DNA Form & Function AT--CG-CG CG GC-Original -TA TA. cell Daughter AT cells -CG-Stages of the cell cycle M = mitosis TA Μ **S** = **DNA** synthesis G = gap The two strands of the parental double helix A unwind, and each -GC--AT-G2 **G1** specifies a new daughter strand ATby base-pairing TA rules S -GC--TA-TA TA GC-GC--TA-GC Old TA

-TA--AT--GC-

New

-TA

DNA: Structure & Replication

Understanding DNA replication – and the resulting transmission of genetic information from cell to cell, and generation to generation – lays the groundwork for understanding the principles of heredity



Understanding DNA structure and replication is a prerequisite for understanding/using the principal tools of molecular biology

DNA: Structure & Replication

Three features of DNA makes it an ideal genetic material

1. Faithful replication

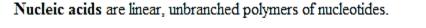
2. Information content

3. Capable of change

DNA: Nucleotides

Overall structure

Nucleotides



Nucleotides consist of three parts:

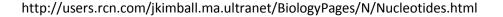
1.

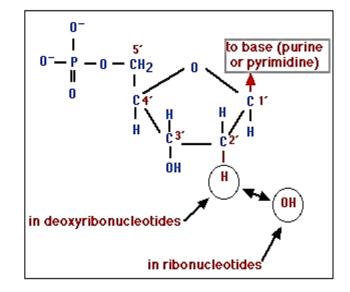
A five-carbon sugar (hence a pentose). Two kinds are found:

- Deoxyribose, which has a hydrogen atom attached to its #2 carbon atom (designated 2'), and
- Ribose, which has a hydroxyl group there.

Deoxyribose-containing nucleotides, the **deoxyribonucleotides**, are the monomers of deoxyribonucleic acids (**DNA**).

Ribose-containing nucleotides, the ribonucleotides, are the monomers of ribonucleic acids (RNA).





DNA: Nucleotides

Bases

2.

A nitrogen-containing ring structure called a base. The base is attached to the 1' carbon atom of the pentose. In DNA, four different bases are found:

- 1. two purines, called adenine (A) and guanine (G)
- 2. two pyrimidines, called thymine (T) and cytosine (C)

RNA contains:

- 1. The same purines, adenine (A) and guanine (G).
- 2. RNA also uses the pyrimidine cytosine (C), but instead of thymine, it uses the pyrimidine uracil (U).

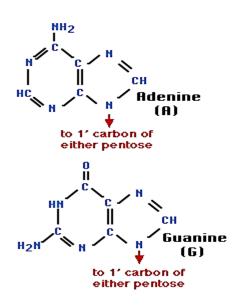
http://users.rcn.com/jkimball.ma.ultranet/BiologyPages/N/Nucleotides.html

DNA: Nucleotides

Bases

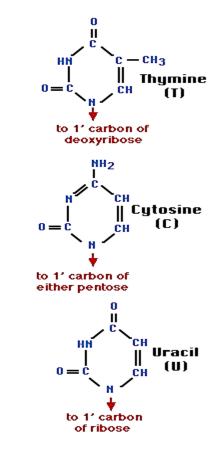
The Pyrimidines

The Purines



http://users.rcn.com/jkimball.ma.ultranet/BiologyPages/N/Nucleotides.html

The combination of a base and a pentose sugar is a nucleoside



DNA: Polymerization

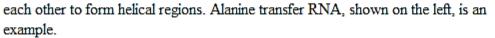
Nucleotide + Nucleotide + Nucleotide +

The polymerization of nucleotides.

The nucleic acids, both DNA and RNA, consist of polymers of nucleotides. The nucleotides are linked covalently between the 3' carbon atom of the pentose and the phosphate group attached to the 5' carbon of the adjacent pentose. The figure on the right shows the polymer structure of DNA.

Most intact DNA molecules are made up of **two** strands of polymer, forming a "<u>double helix</u>".

RNA molecules, while single-stranded, usually contain regions where two portions of the strand twist around



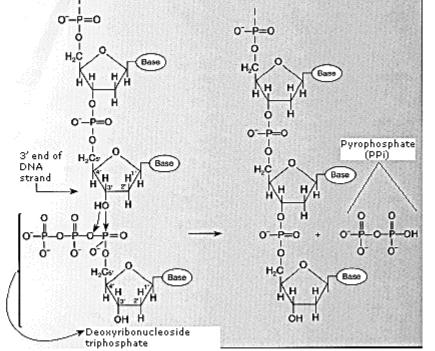
The two strands of DNA and the helical regions of RNA are held together by base pairing.

The polymerization of DNA is described more fully in DNA Replication.

