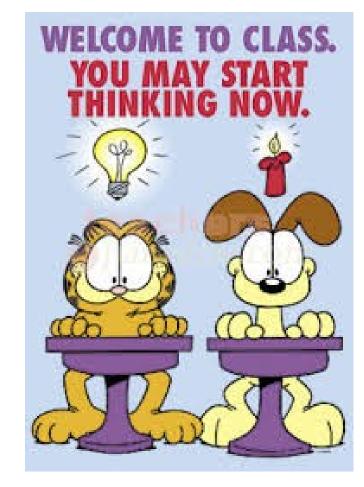
Yeah !! , it's the <u>Second</u> <u>year</u> !!!!!!

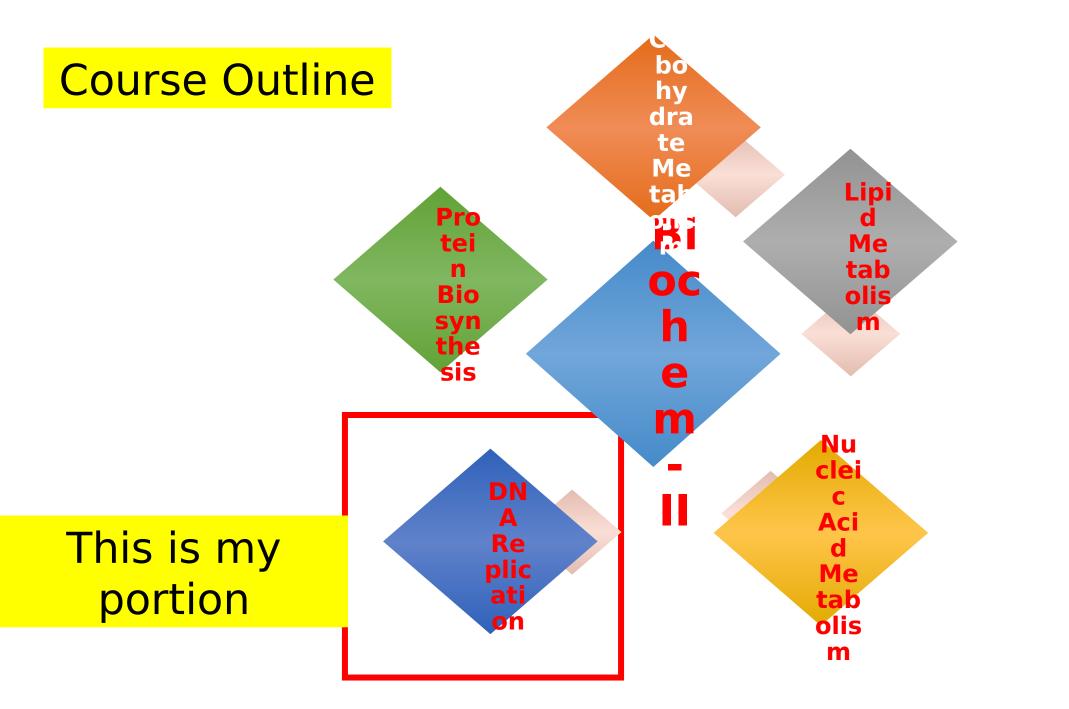






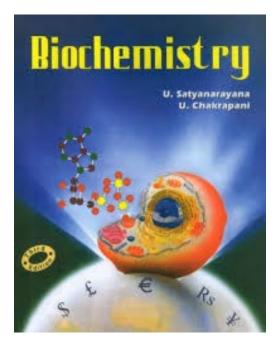


Shariq

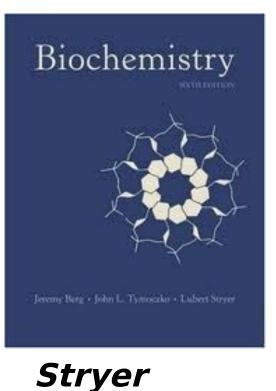


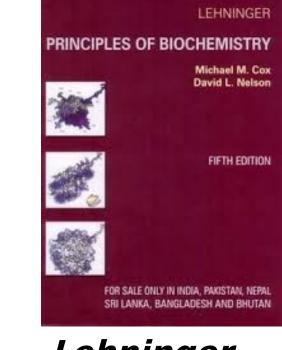
Books we will refer do

Shariq



Satyanara yana Primary Book





Lehninger

Excellent

Reference

We have a new "Kool" eblackboard !!!

Week1	DNA Replication Animation Kool animation from Harvard University on DNA Replication	
www.padlet.com/shariqasad/bi ochem		
I'll be posting my lecture notes, exciting article & any other links I find useful		

Biochemistry - II

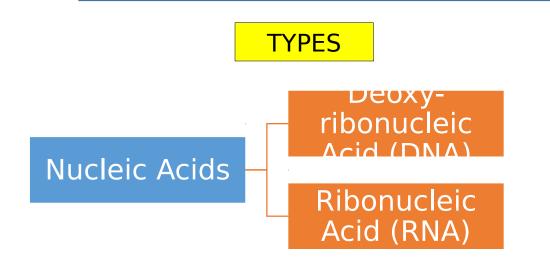
NA Replication - 1

Dr. Shariq Syed



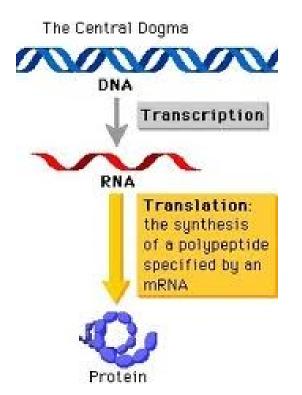
Shariq

What are "Nucleic Acids" ??

