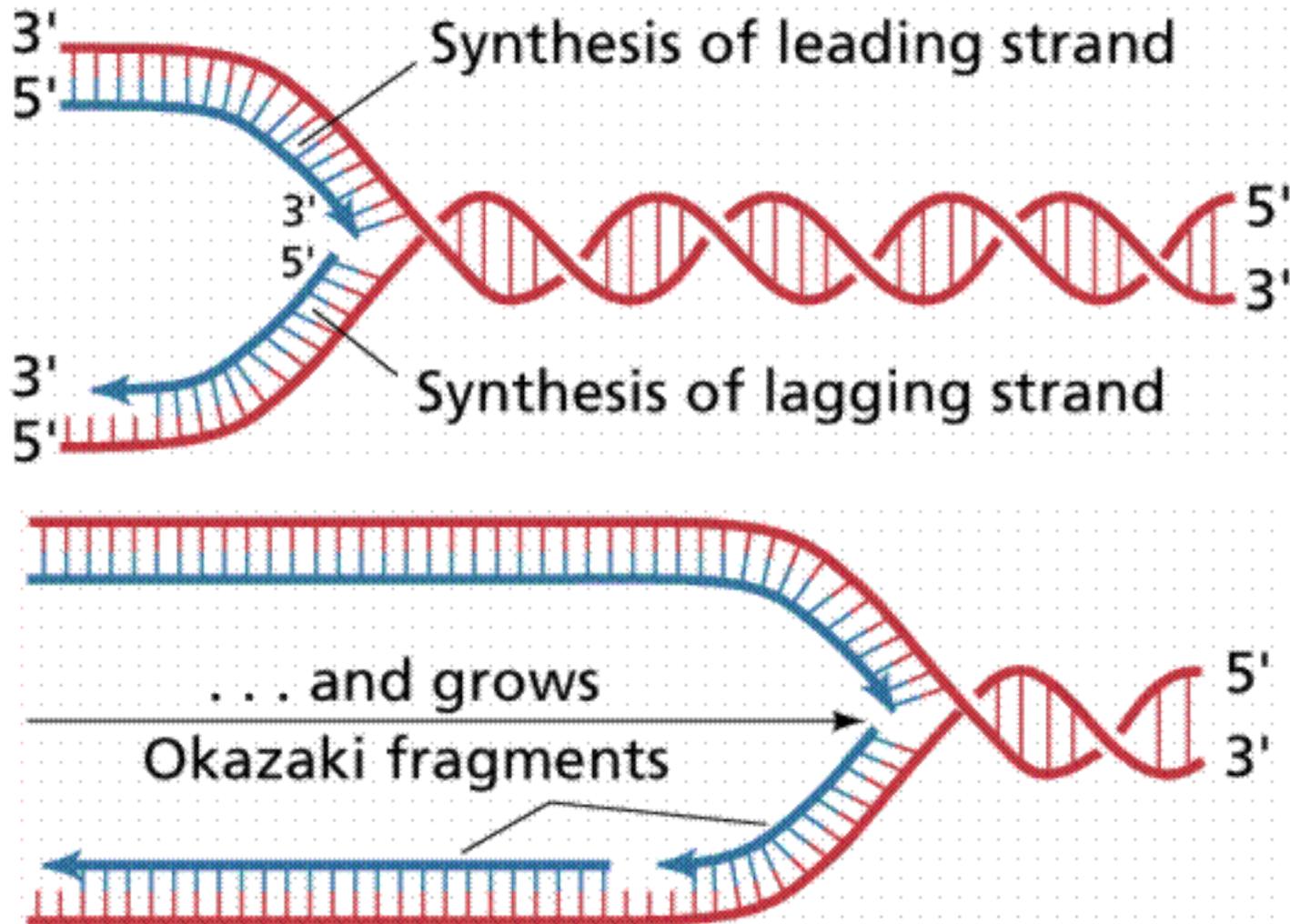


Okazaki



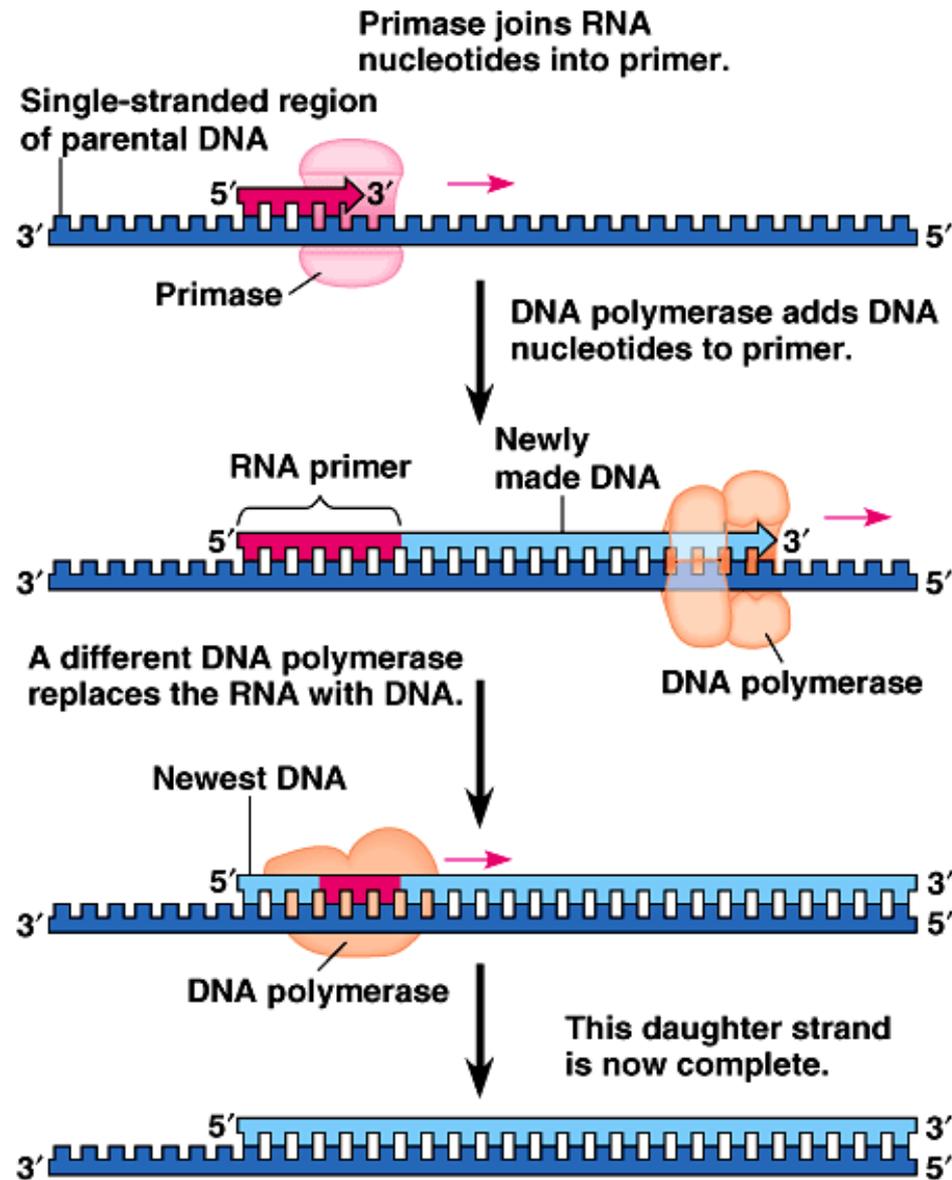
Lagging strand
- Okazaki fragments
- joined by **ligase**
- “spot welder” enzyme

