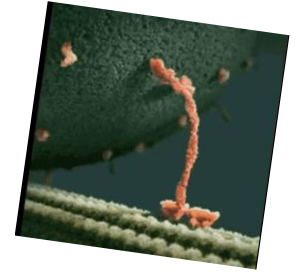




NanoMachines: The Tiny Biological Gadgets that Animate Life



I Love Lucy (1952)



Session 4
DNA and Proteins

OLLI at Illinois
Spring 2023

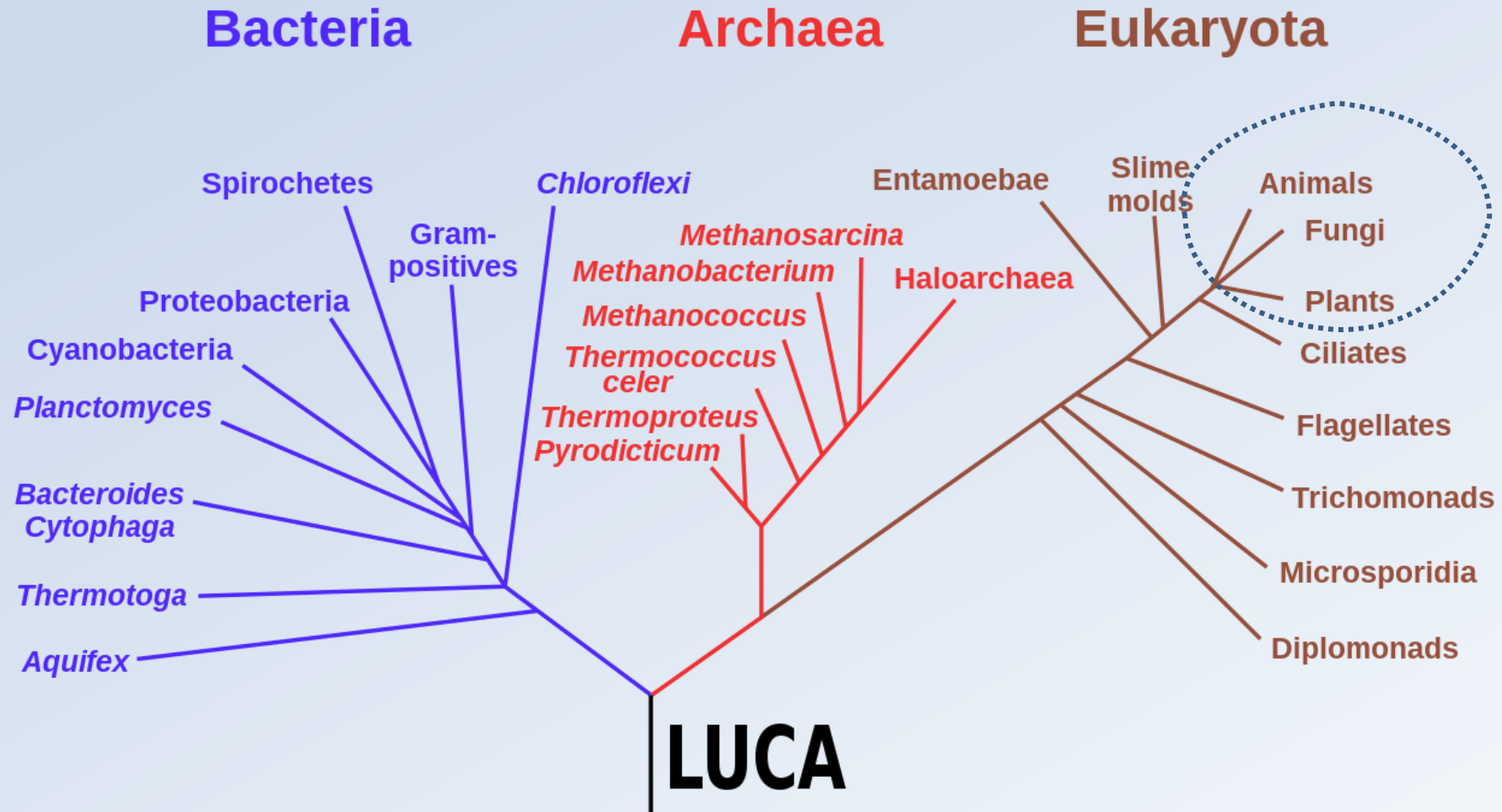
D. H. Tracy

Course Outline

1. The nanoscopic world: Building blocks, energy flow, how we observe it
2. Energy harvesting, storage and consumption
3. Movement, motors and locomotion
4. **Genetic information processing, protein manufacturing**

“LUCA”

Last Universal Common Ancestor



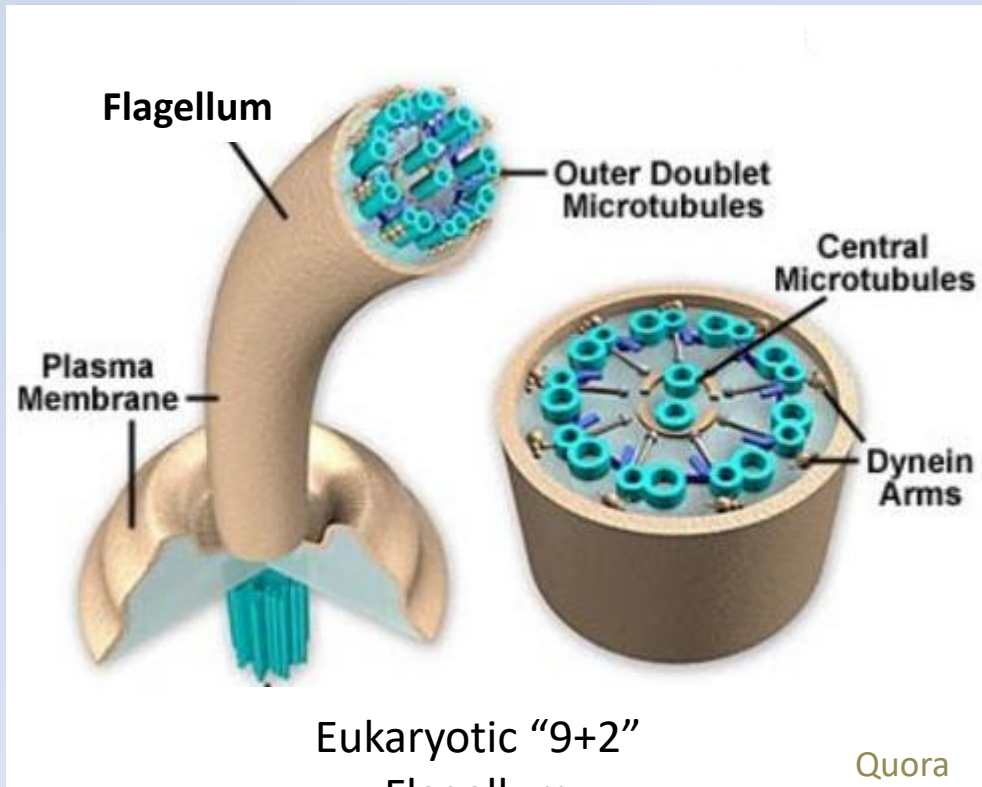
Session 4 Outline

- Finish Molecular Motors
- DNA & RNA: The Central Dogma
- DNA Replication
- RNA Blueprints: Transcription
- Protein Factories: Ribosomes

Two very different ways to build a Flagellum

Eukaryotic Flagella (e.g. Sperm or Algae)