Alanine

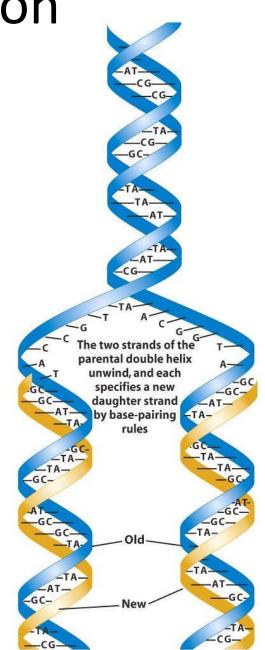
5'

3'



DNA: Replication

- Complementary base pairing and the double helix
- Replication is semiconservative



- 1. DNTPs: dATP, dGTP, dTTP, and dCTP
- 2. Template DNA (a pre-existing single strand)
- 3. DNA polymerase

1. DNTPs

One (as shown in the first figure), two, or three **phosphate** groups. These are attached to the 5' carbon atom of the pentose. The product in each case is called a **nucleotide**.

Both DNA and RNA are assembled from nucleoside triphosphates.

For DNA, these are dATP, dGTP, dCTP, and dTTP.

For RNA, these are ATP, GTP, CTP, and UTP.

In both cases, as each nucleotide is attached, the second and third phosphates are removed.

The nucleosides and their mono-, di-, and triphosphates

	Base	Nucleoside	Nucleotides		
DNA	Adenine (A)	Deoxyadenosine	dAMP	dADP	dATP
	Guanine (G)	Deoxyguanosine	dGMP	dGDP	dGTP
	Cytosine (C)	Deoxycytidine	dCMP	dCDP	dCTP
	Thymine (T)	Deoxythymidine	dTMP	dTDP	dTTP
RNA	Adenine (A)	Adenosine	AMP	ADP	ATP
	Guanine (G)	Guanosine	GMP	GDP	GTP
	Cytosine (C)	Cytidine	CMP	CDP	CTP
	Uracil (U)	Uridine	UMP	UDP	UTP

http://users.rcn.com/jkimball.ma.ultranet/BiologyPages/N/Nucleotides.html

2. Template DNA (a pre-existing single strand)

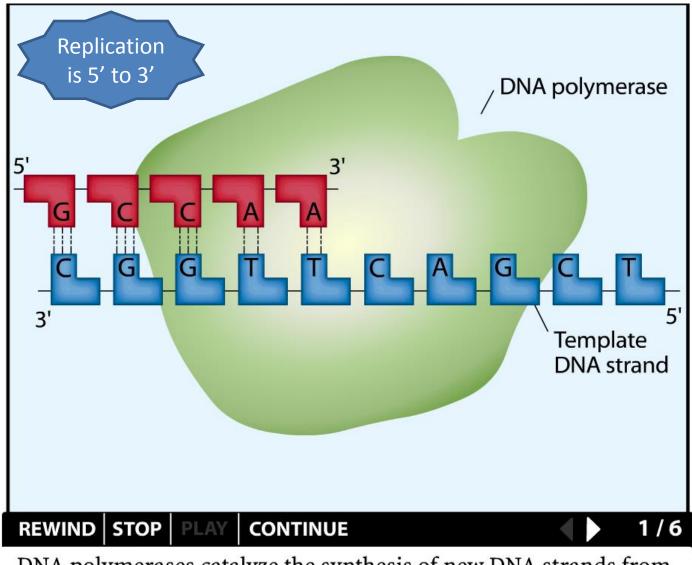
ATCGGTCAACGTTAAAGTTAGCGG

3. DNA Polymerases

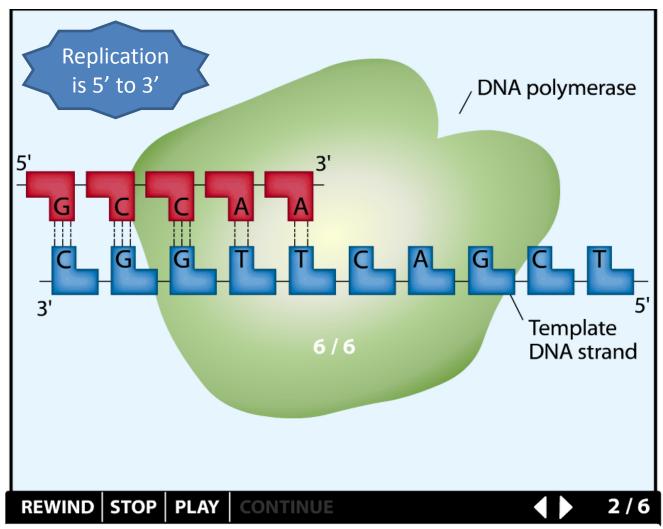
- There are multiple forms of DNA polymerase
- Different forms have different activities
 - Replicases have direct roles in replication
 - > Others have secondary roles in replication and/or repair synthesis.
- DNA replication polymerization of deoxyribonucleotides
 - Polymerases catalyze the formation of a phosphodiester bond between the 3'-OH of the deoxyribose on the last nucleotide and the 5' phosphate of the dNTP precursor
 - Process is repeated forming a synthesized DNA chain

3. DNA Polymerases (continued)

- Polymerase binds to the DNA template strand and moves along as the synthesized polynucleotide chain grows
 - At each template base, the dNTP precursor is identified that can base pair with it
 - \succ The frequency of error is low, but errors can occur.
- Polymerases can have exonuclease activity (removal of nucleotides from the 3' end of the chain)
 - This is a proof-reading mechanism
 - An unpaired nucleotide from the 3'OH end of the growing chain triggers exonuclease activity
 - The unpaired nucleotide is cleaved from the end of the growing chain by the polymerase.