Okazaki fragments

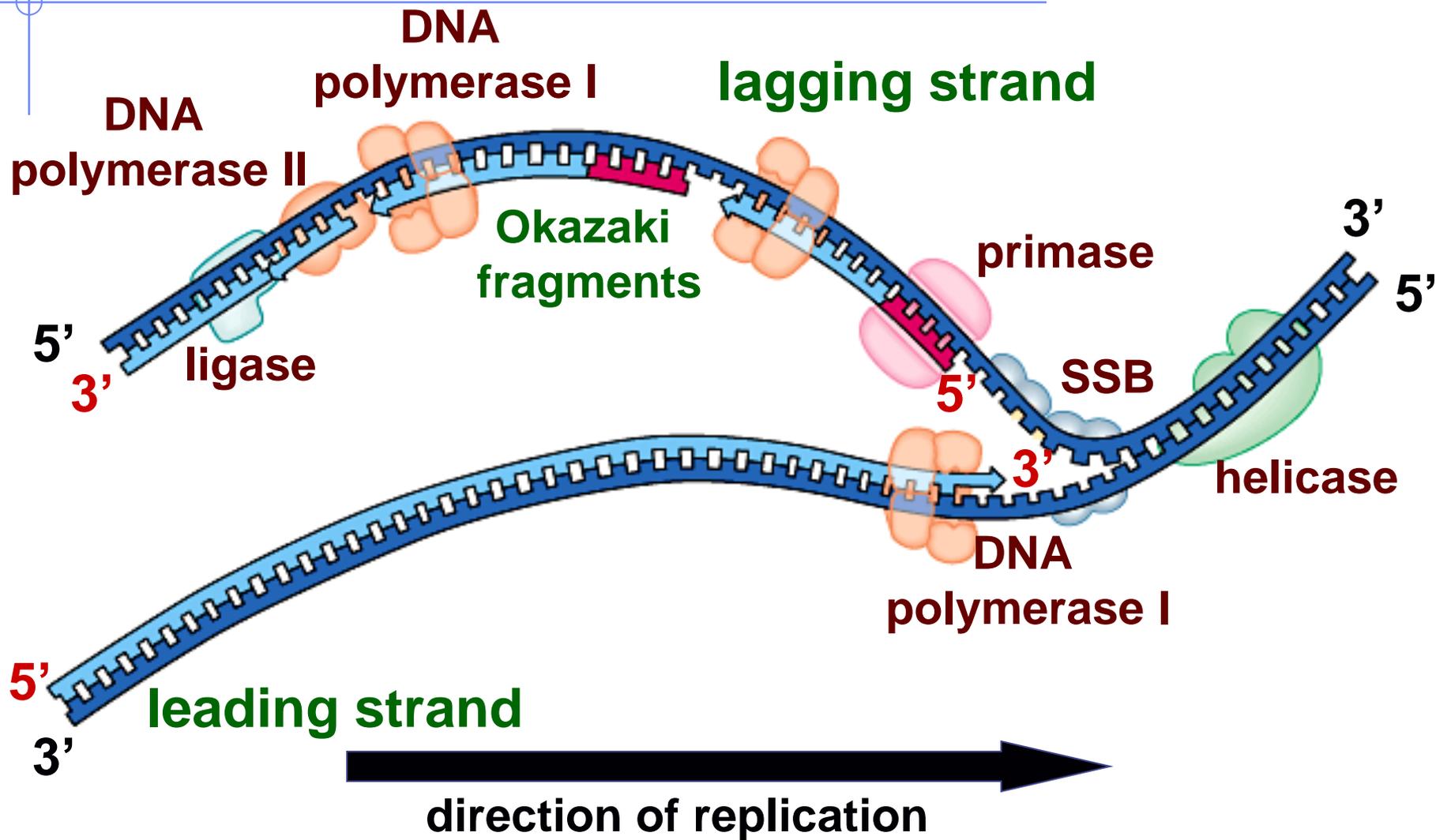


Step 4: Polymerase II cleans up primers

DNA polymerase II removes sections of RNA primer and replaces with DNA nucleotides