Complex Flagellum **Bends**
using Dynein actuators



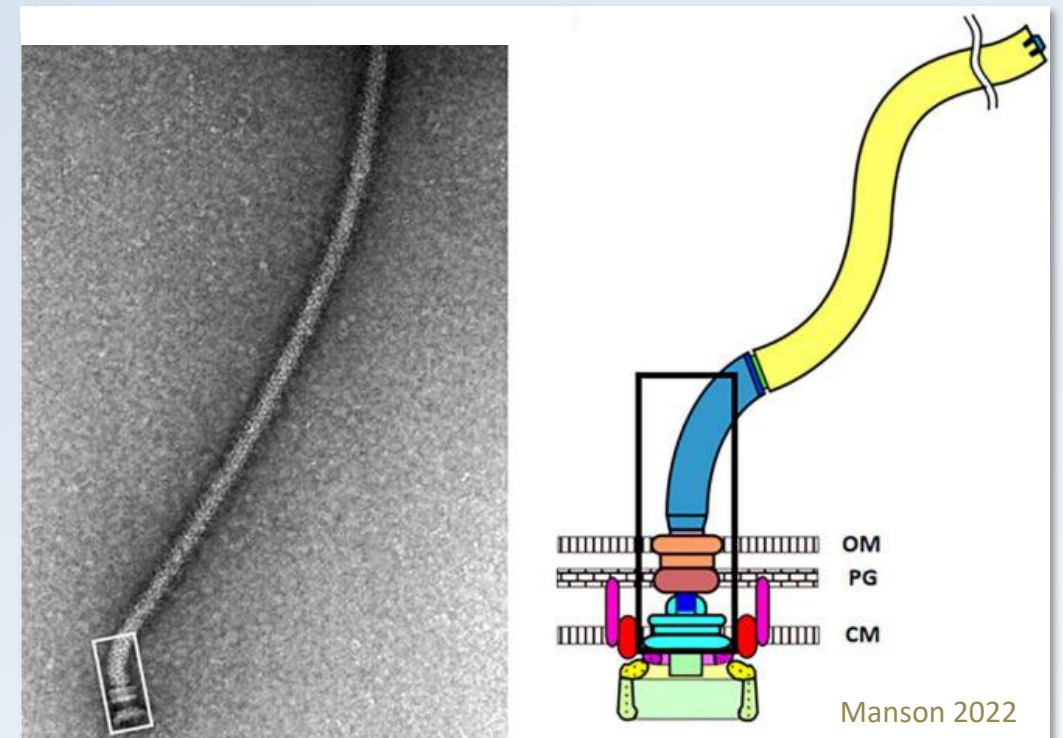
Eukaryotic "9+2"
Flagellum

Quora

3/24/2023

Prokaryotic Flagella (Bacteria & Archaea)

Simple Flagellum **Spins**
using a Rotary Motor

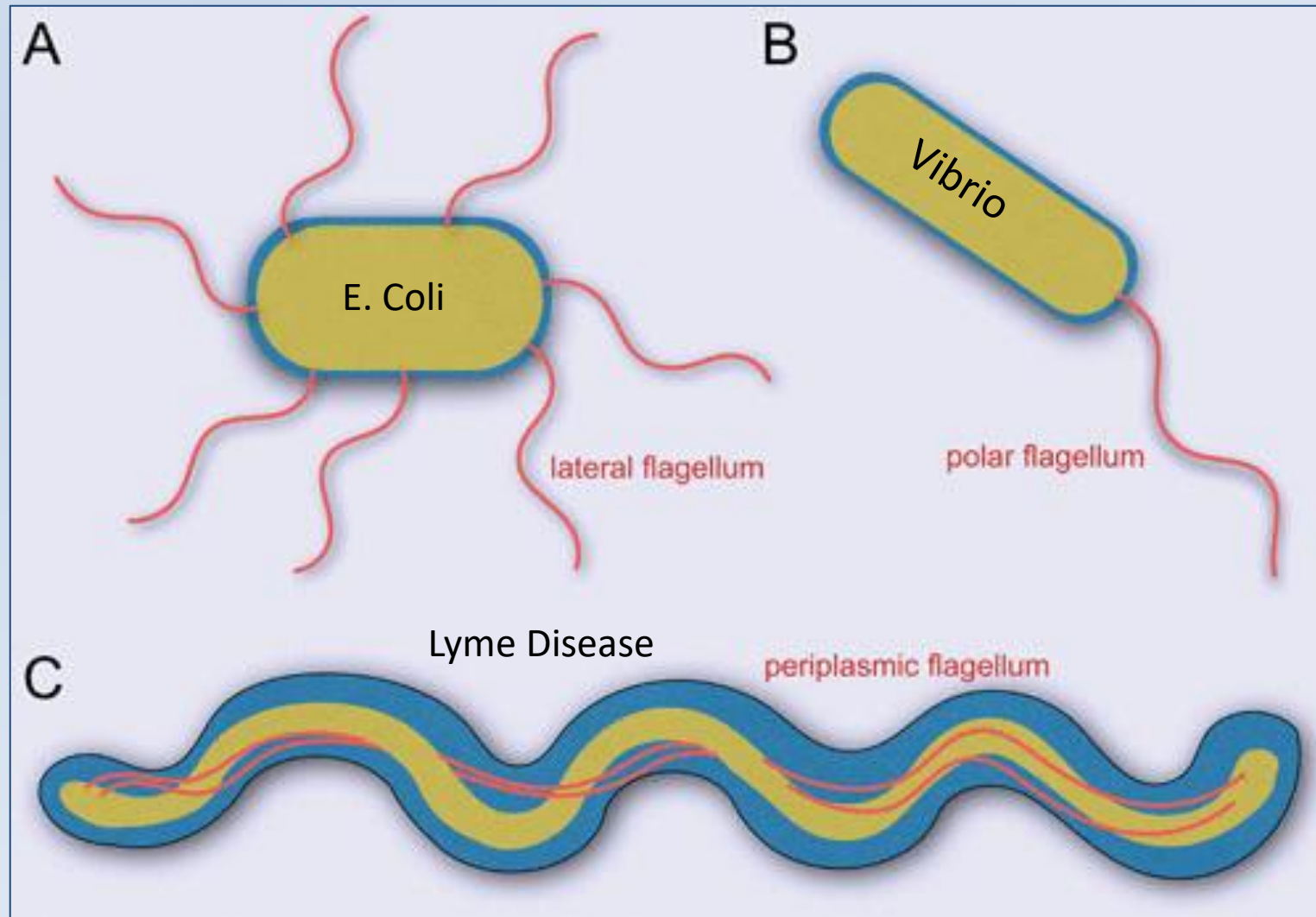


Manson 2022

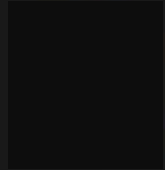
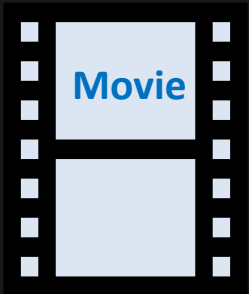
nanoMachines 4