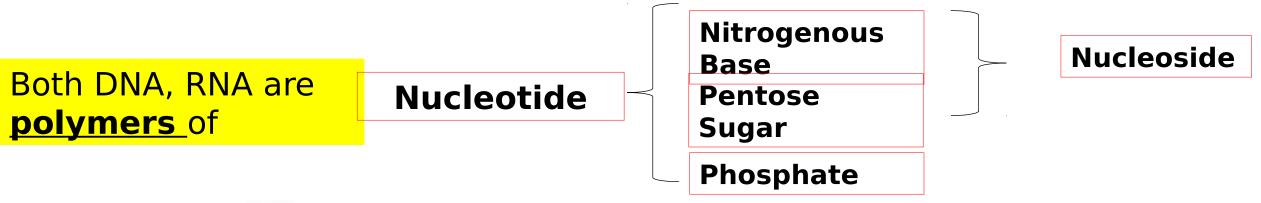


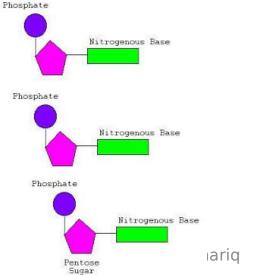
What's job of nucleic acid ??

- Storage
- Transmission of GENETIC

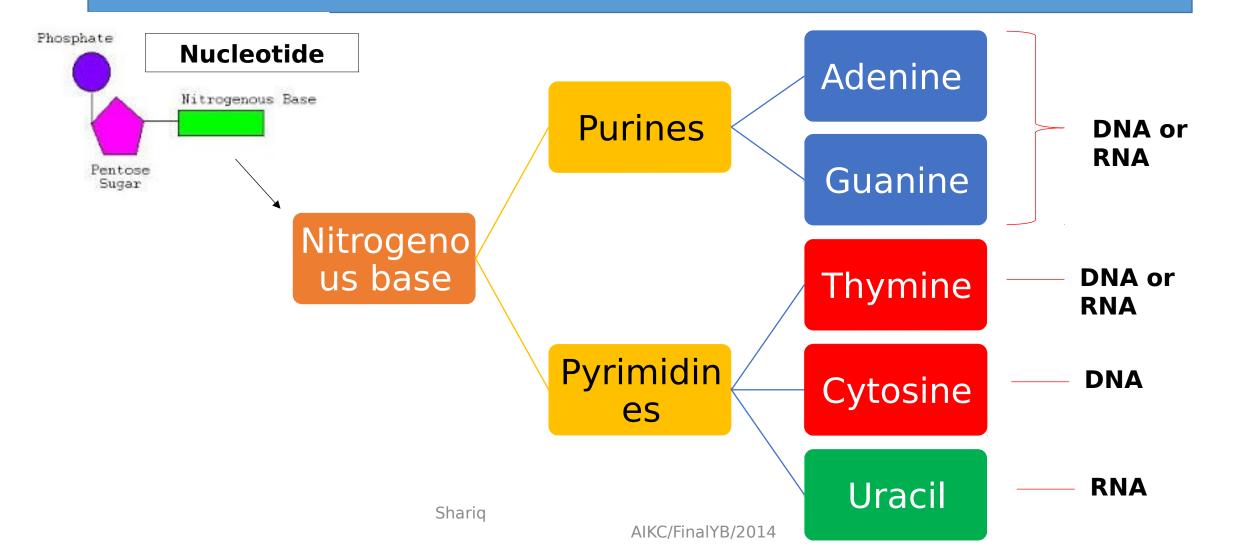


What's the composition/structure of Nucleic Acid ??