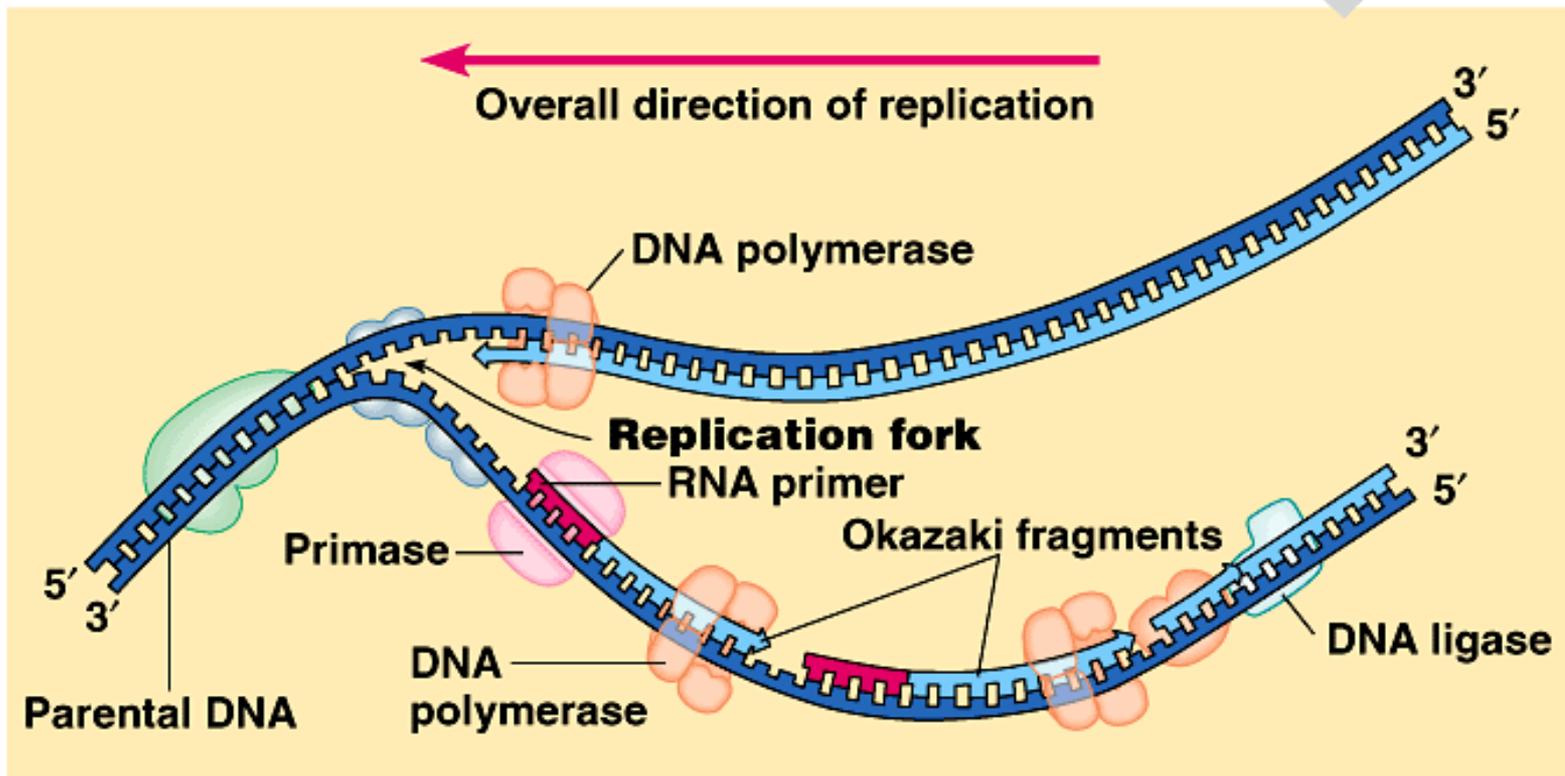
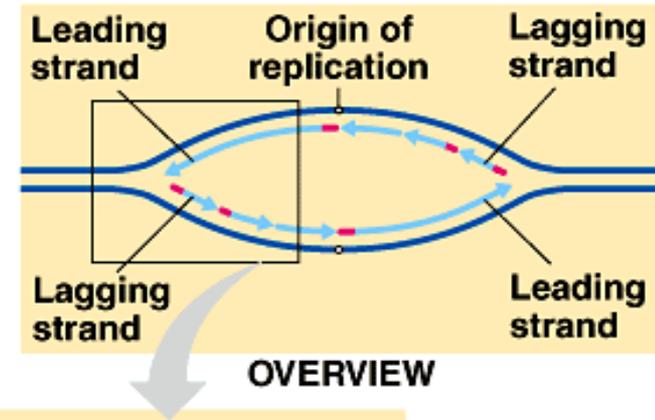
Step 5: Ligase links all DNA fragments



Replication bubble

Adds 1000 bases/second!