26

Bacterial Flagella take many forms

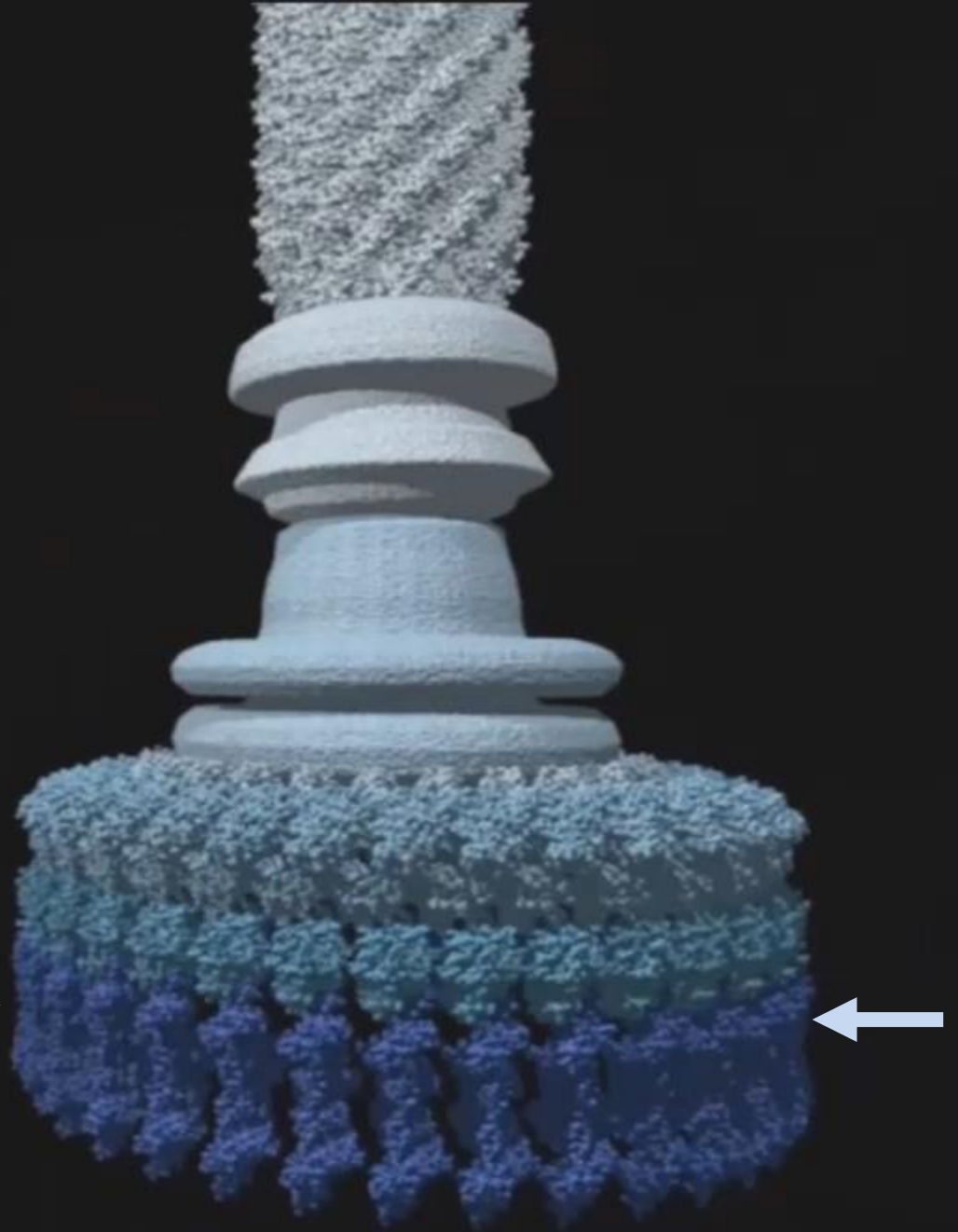


The Bacterial Flagellar Motor



ATP Synthase

~ 50nm

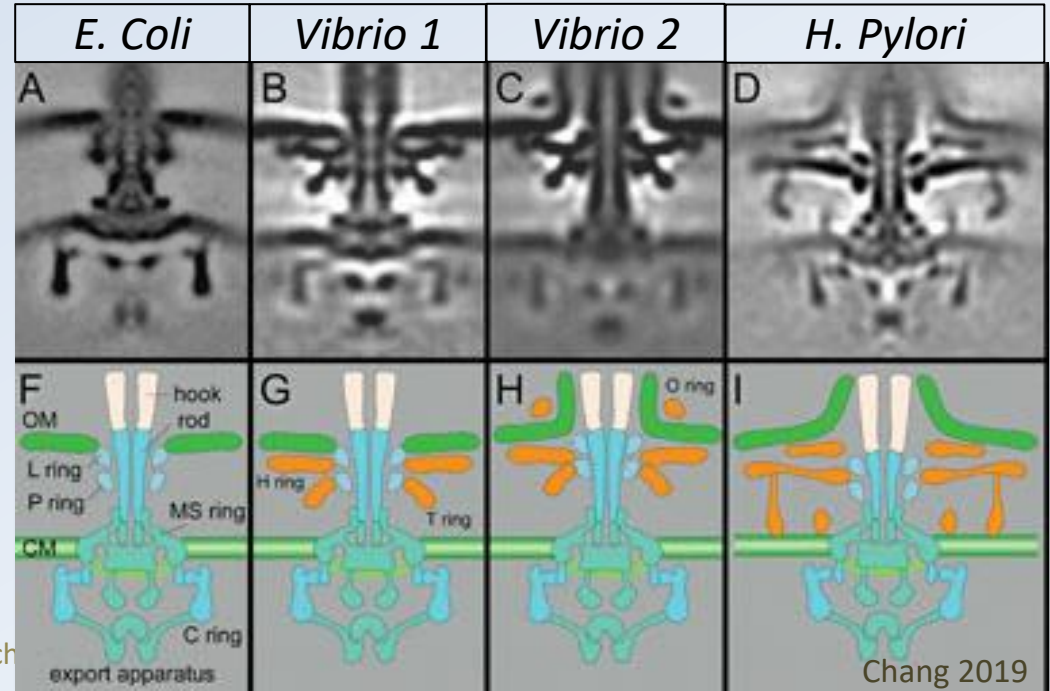
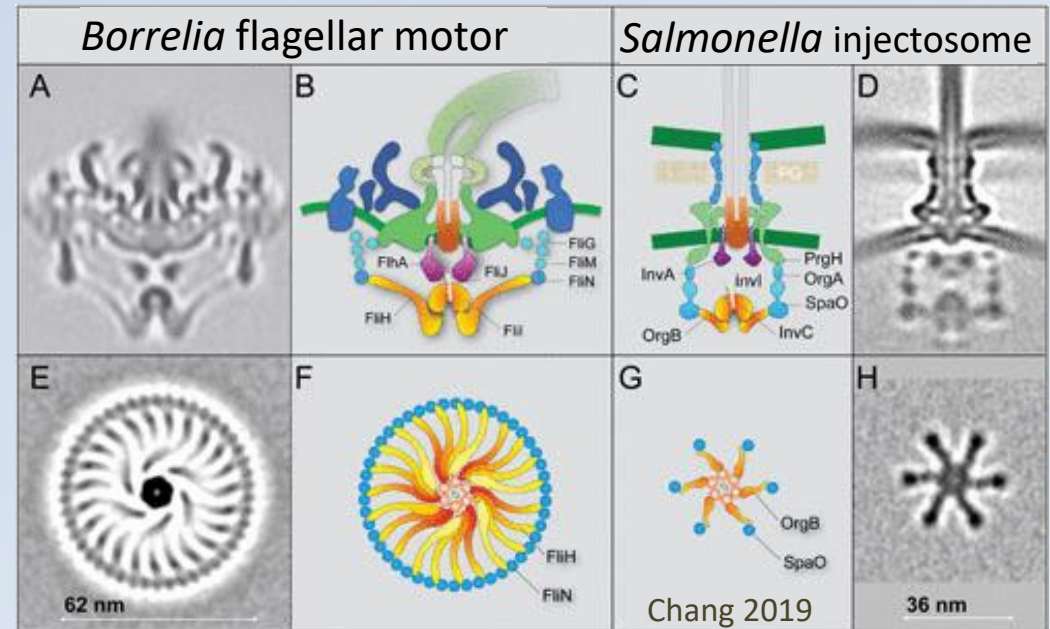
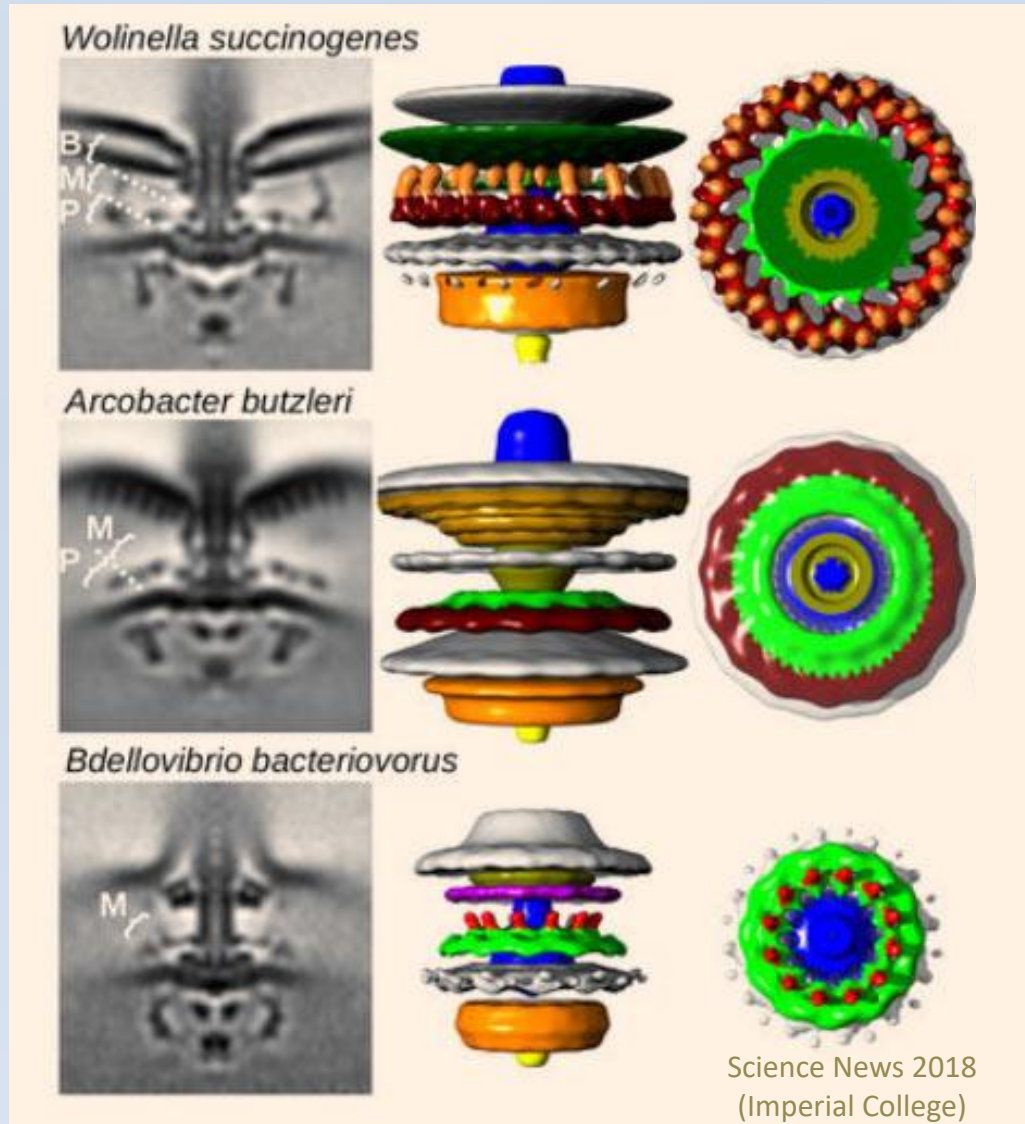