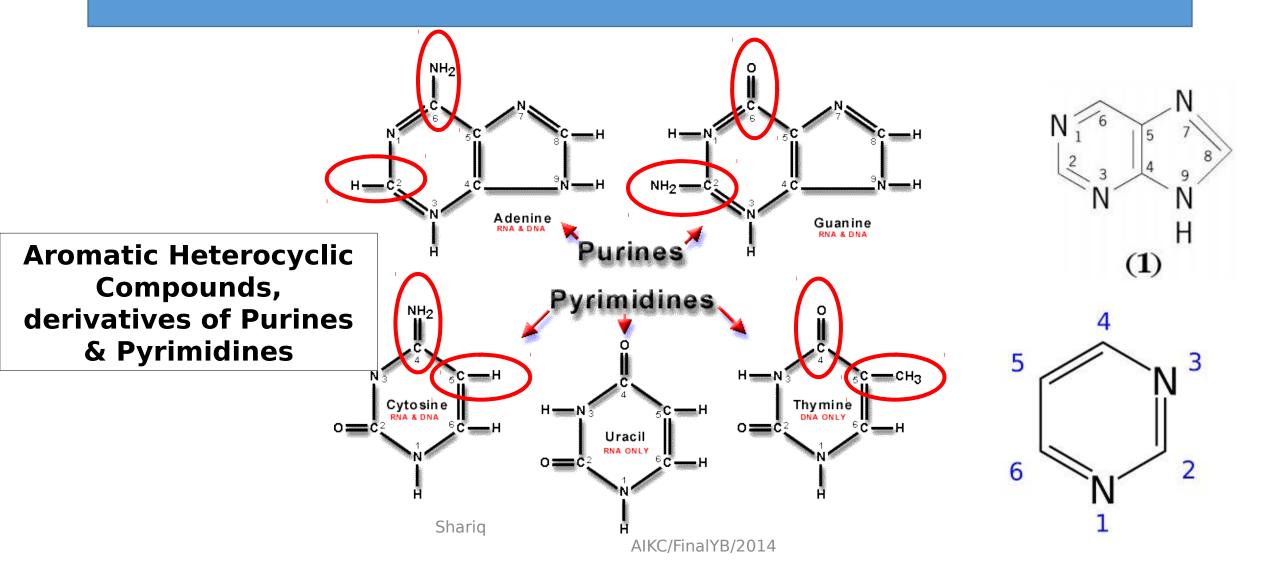




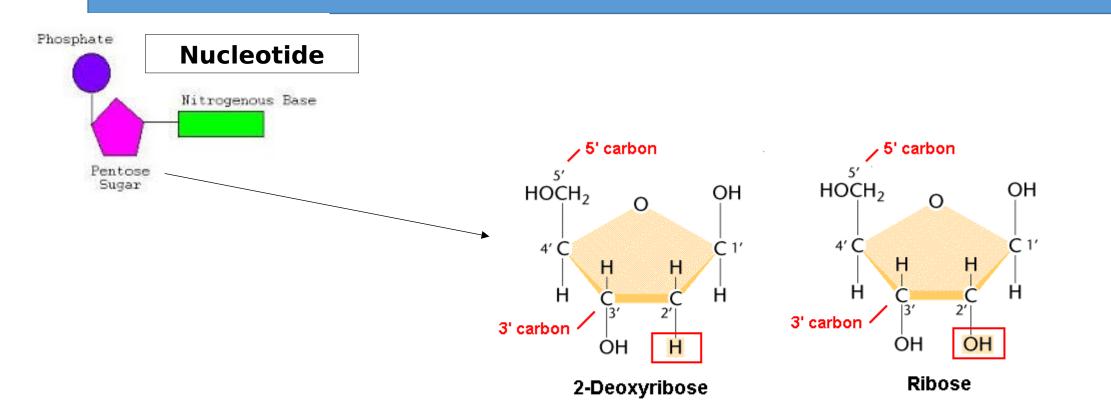
Nitrogenous Base



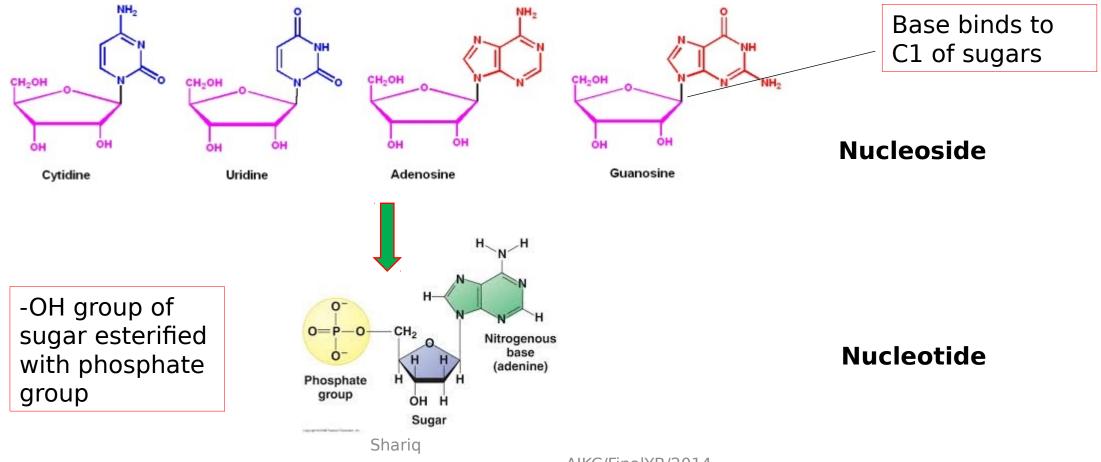
Nitrogenous Base



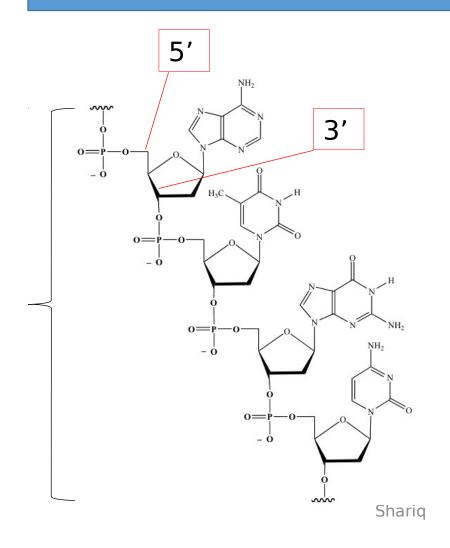
Pentose Sugars



Nucleotide

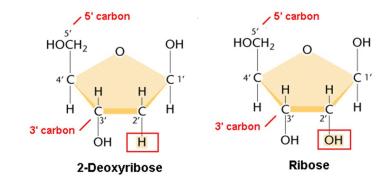


Nucleotide



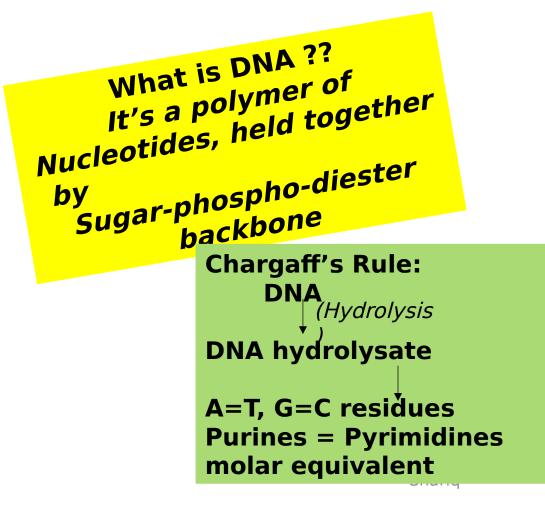
Esterification of –OH group happens at both 3' & 5' position of sugars

This "dual" esterification provides a phosphate backbone to nucleotide polymer



AIKC/Fin

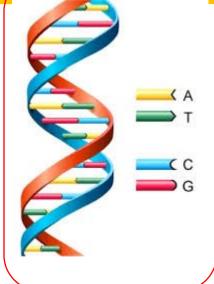
Structure of DNA



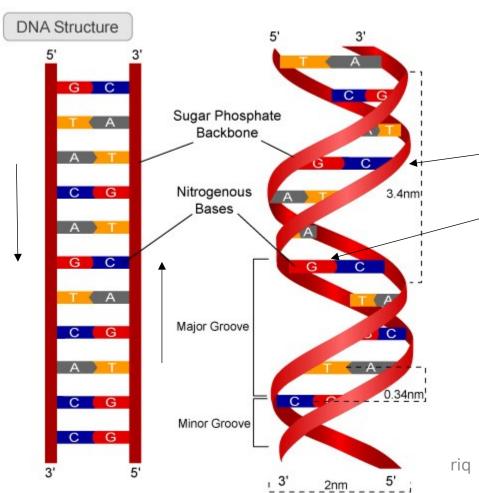
<u>Watson & Crick proposed a</u> Double-Helix structure for DNA