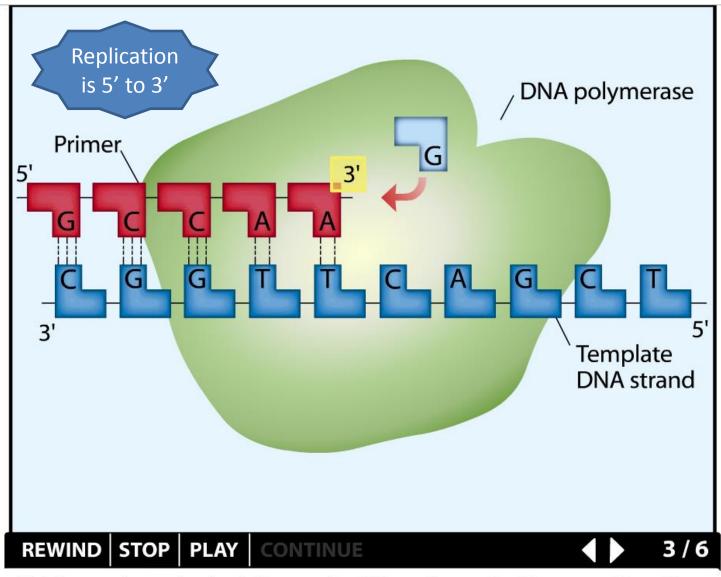


DNA polymerases catalyze the synthesis of new DNA strands from a DNA template.

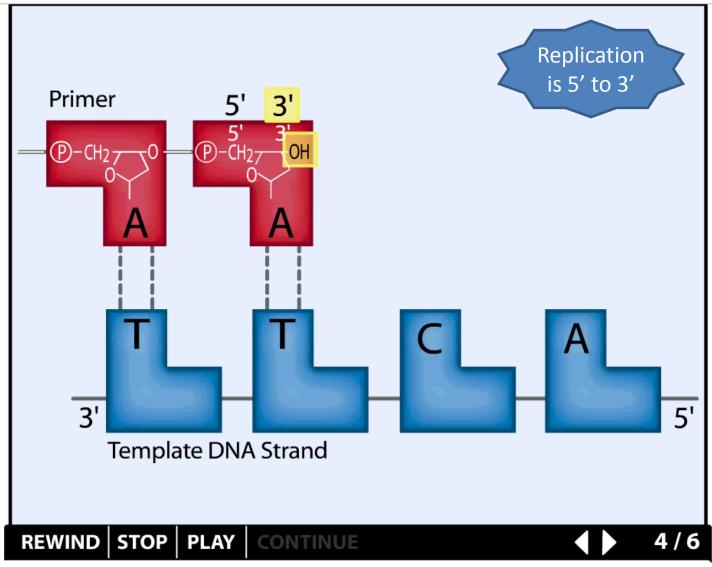


DNA polymerases require a pre-existing RNA or DNA strand, the primer, to initiate new DNA synthesis.

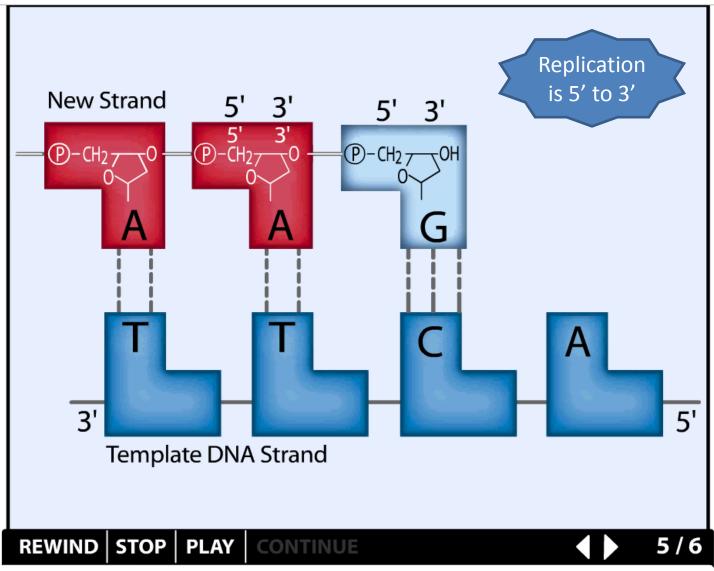
These polymerases add deoxyribonucleotides only to the 3' end of a growing strand.