Stewart Lab
Callum Smits
ca 2018

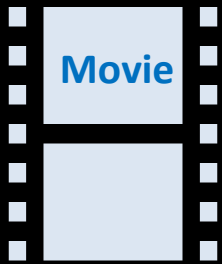
3/24/2023



Bacterial Flagellar Motors come in many variations



Bacterial Flagella



proton
flow

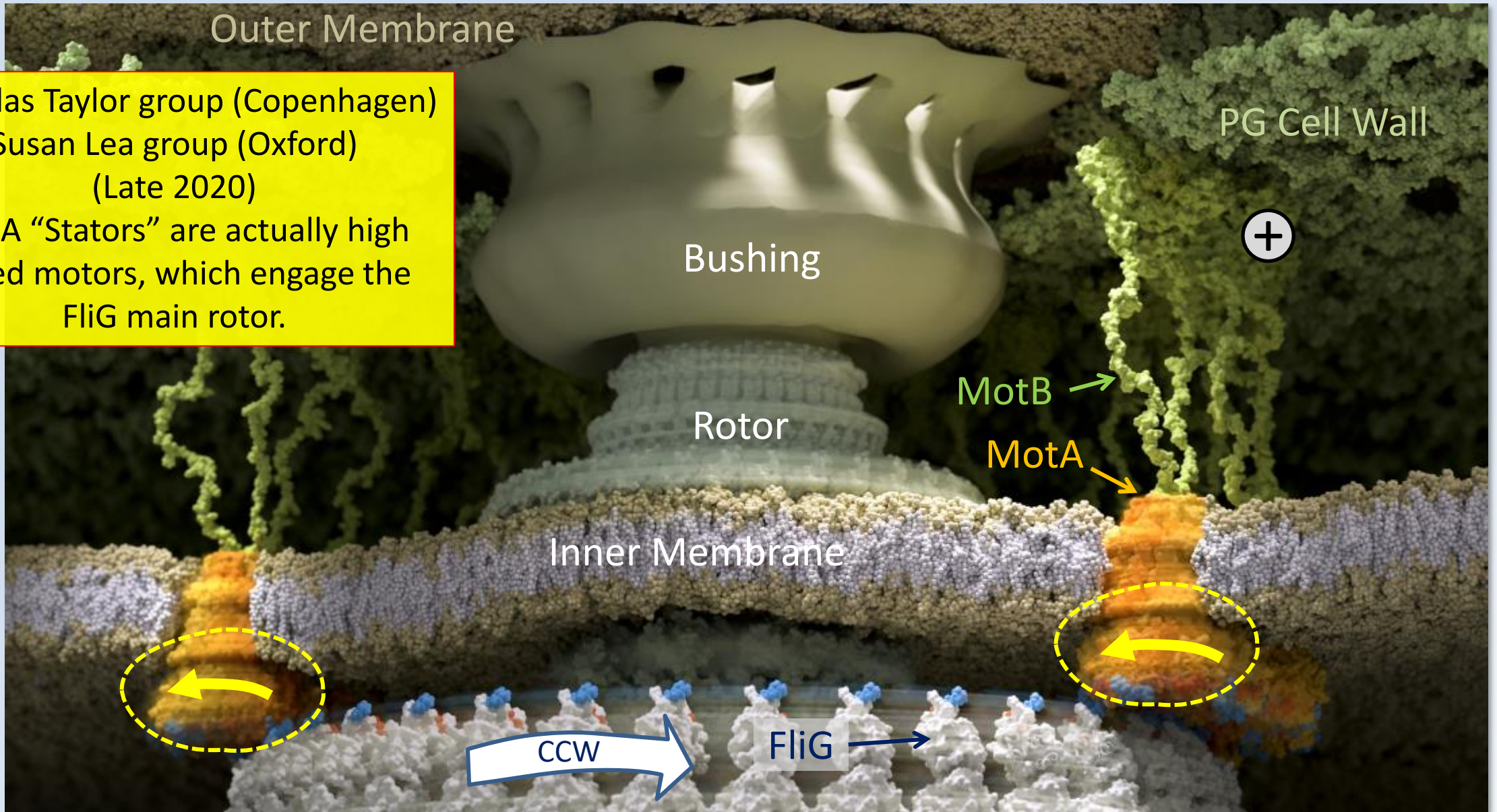
MotB

MotA

Biology Help
ca 2018



Nicholas Taylor group (Copenhagen)
Susan Lea group (Oxford)
(Late 2020)
MotA “Stators” are actually high speed motors, which engage the FliG main rotor.