Twisted Ladder structure



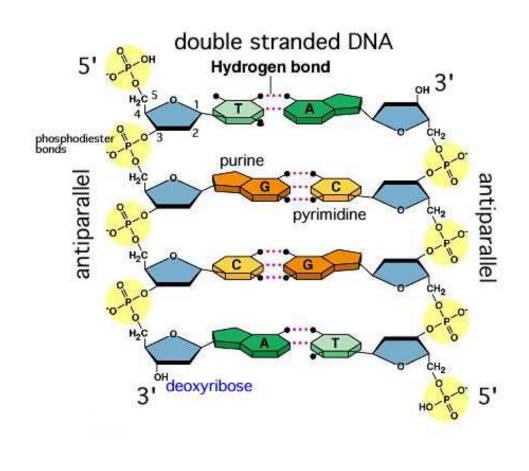


Double Helix Structure of DNA



- <u>Two</u> polynucleotide chains twisted around each other on common axis
- The Two strands run **anti-parallel**
- Hydrophilic phosphate chain outside
- Hydrophobic bases form inside core
- Two chains held together by H-bonds
- A=T, G C
- Proposed structure agrees to Chargiff's rule

Double Helix Structure of DNA

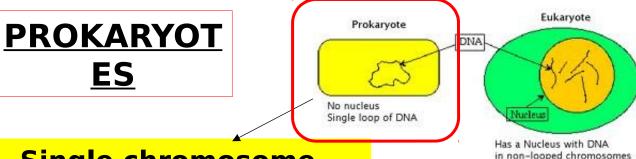


 Two chains run anti-parallel, 5' end, 3' end

- Space orientation <u>only</u> permits purine=pyrimidine H-bonds
- Purine=purine would not fit in structure while

 Pyrimidine=pyrimidine would be too far to form H-bonds

How is DNA organized in a CELL



Single chromosome

ES

Double stranded circle

(DNAL Drotoins - cation

capsul

Chromosome are packed in form of nucleoids

cell wall

nucleoid (DNA)

ribosomes

flagellum

DNA Binding Protein

Cell Envelope

Nucleoid Organization:

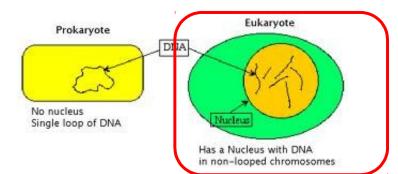
DNA Origin

Domain

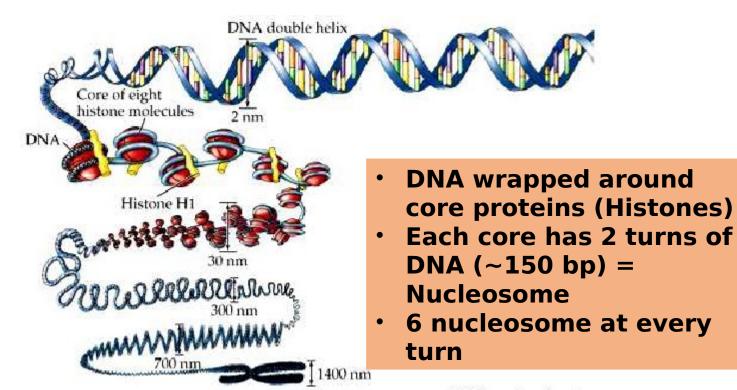
 The NUCLEOID is an irregularly-shaped region within the cell of a prokaryote that contains all or most of the genetic material.

Shariq

How is DNA organized in a CELL

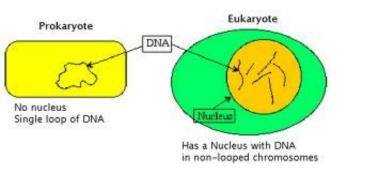


- Eukaryote = DNA in clearly defined nucleus
- DNA + Proteins = chromatin which gets organized as chromosome

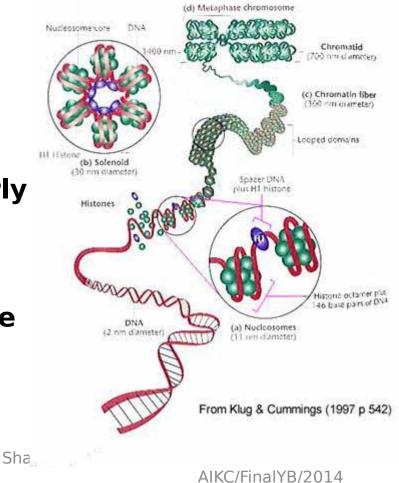


4 2001 Sinauer Associates, Inc.

How is DNA organized in a CELL

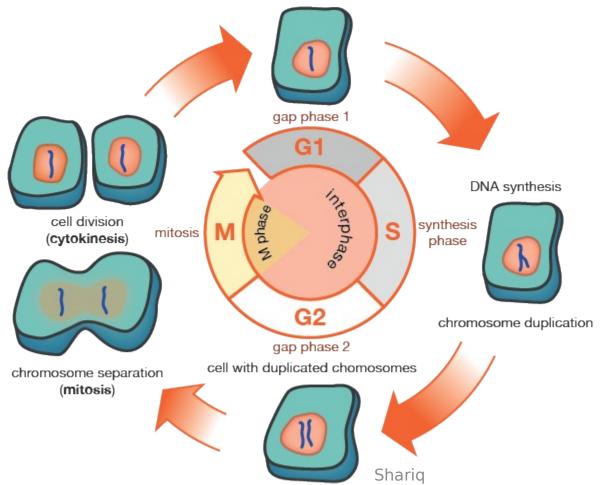


- Eukaryote = DNA in clearly defined nucleus
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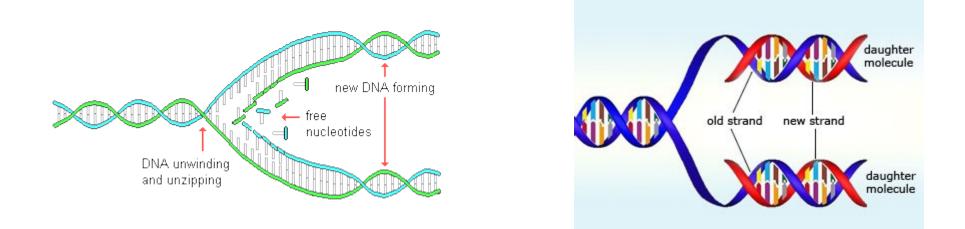
- DNA wrapped around core proteins (Histones)
- Each core has 2 turns of DNA (~150 bp) = Nucleosome
- 6 nucleosome at every turn

When does DNA Replication occur during Cell Cycle ??