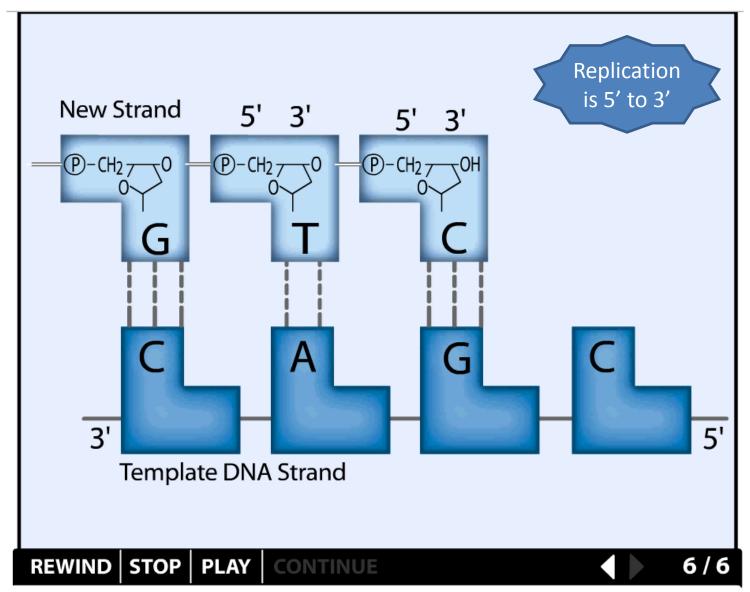
Taking a closer look at the nucleotide polymerization reaction, the 3' end of the primer contains a free 3'-hydroxyl group.



The 3' hydroxyl reacts with the 5' end of the next free nucleotide to be added.



Free nucleotides continue to be added to the growing DNA strand by the same type of reaction.

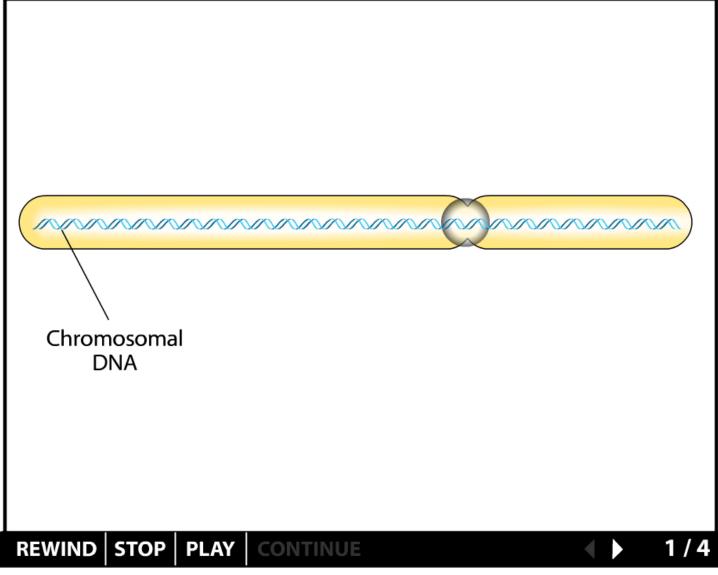


Overall, the new DNA strand grows in the 5' \rightarrow 3' direction.

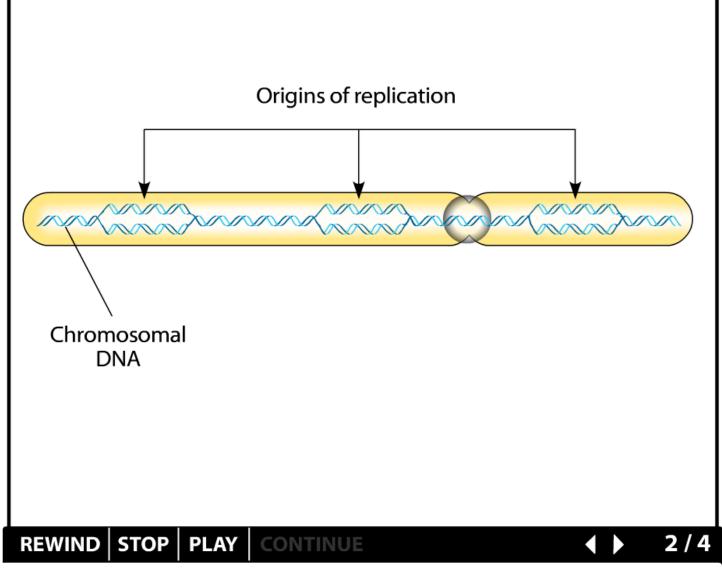
 Replication begins at a fixed point, called the origin, and proceeds bi-directionally. In a higher plant chromosome there are thousands of origins.