- Which direction does DNA build?
- List the enzymes & their role



Replication Speed & Accuracy

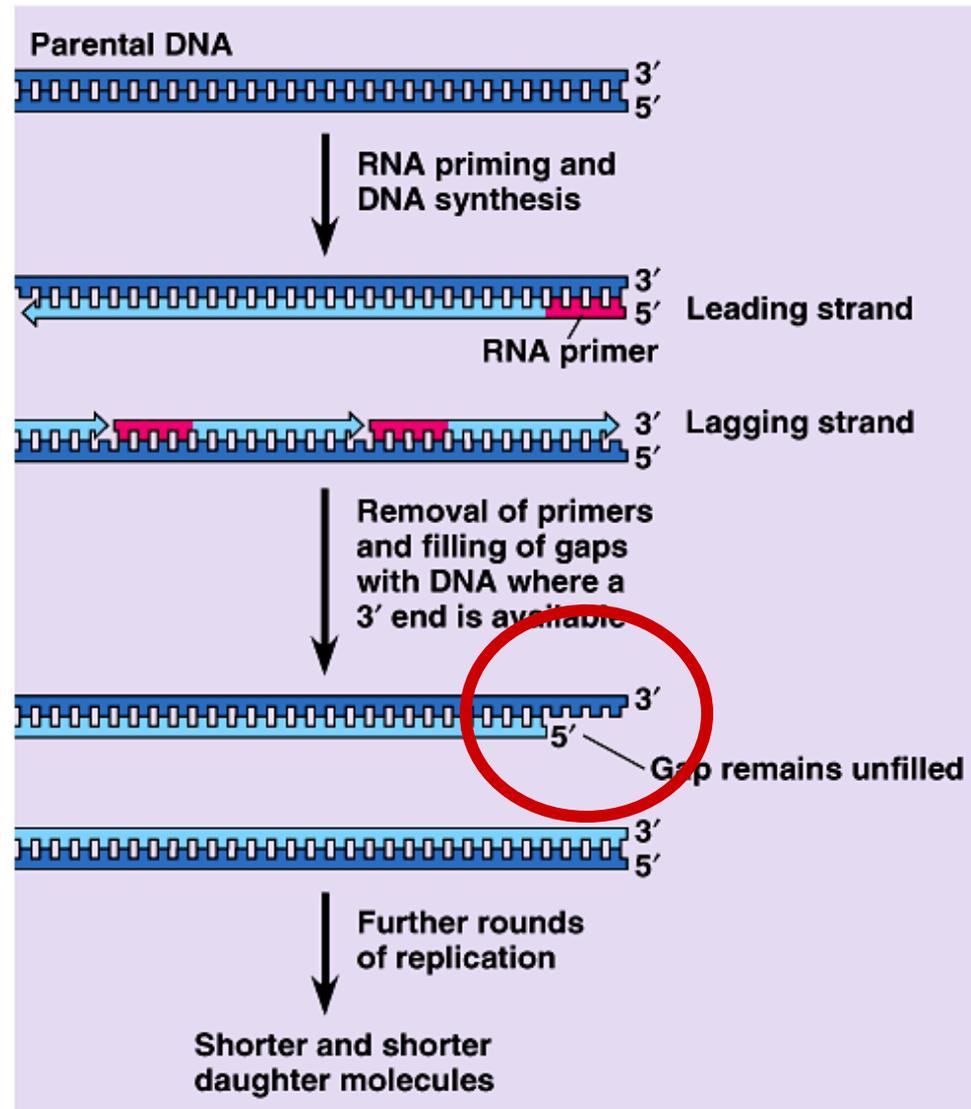
- Because the DNA molecule is copied so quickly (~50 new letters per second in mammals), occasionally the wrong letters are paired together
 - ◆ Most of these mismatch errors are fixed by the Polymerase enzyme as it backs up and inserts the correct matching base.
 - ◆ However, 1 mistake or mismatch every 1 billion letters does not get fixed and remains to be copied
- An army of repair enzymes constantly proofread the new DNA strands and fix most (99.9%) mistakes.
- Any unfixed DNA mistake = mutation

Replication enzymes

- *helicase*
- *DNA polymerase I*
- *primase*
- *DNA polymerase II*
- *ligase*
- *Repair Enzymes*

And in the end...