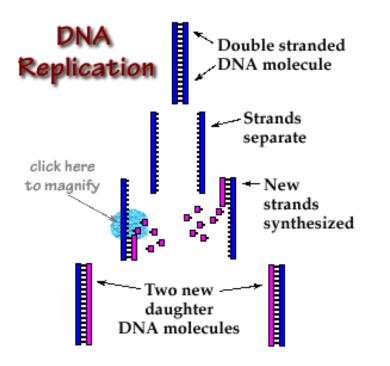


- DNA synthesis takes place in <u>S-phase</u>
- Entire process take 8-10 hrs
- Large number of DNA Synthesizing enzymes (500-1000) are involved

DNA Replication; Short story



DNA Replication is *Semi*conservative



What happens during replication ??

- Parent strand is separated
- Creating two separate strand
- These single strand act as template for new strand
- "Semi-conservative" because half of parent DNA retained in daughter DNA

Site of DNA Replication

Site of Replication:

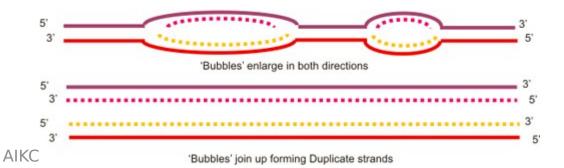
Location where the DNA "unzips" Single location for prokaryotes, multiple for eukaryotes Region rich in A=T sequence since easy to break A=T bond

Specific protein "dna A" opens up DNA

- Opening of DNA leads to formation of <u>replication</u> <u>bubbles</u>
- Multiple in case of eukaryotes (Important for rapid replication)



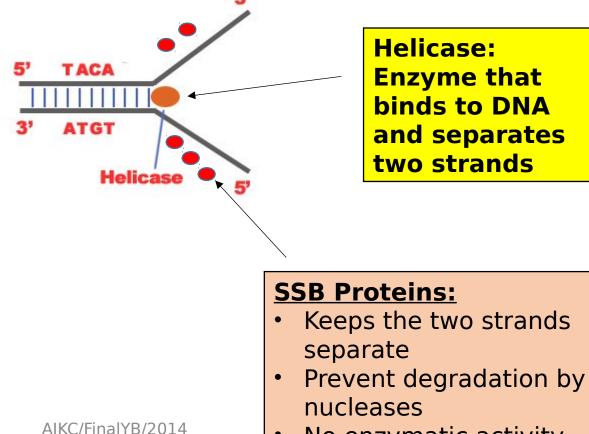
'Bubbles' form in DNA double strand



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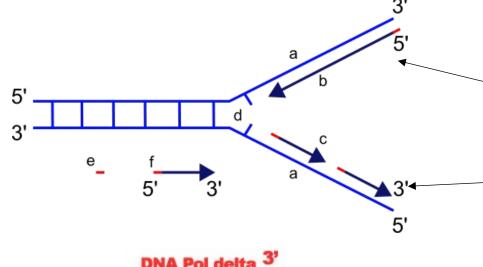
Site of Replication:

- Separation of DNA strands results in formation of <u>replication fork</u>
- Replication fork moves forward along parent DNA, daughter DNA is synthesized

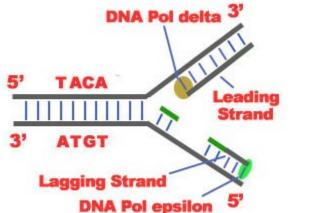


No enzymatic activity

Shariq



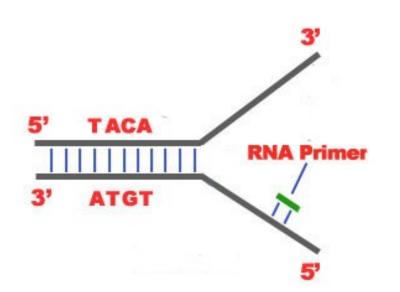
Shariq



 DNA replication occurs only in one direction 5'-3'

At one strand there is Continuous synthesis of DNA (5'-3' direction)

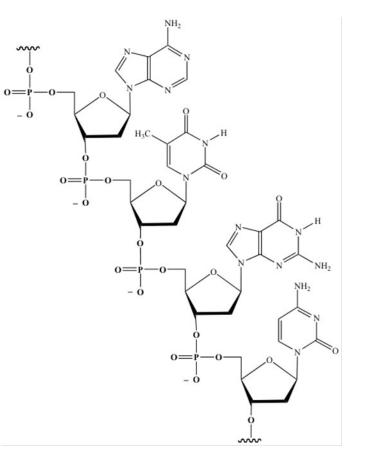
At another strand there is Discontinuous synthesis

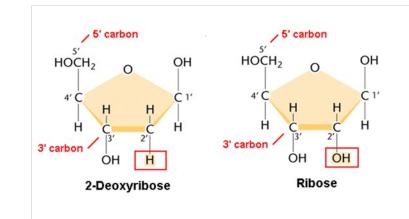


- **RNA Primase** binds to initiation point of the 3'-5' parent chain
- RNA Primase synthesis short RNA (~ 5nucleotides) which bind to the DNA nucleotides of the 3'-5' strand
- RNA nucleotides are the primers (starters) for the binding of DNA nucleotides
- These RNA primers are later removed by hydrolysis & replaced by DNA