Consider

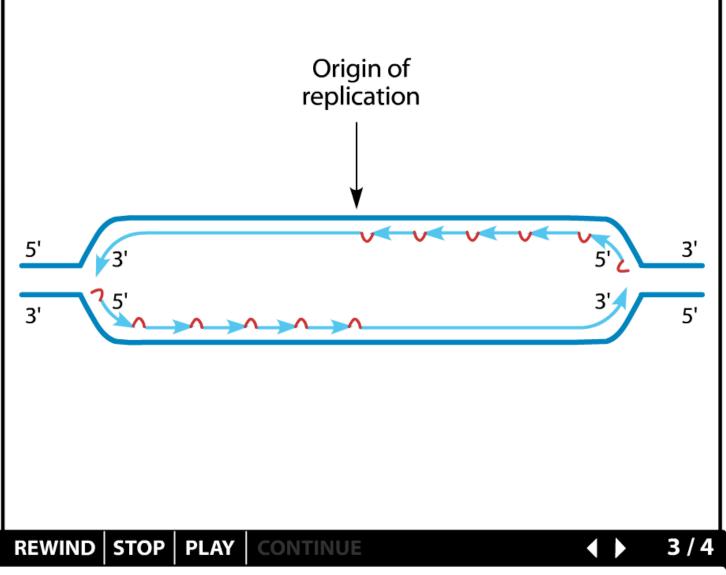
The size of the genome The rate of DNA replication The length of the S phase



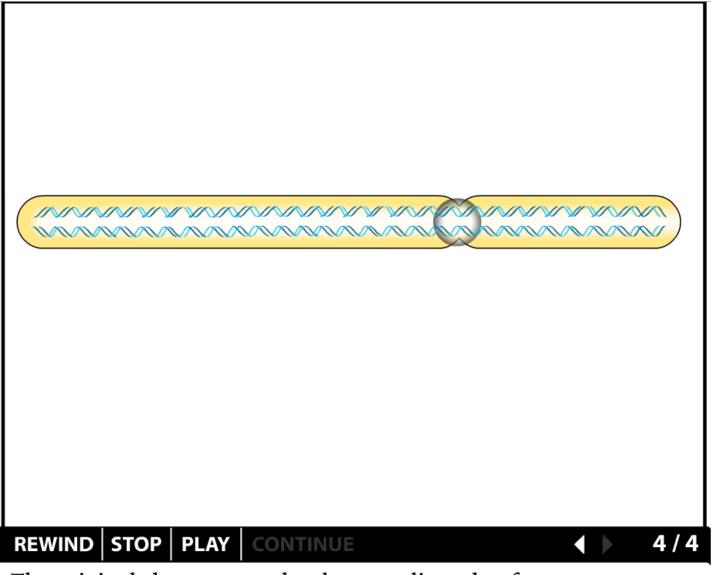
A eukaryotic chromosome contains multiple origins of replication.



At each origin of replication, DNA synthesis proceeds bidirectionally. Two replication forks move outward in opposite directions.



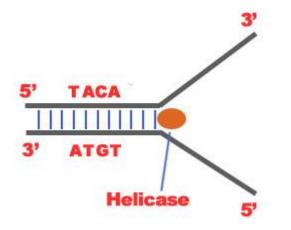
The replicating DNA helices from each origin elongate and eventually join each other.



The original chromosome has been replicated to form two identical daughter DNA molecules, also called sister chromatids.

Replication: 2-Unwinding, 3-Stabilization

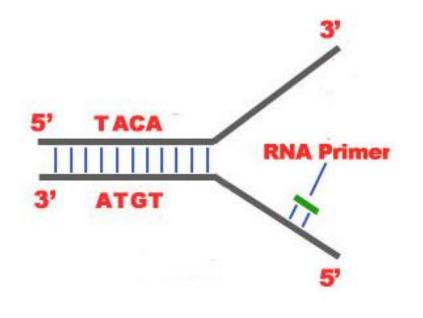
2. Unwinding: The DNA helix needs to be opened up. This is accomplished by <u>helicase</u> enzymes, which break the hydrogen bonds holding the two strands of the helix together. <u>Gyrase</u> facilitates helicase action by relieving tension in coiled DNA



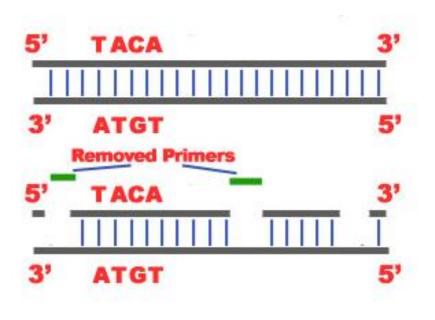
3. Stabilization: The unwound DNA is stabilized by a protein (single strand binding protein (SSB)), which speeds up DNA replication.