- Ends of chromosomes are eroded with each replication
 - ◆ an issue in aging?
 - ◆ ends of chromosomes are protected by telomeres



Telomeres

- Expendable, non-coding sequences at ends of DNA

- ◆ short sequence of bases repeated 1000s times

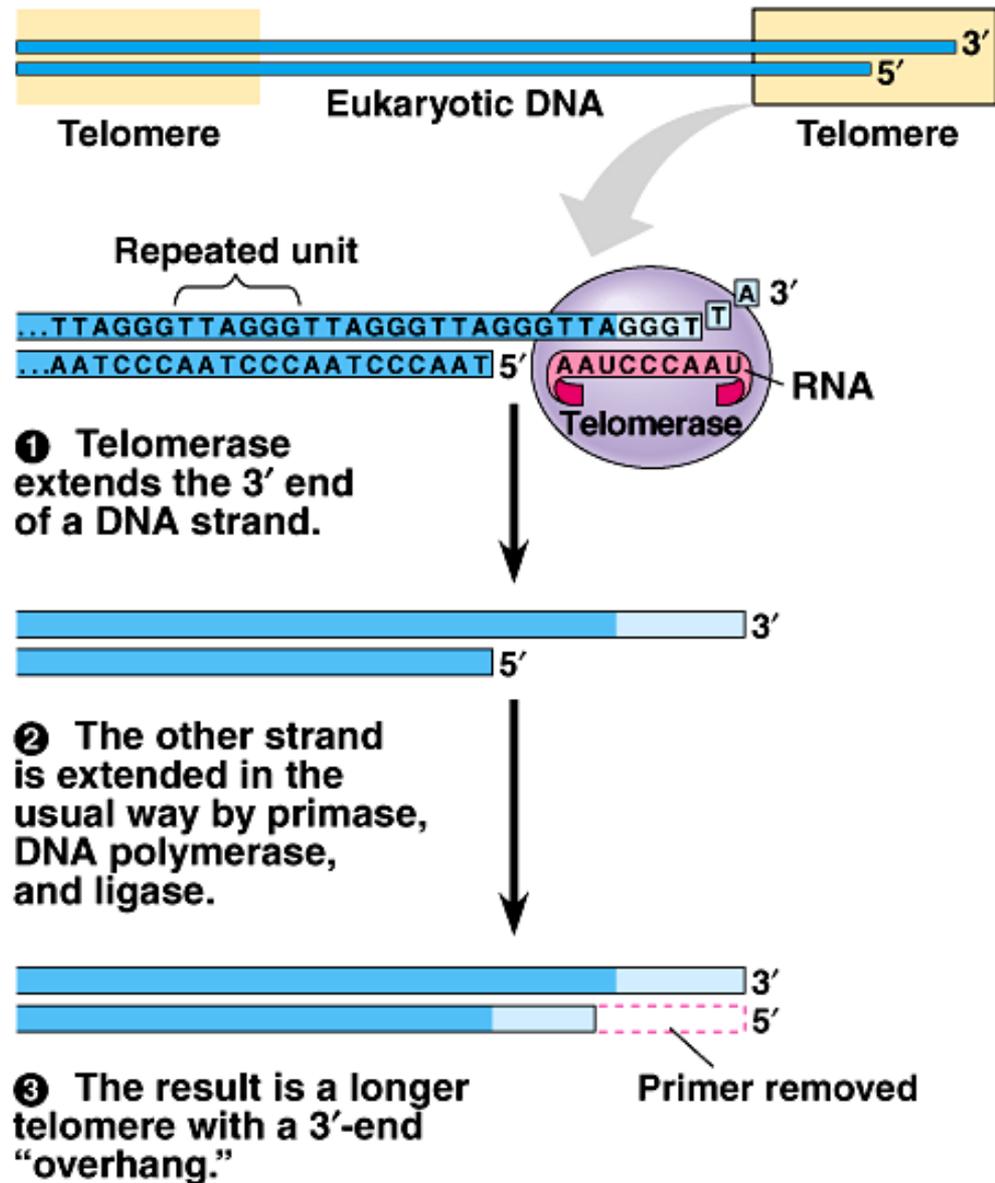
- ◆ TTAGGG in humans

- Telomerase enzyme in certain cells

- ◆ enzyme extends telomeres

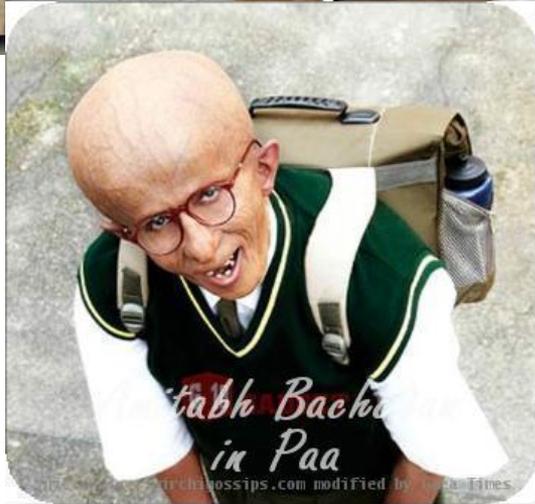
- ◆ prevalent in cancers