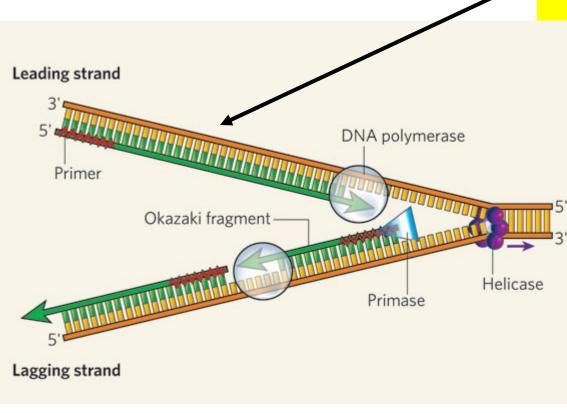
Shariq

DNA Polymerase can add nucleotide ONLY to 3' end





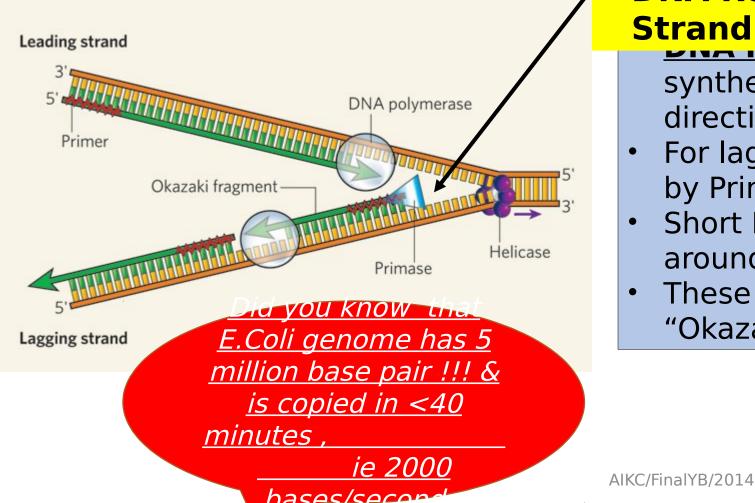
(Klug & Cummings 1997)



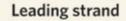
Shariq

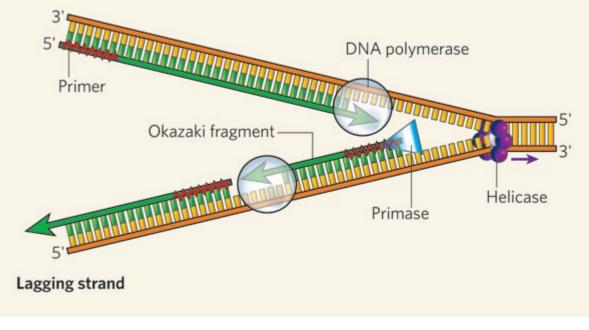
 DNA Replication at "Leading" Strand

- DNA Polymerase catalyses synthesis of new DNA ONLY in 5'-3' direction
- Synthesis occurs continuously at leading strand
- Incoming nucleotides are appropriately added by DNA polymerase to 3' end of growing chain
- Nucleotide in triphosphate form, when added to chain single Ppi is removed



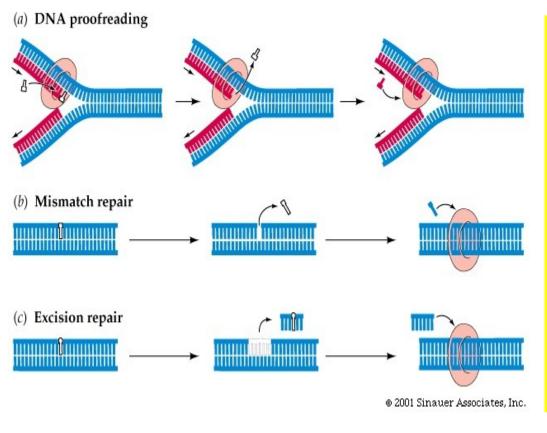
- DNA Replication at "Lagging" Strand
 - synthesis of new DNA **ONLY** in 5'-3' direction
 - For lagging strand, several RNA primers by Primase enzymes formed
 - Short DNA sequences synthesized around these primers
 - These short DNA fragments are called "Okazaki" fragments





- There are multiple RNA primers on lagging strand & one on leading strand
 - DNA synthesis continues till the strand is near to primers
 - DNA polymerase-I removes primer, synthesizes fresh DNA portion
- DNA ligase enzyme seals the two DNA ends

Proof reading DNA !



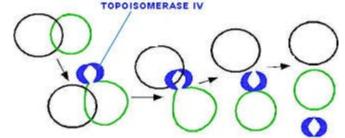
- As they add new bases to a growing strand, DNA polymerases-III make a proofreading check
- When a DNA polymerase recognizes an error, it removes the wrong nucleotide and tries again.
- The error rate of DNA polymerase on each attempt is only about 1 in 10,000, so the second attempt at matching the template is very likely to be successful.
- This proofreading function reduces the overall error rate to about one base in a billion (one in 10⁹⁾

Termination of replication

- DNA has a certain base pair region called "Ter"
- When replication fork enters, it cannot leave (it's a trap !)
- These supercoiled chromosomes are called "catenated"

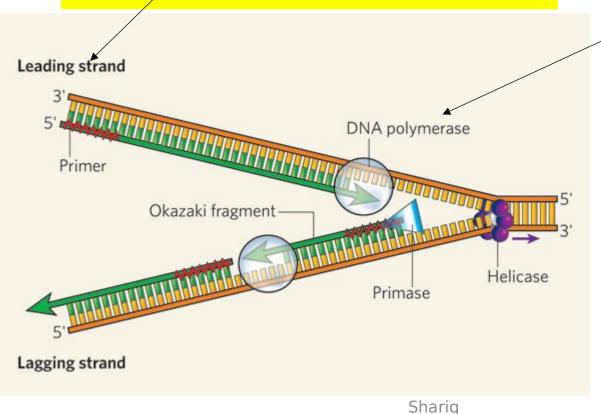
Role of Topoisomerase

- Topoisomerases are enzymes that regulate the overwinding or underwinding of DNA
- During DNA replication, DNA becomes overwound ahead of a replication fork
- Topoisomerases bind to either single-stranded or double-stranded DNA and cut the phosphate backbone of the DNA