Replication: 4 - Priming

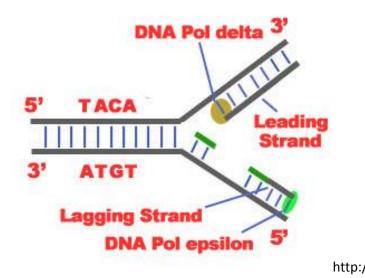
- Primases form a short RNA primer
- DNA polymerases use the primer to synthesise a new chain
 - Polymerases cannot start synthesis on their own
- RNA primers subsequently removed by exonuclease activity of a polymerase and replaced with DNA

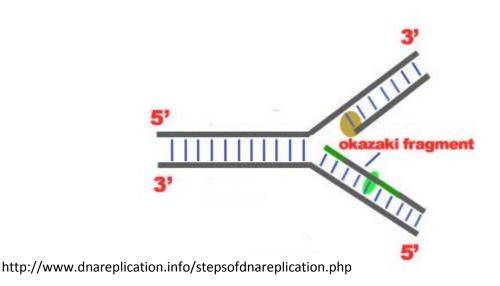


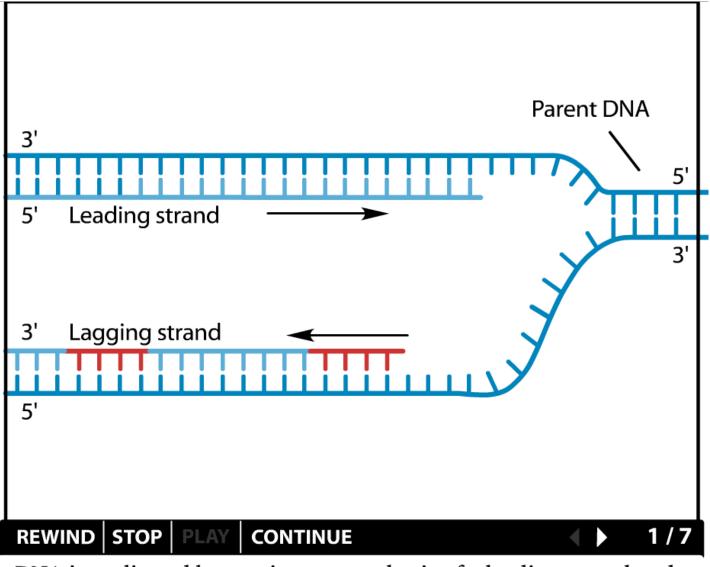
http://www.dnareplication.info/stepsofdnareplication.php



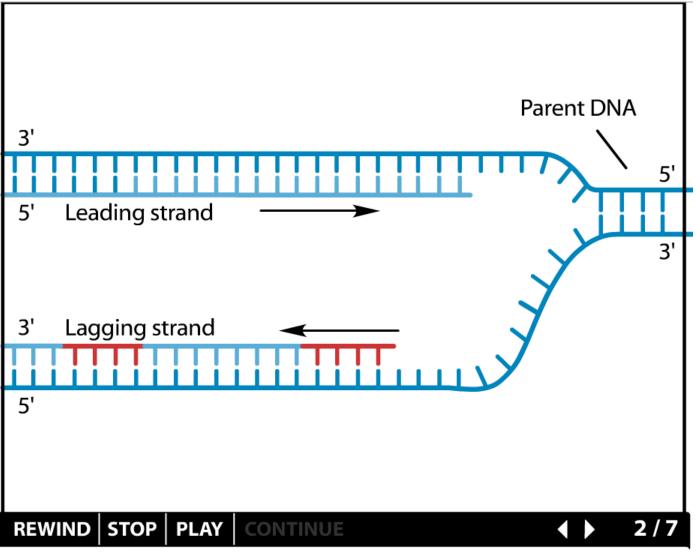
- DNA polymerases synthesize new chains only from 5' to 3'
 DNA molecule is antiparallel and DNA synthesis is semi-conservative.
- DNA synthesis is continuous on 5'-3' strand leading strand
- Synthesis discontinuous on 3'-5' strand lagging strand
 - Multiple priming sites on the lagging strand
 - DNA therefore formed in fragments on lagging strand Okazaki fragments