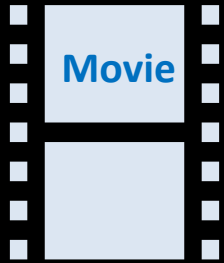
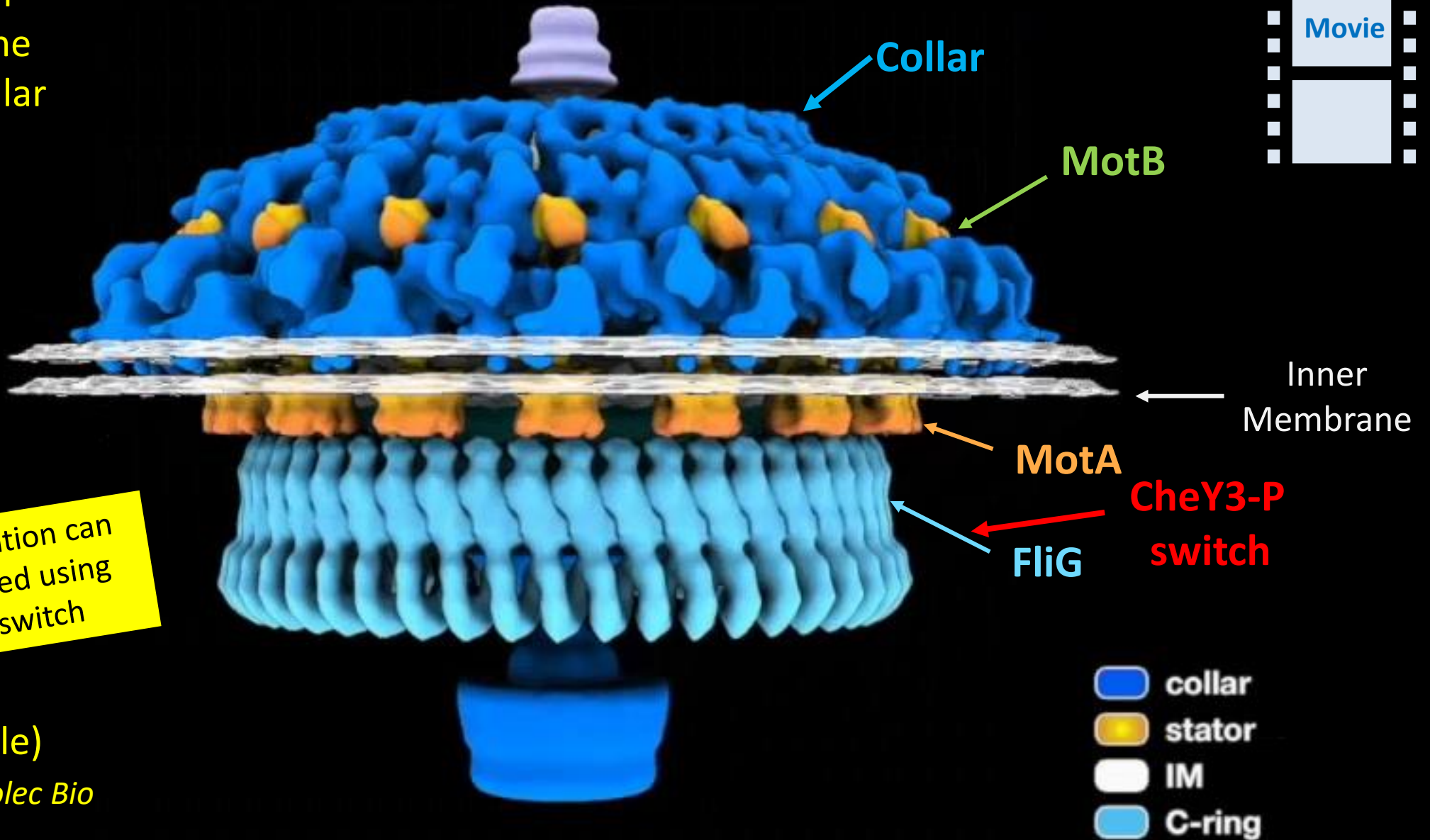


Molecular Mechanism for Rotational Switching in the Bacterial Flagellar Motor

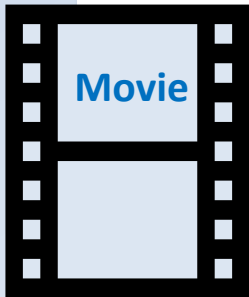
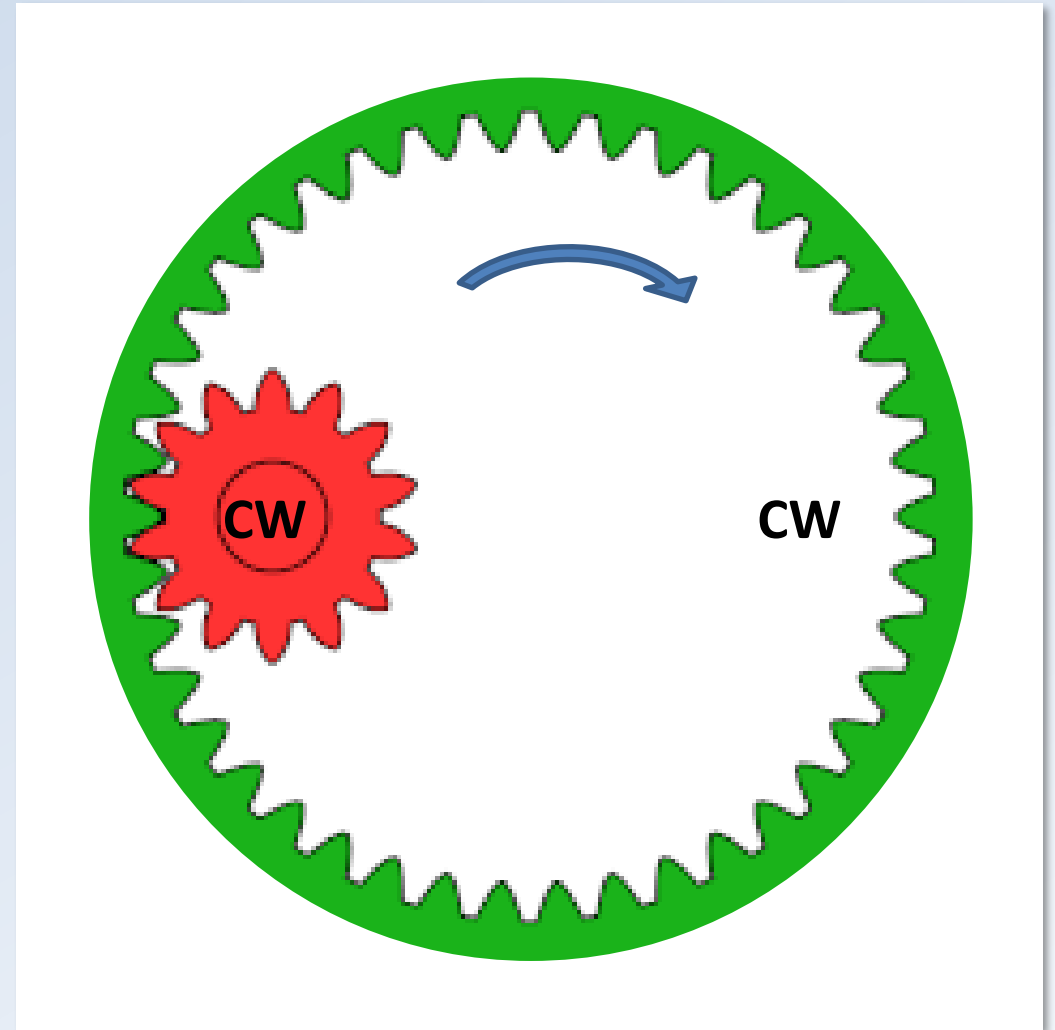
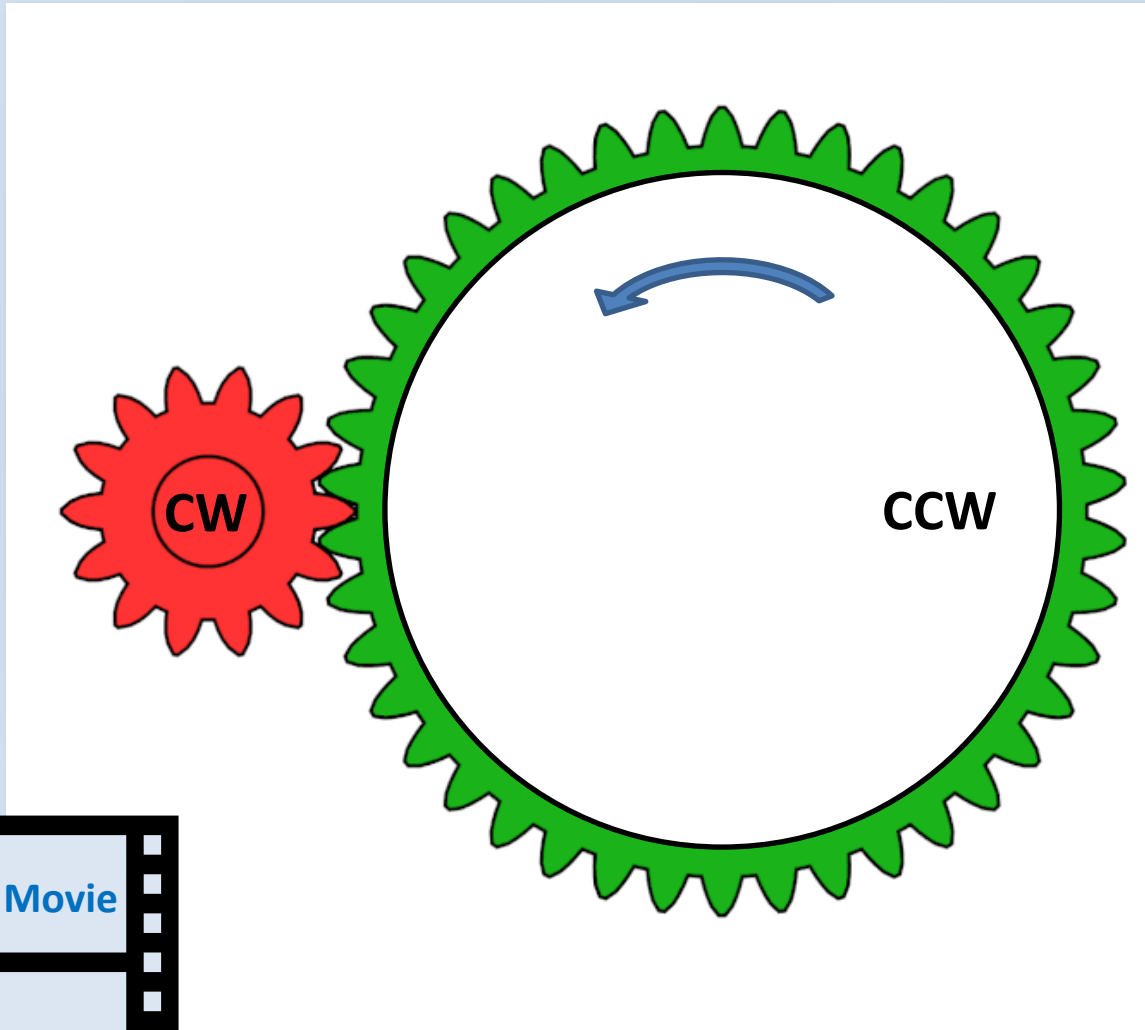
Motor from *B. burgdorferi* (Lyme disease spirochete)