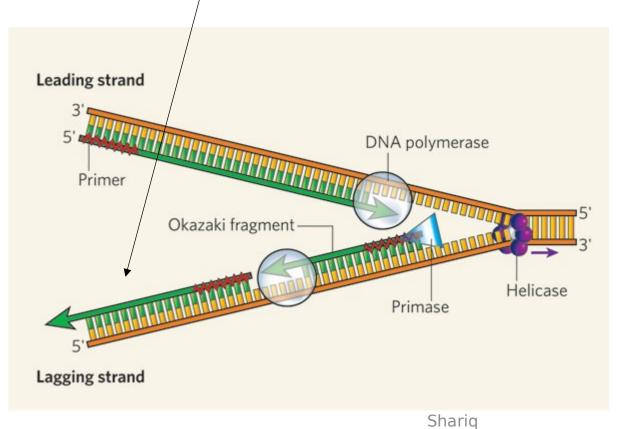
 This intermediate break allows the DNA to be untangled or unwound, and, at the end of these processes, the DNA backbone is resealed again

Replication at Leading Strand



- <u>DNA polymerase- δ</u> along with clamp slider <u>PCNA</u> (proliferating cell nuclear antigen)
- PCNA forms a ring around DNA to which polymerase binds

Replication at Lagging Strand

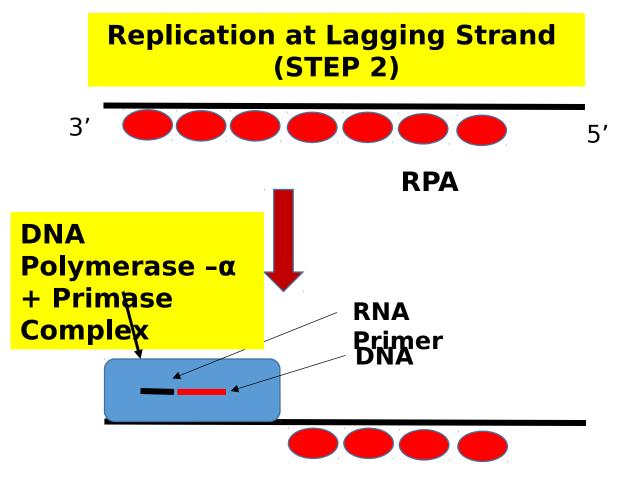


 DNA synthesis at lagging strand is slightly a <u>COMPLEX (multi-step)</u> compared to prokaryotes

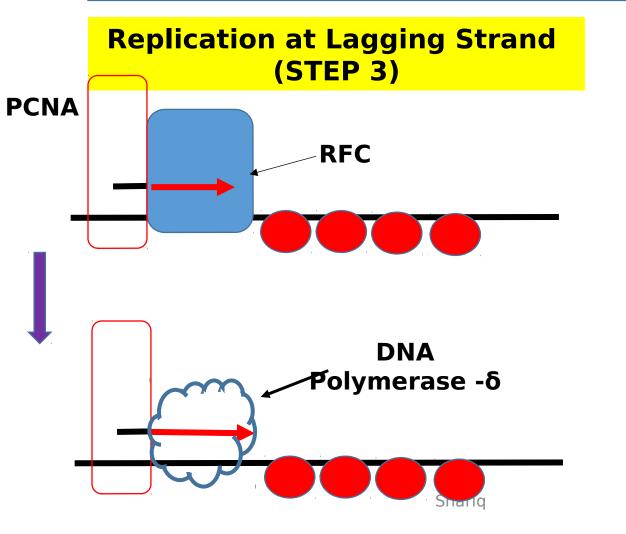
(STEP 1) Single Strand of DNA 3' 5'

Replication at Lagging Strand

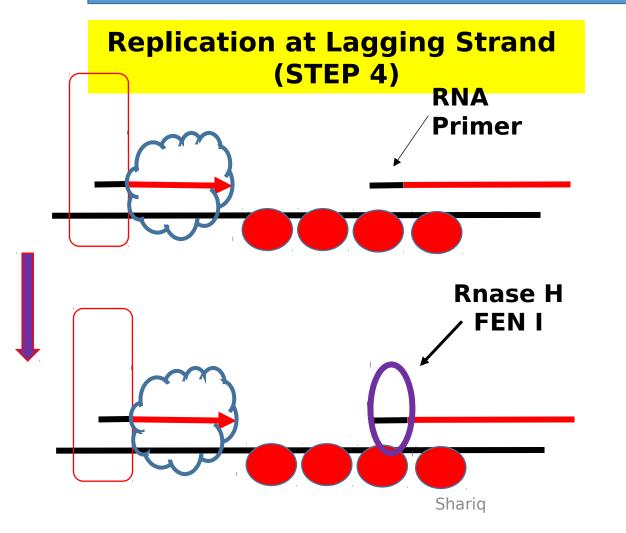
- Parental DNA separated by Helicase
- Exposed single strand is stabilized by *Replication Binding Protein* (RPA)



- Primase forms complex with DNA <u>Polymerase-α</u>
- Primase produces RNA primer (10 nucleotide)
- DNA Polymerase-α synthesizes short DNA (20-30 nucleotide)
- Primase+Poly complex dissociates

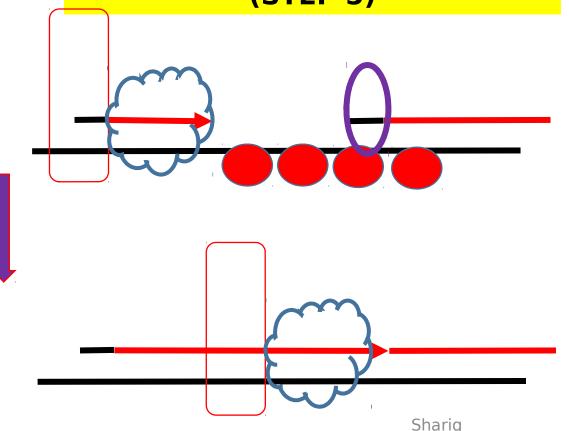


- RFC (Replication Factor) binds to elongated primer
- PCNA also gets added to RFC
- DNA Polymerase –δ binds to RFC (acts as sliding clamp)
- DNA Polymerase –δ elongates DNA fragment (150-200 bp)