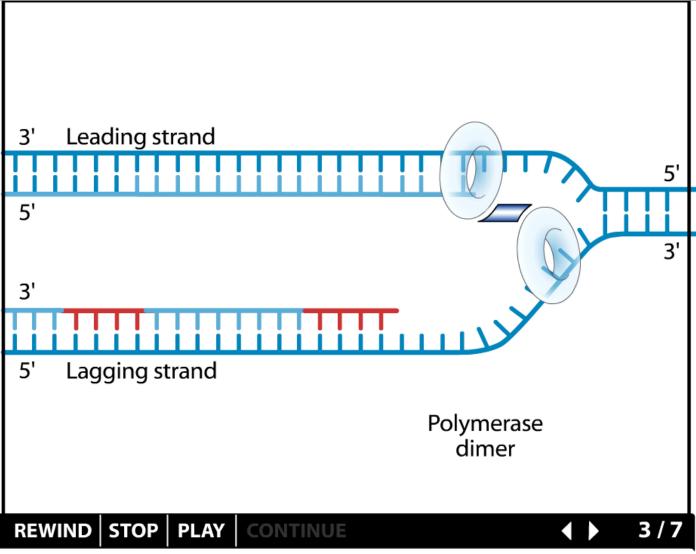




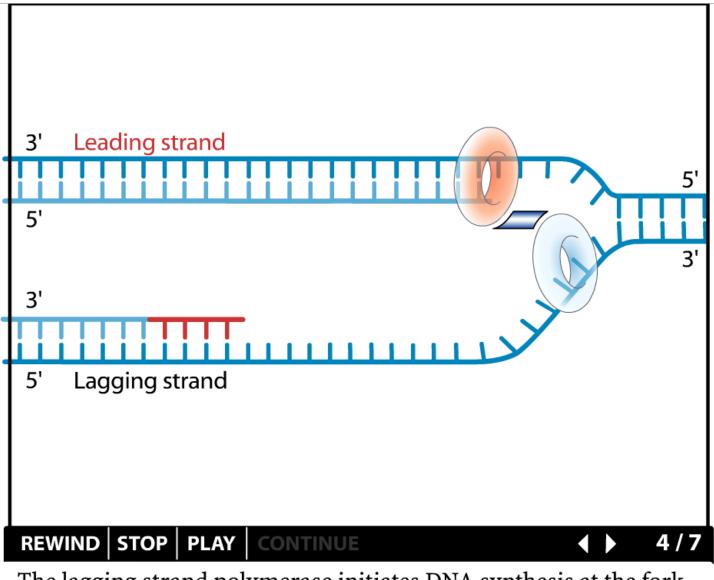
DNA is replicated by continuous synthesis of a leading strand and discontinuous synthesis of a lagging strand.



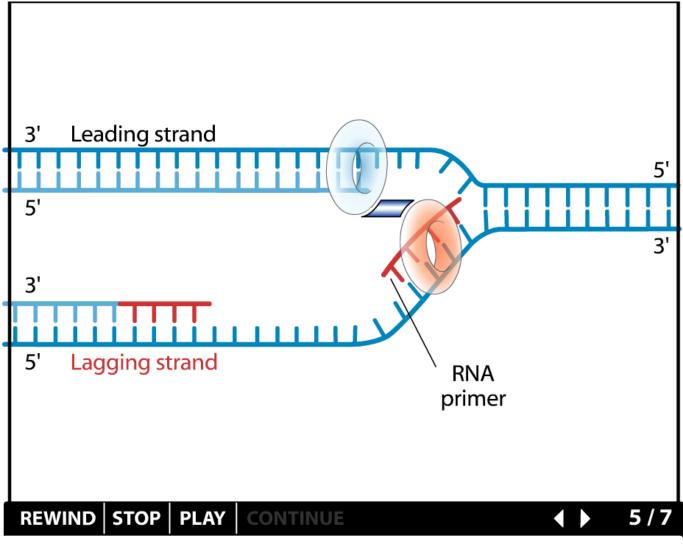
Coordination between leading and lagging strand synthesis is achieved by the dimerization of DNA polymerase molecules at the replication fork.



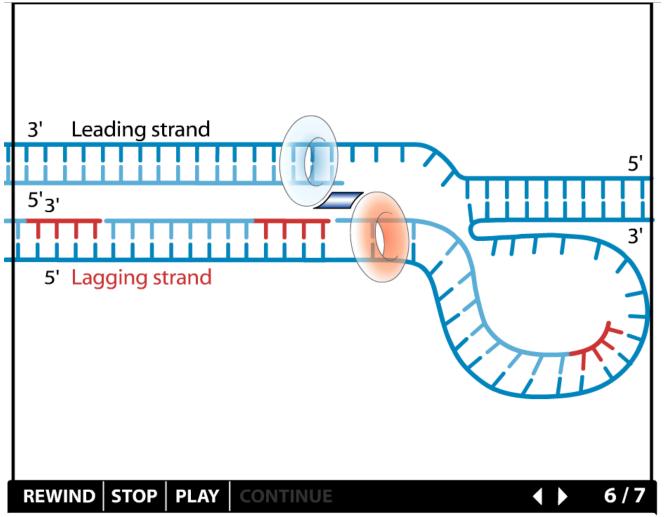
The DNA polymerase dimer moves with the replication fork. The polymerase at the leading strand template remains attached to the DNA, continuously synthesizing the leading strand.



The lagging strand polymerase initiates DNA synthesis at the fork, from an RNA primer made by a primase complex.

