

### DNA Replication

-happens in SPhase - proceeds bidirectional

-copying a DNA molecule

-semiconservative

-DNA unwinds and unzips 2 strands

- Daughter DNA

*parent strand and newly synthesized*

-unzip = supercoiling

### SUPERCOILING

**DNA** -enzyme topoisomerase

**Gyrase** -creates a nick to relieve torsional strain and release

**Helicase** -Destabilizing protein  
-promotes unwinding by binding to RF

**SSB** -stabilize single stranded regions

### DNA POLYMERASE

-polymerization at 5' to 3' |add in 3' end

-requires template to copy |complementary strand

#### Types of dNTPs

dATP, dGTP, dCTP, dTTP

#### Primer

Organisms RNA, DNA in some organisms

Viruses Protein

DNTP - deoxynucleotide triphosphate adenine (dATP), cytosine (dCTP), guanine (dGTP), and thymine (dTTP)

### DNA POLYMERASE

-polymerization at 5' to 3' |add in 3' end

-requires template to copy |complementary strand

#### Types of 4dNTPs

dATP, dGTP, dCTP, dTTP

#### Primer

Organism RNA, but DNA in some organism

Virus Protein

DNTP - deoxynucleotide triphosphate adenine (dATP), cytosine (dCTP), guanine (dGTP), and thymine (dTTP)

### Models of DNA Replication

SemiConservative	Conservative	Dispersive
------------------	--------------	------------

-parent & new synthesis	-both original and new strands	-mosaic/hybrid
-------------------------	--------------------------------	----------------

-unwind each other	-rebase pair
--------------------	--------------

### Theta Replication

-present in bacteria (prokaryotic)	- circular form
------------------------------------	-----------------

### LAGGING AND LEADING STRAND

Lagging Strand	Leading Strand
----------------	----------------

-3' to 5'	-5' to 3'
-----------	-----------

-discontinuous	-continuous
----------------	-------------

-okazaki fragments	-no short fragments
--------------------	---------------------

-each fragments requires its own set of primers	- requires only one primer
---	----------------------------

-requires DNA ligase	-no DNA ligase
----------------------	----------------

### LAGGING AND LEADING STRAND (cont)

-grows away from RF	-grows in direction of RF
---------------------	---------------------------

-synthesis of new strands SLOW	-synthesis of new strands FAST
--------------------------------	--------------------------------

RF - Replication Fork

### REPLICATION ORIGINS

-unique DNA segments	contains multimeric origin-binding protein
----------------------	--

-ATrich stretch	less energy to melt A-T bp due to Hbonds
-----------------	--

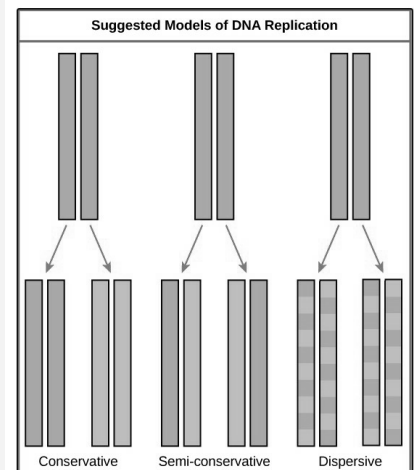
### PROKARYOTIC DNA POLYMERASE

DNA polymerase I	-RNA primer removal, DNA repair
------------------	---------------------------------

DNA polymerase II	-DNA repair
-------------------	-------------

DNA polymerase III	-Chromosome replication
--------------------	-------------------------

### MODELS OF DNA REPLICATION



### PROTEINS INVOLVED IN DNA REPLICATION

Protein	Activity
DNA Polymerase	-initiates -add nucleotide at 3' end
DNA Helicase	-unzip/unwind DSDNA -separates DNA strands
SSBS	-stabilizer -preventing bp from reforming before lagging strand is replicated
DNA Topoisomerase	-re-seal -unravel twists in DNA -reduce torsional strain & positive supercoil
Sliding Clamp	-keeps DNA polymerase attached to template
Clamp Loader	-lock sliding clamp to DNA (uses ATP hydrolysis)
Primase	-synthesis RNA primers in lagging strand
DNA Ligase	-sealed/connect okazaki
SSBS - Single Strand DNA Binding Protein DSDNA- Double Strand DNA	



By **elrchqt**  
[cheatography.com/elrchqt/](https://cheatography.com/elrchqt/)

Not published yet.  
Last updated 22nd August, 2023.  
Page 2 of 2.

Sponsored by **ApolloPad.com**  
Everyone has a novel in them. Finish Yours!  
<https://apollopad.com>