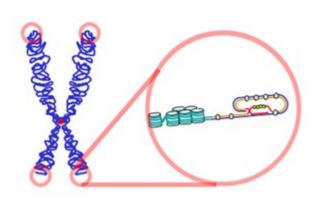
- Replicating DNA approaches primer of previous DNA fragment
- RNA primer is removed by enzymes *Rnase + FEN I*
- Gap filled by new DNA synthesized by DNA polymerase -δ
- Small nicks finally sealed by DNA Ligase

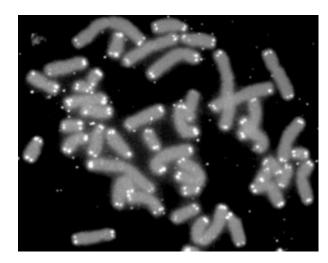
Replication at Lagging Strand (STEP 5)



- Replication DNA approaches primer of previous DNA fragment
- RNA primer is removed by enzymes *Rnase + FEN I*
- Gap filled by new DNA synthesized by DNA polymerase -δ
- Small nicks finally sealed by DNA Ligase

Telomeres





- Telomeres are end of chromosomes (*Telo* = end)
- Special structures to prevent continuous loss of DNA
- Protect ends of DNA
- Prevent chromosome fusing with others
- Sequence of TTAGGG is repeated X1000
- Are disposable buffers at the ends of chromosomes which are truncated during cell division
- They protects the <u>genes</u> before them on the chromosome from being truncated instead

Shariq

Telomerase

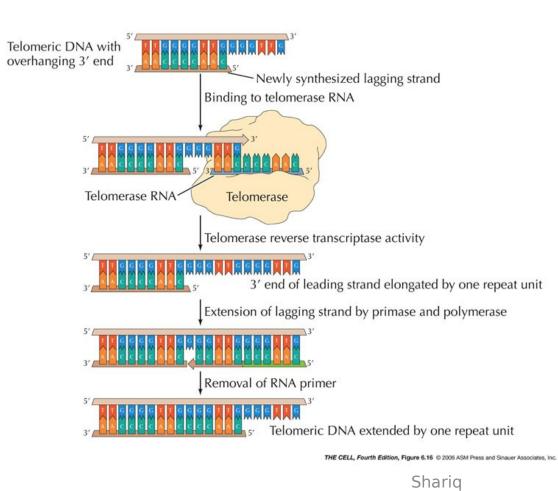
Last fragment Previous fragme **END-REPLICATION** PROBLEM **RNA** primer Lagging strand 5' **DNA Pol delta Primer removed but** Removal of primers and cannot be replaced replacement with DNA with DNA because where a 3' end is available no 3' end available TACA for DNA polymerase Leading Strand ATGT Second round of replication Lagging Strand **DNA Pol epsilon** New leading strand 3 New lagging strand 5'

iq



- DNA polymerases operate only in the 5' to 3' direction
- Synthesis of the lagging strand occurs through a "backstitching" mechanism that produces short fragments of DNA
- Without its complement, the hanging piece of unpaired DNA from the parent strand might be recognized by the cell as a broken piece of DNA and then damaged or even chopped off in the cell's

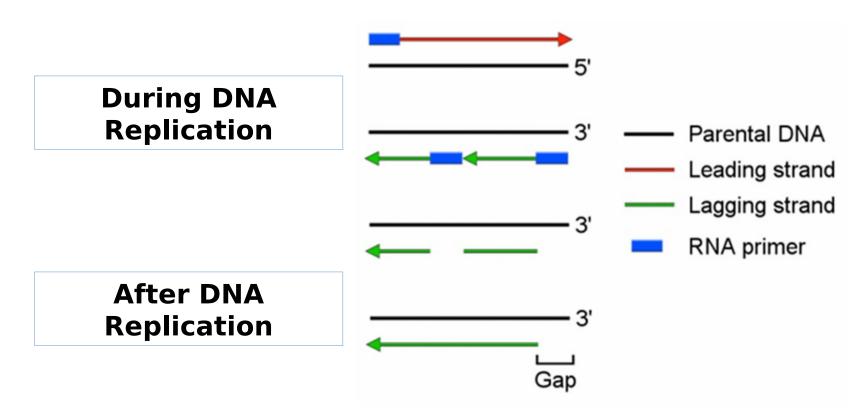
Telomerase

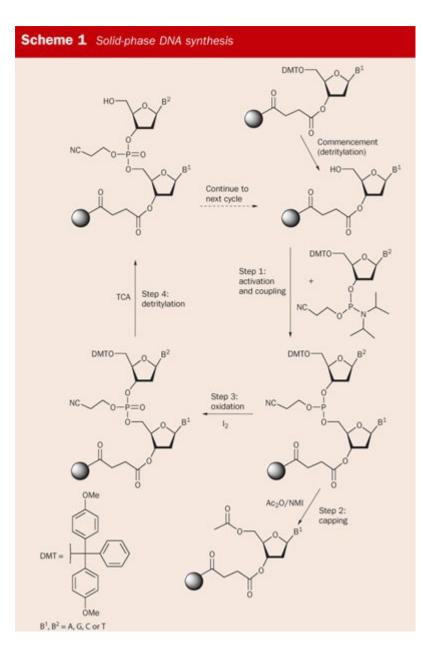


- Telomerase: <u>Unique protein +RNA</u> <u>template</u>
- Reverse transcriptase activity (Create DNA from RNA)
- DNA created is based on telomere template (TTAGGG) at leading strand
- Synthesis of DNA on lagging strand by DNA Polymerase
- At the end RNA primer is removed
- Some shortening will NOT create a problem
- Since Telomere do not encode proteins they Do NOT have to be of same length AIKC/FINALYB/2014

Telomerase

<u>"END-REPLICATION"</u>PROBLEM





http://www.atdbio.com/content/17/Solid-phase-oligonucleotidesynthesis