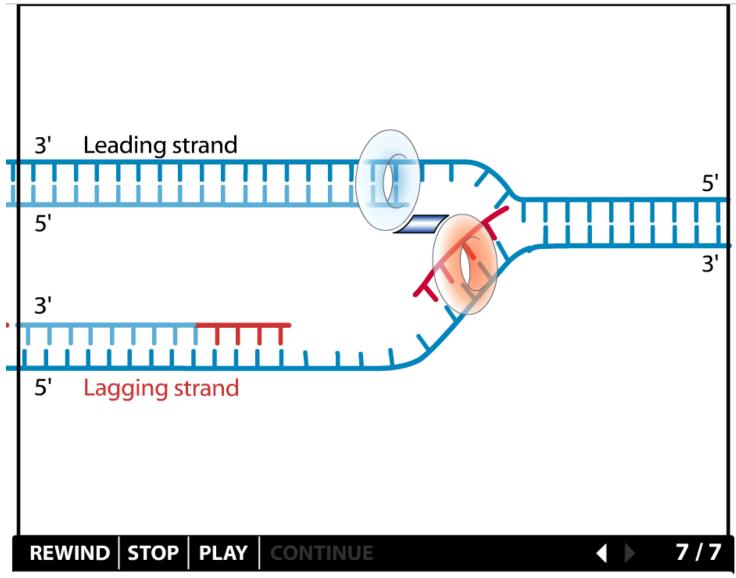


The polymerase elongates the lagging strand in a direction opposite the fork, but stays bound at the fork. As a result, the newly synthesized lagging strand fragment loops out between the polymerase and the fork.



Once the polymerase completes an Okazaki fragment, it dissociates from the DNA template.

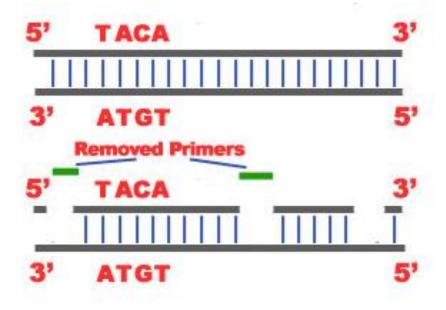
A new primer is produced at the fork. The polymerase reassociates with the template at this position, to continue synthesis of the lagging strand.



By this mechanism, the two polymerases can add nucleotides to the growing strands at the same time, and at rates up to 1000 bp/s.

Replication: 6 Termination

- DNA Pol I exonuclease removes RNA Primers in lagging strand
- DNA Polymerase adds complementary nucleotides to fill the gaps
- **DNA Ligase** adds phosphate in the remaining gaps of the phosphate sugar backbone
- Can't backfill RNA primer site at telomere of lagging strand
 - Telomere shortening during replication
 - Role of Telomerase



The telomerase solution

https://www.youtube.com/watch?v=vtXrehpCPEE

The Plant Cell, Vol. 16, 794-803, April 2004, www.plantcell.org © 2004 American Society of Plant Biologists

HISTORICAL PERSPECTIVE ESSAY

Plant Telomere Biology

Analysis of telomeres, the nucleoprotein 1938) and concluded that "... the terminal complexes that physically cap and protect gene must have a special function, that of studies that these chromatin bridges eventhe ends of eukaryotic chromosomes, has sealing the end of the chromosome, so to tually snapped, and the sister chromatids a long and intriguing history. The recent speak, and that for some reason a chromo-then fused to recreate a dicentric chromoresurgence of plant telomere biology some cannot persist indefinitely without some (McClintock, 1938). The question she prompted us to recap this history to pro- having its ends thus sealed." After explain- was asking now was whether this chromovide background and context for current ing that one difference between the termi-some breakage-fusion-bridge cycle would investigations addressing how plants main- nal gene and the others is that it is unipolar, continue indefinitely. To answer the questain a stable genome. Although many of the with genes on only one side of it, Muller tion, McClintock placed her dicentric-genfundamentals of telomere biology were first deduced that the bipolar genes in the erating chromosome 9 in a background uncovered in ciliates or fungi, telomere interior of the chromosome "... cannot be with a normal chromosome 9 carrying reresearch in Arabidopsis allows us to ask made into properly functioning unipolar cessive alleles for plant color (yellow green, basic questions in a multicellular organism ones by the simple expedient of tearing yg) and seed characteristics (waxy endowith complex development and excellent them loose from their connections on one sperm, wx; colored aleurone, c; and genetic tools. This powerful combination of side" (Muller, 1938). Muller coined the term shrunken endosperm, sh) in the region advantages is unsurpassed in other organ-telomere for this terminal gene from the between the two centromeres. Any break-

McClintock knew from her previous

Replication: 7 Proof-Reading

- Mis-matched bases can be added in error during polymerization
- Specific polymerases with 3'-5' exonuclease activity identify and remove any detected mis-matches on the synthesized strand
- Proofreading starts at the 3' end of the synthesized strand and proceeds 3' -> 5'

Two exact copies of original, each with one original and one new strand