Rotor rotation can be reversed using CheY3-P switch

Chang *et al* (Yale)
Nature Structural Molec Bio
Nov 2020

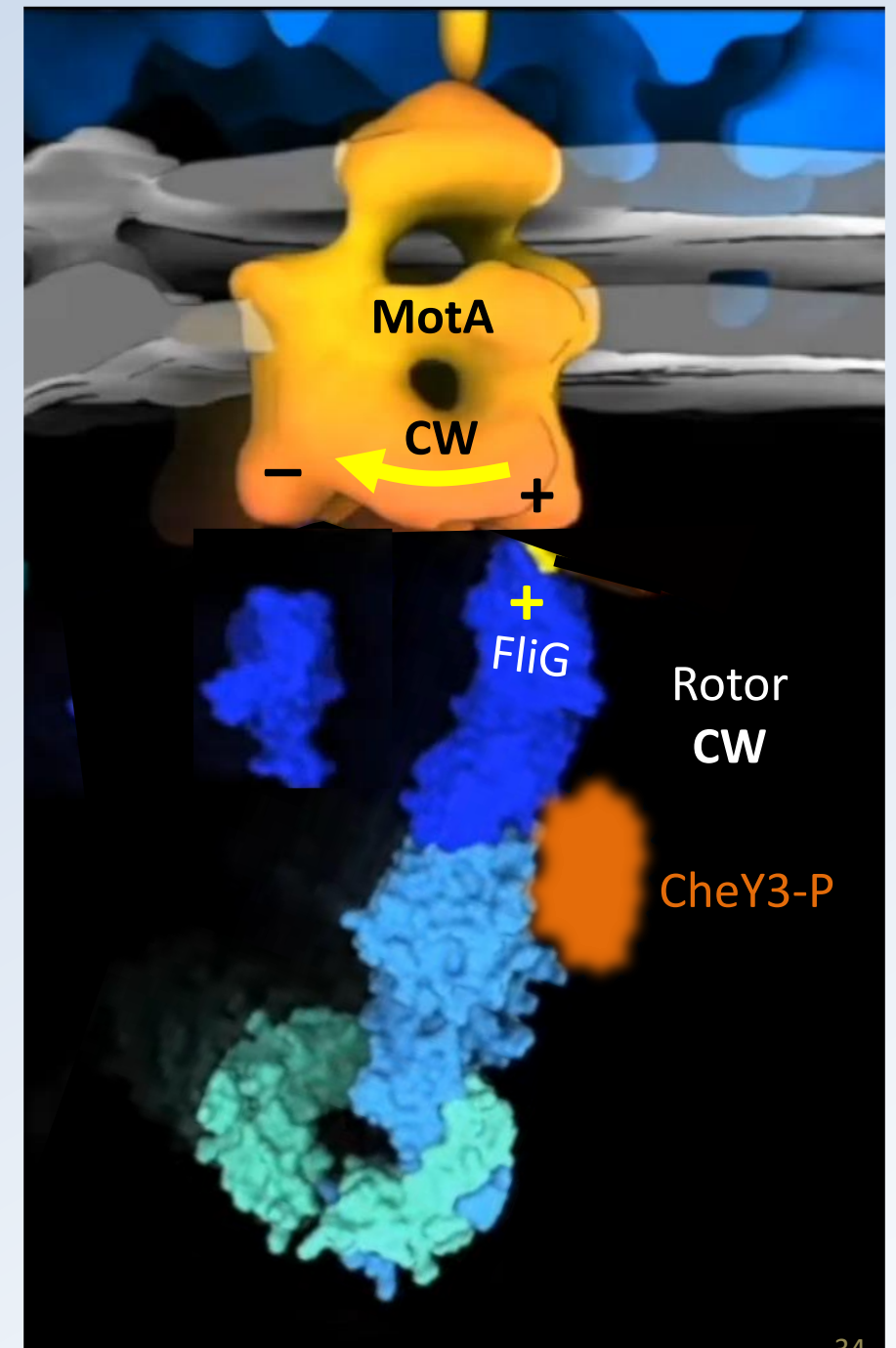
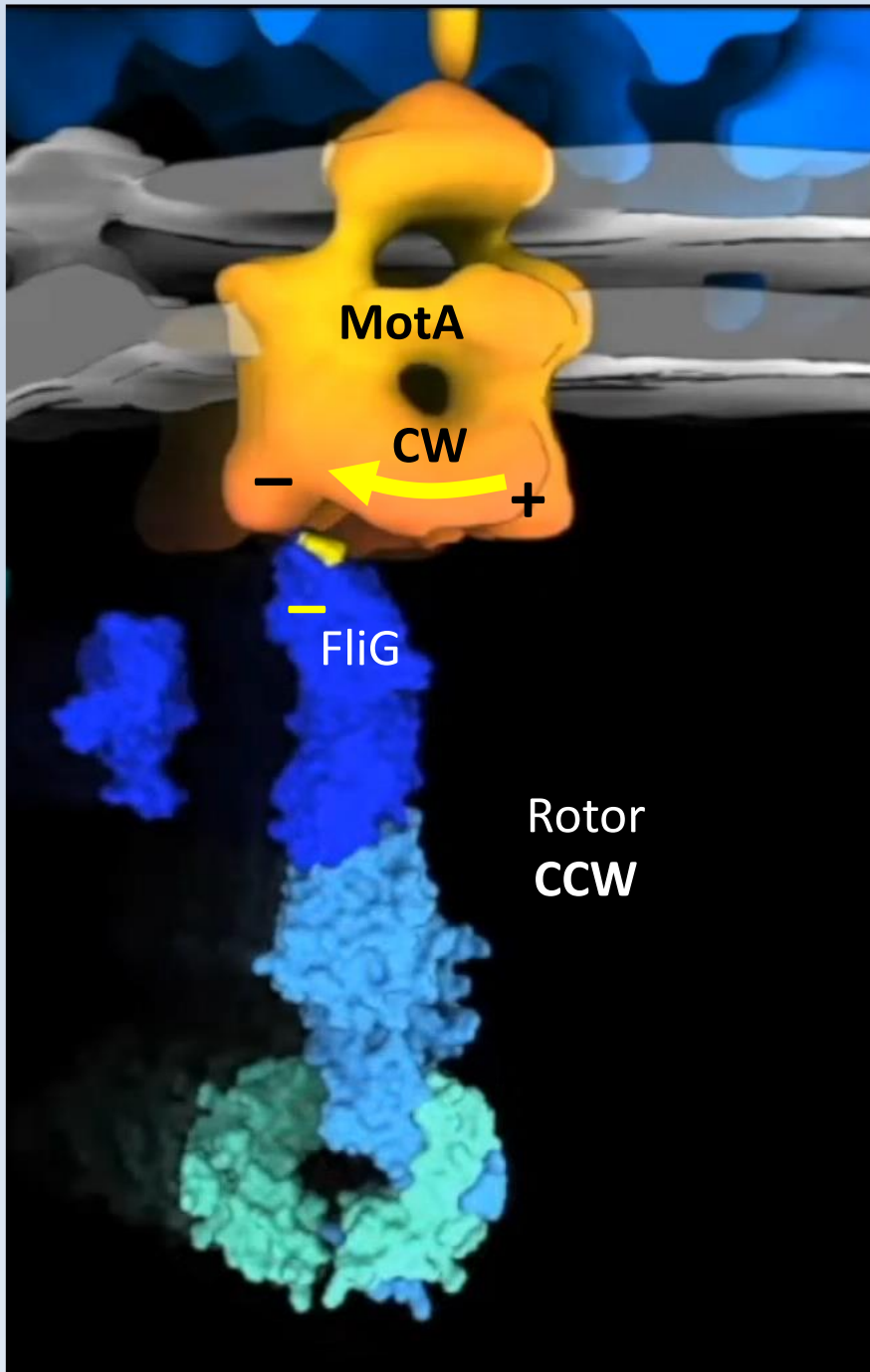