- Why?

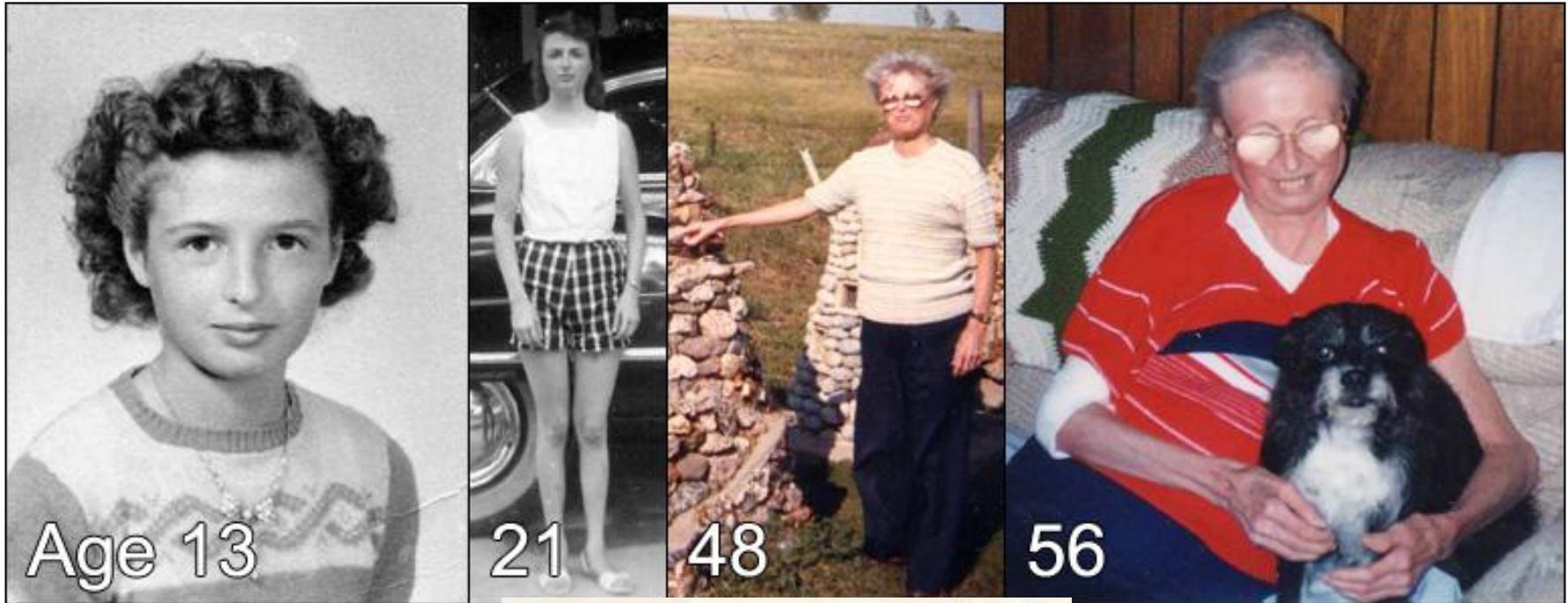


(b)

Progeria: One type of premature aging



Werner Syndrome: another type of premature again



Taking its toll. As a teenager (left) this Japanese American looked normal, but by age 48, the effects of Werner's syndrome were readily apparent. [Image credit: William and Wilkens Publishing Inc.]

Measuring Antioxidant levels in the skin



Point of Senescence
(SOX)