Shifting into Reverse



Shifting Gears into Reverse

CheY3-P bends the FliG rotor units outward so they engage the outer sides of the MotA motors



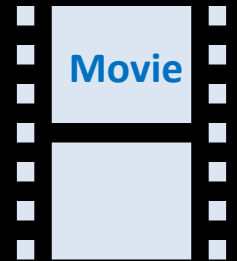
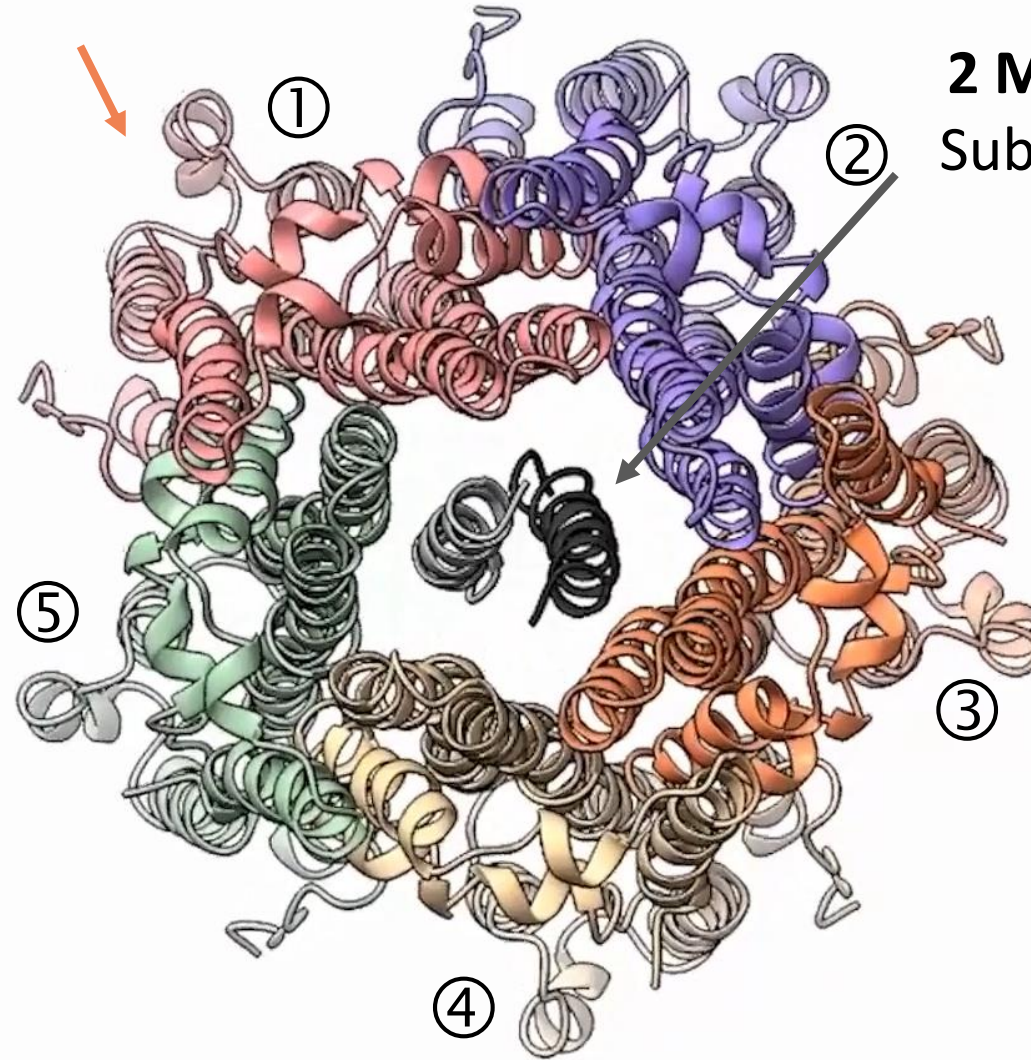
The Bacterial
Flagella
MotA/MotB
Motor in Action
from numerical
modeling

MotA Pentamer
rotates 36° for
each proton
flowing through

Santiveri et al, Cell **183** (2020)
*Structure and Function of
Stator Units of the BFM
Video S5*

5 MotA Subunits

2 MotB
Subunits





INFORMATIONAL

Molecular Machines

Selected List of Molecular Machines

I. MOLECULAR MACHINES THAT SCIENTISTS HAVE ARGUED SHOW IRREDUCIBLE COMPLEXITY

1. BACTERIAL FLAGELLUM

The flagellum is a rotary motor in bacteria that drives a propeller to spin, much like an outboard motor, powered by ion flow to drive rotary motion. Capable of spinning up to 100,000 rpm,¹³ one paper in *Trends in Microbiology* called the flagellum “an exquisitely engineered chemi-osmotic nanomachine; nature’s most powerful rotary motor, harnessing a transmembrane ion-motive force to drive a filamentous propeller.”¹⁴ Due to its motor-like structure and internal parts, one molecular biologist wrote in the journal *Cell*, “[m]ore so than other motors, the flagellum resembles a machine designed by a human.”¹⁵ Genetic knockout experiments have shown that the *E. coli* flagellum is irreducibly complex with respect to its approximately 35 genes.¹⁶ Despite the fact that this is one of the best studied molecular machines, a

Questions?

Building Cells from a DNA Blueprint

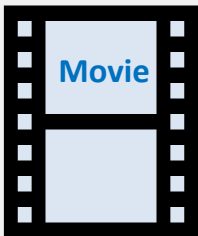
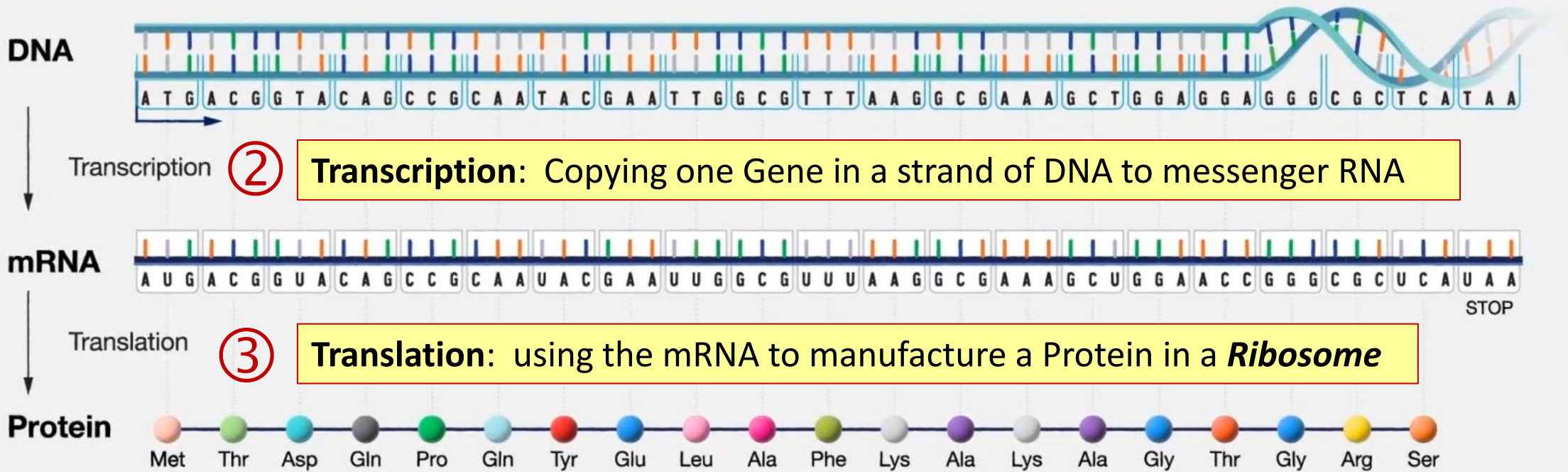




The Central Dogma

Central Dogma

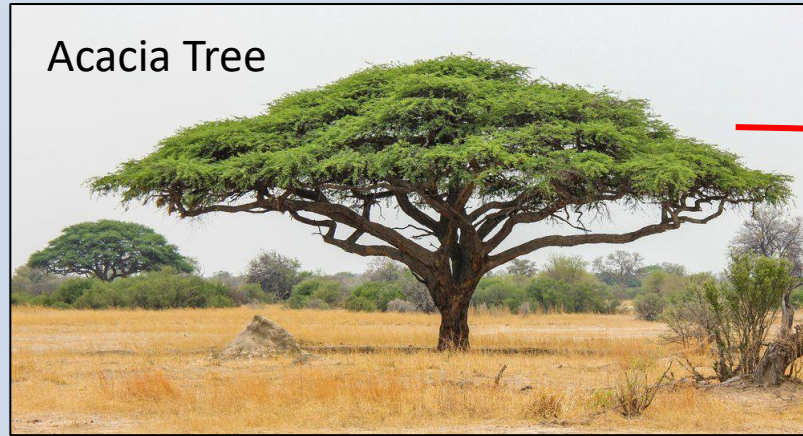
① **Replication:** Duplicating DNA when cells divide.



Structure of DNA and RNA



Nucleic Acids

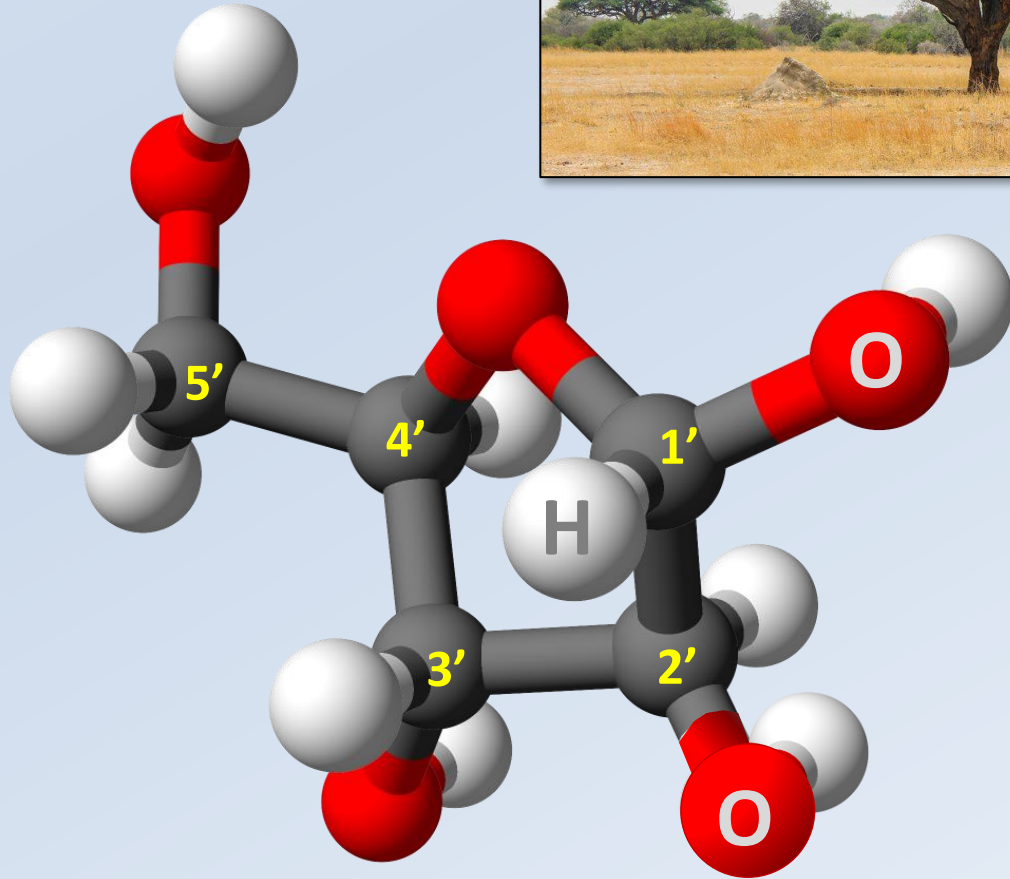


A similar 5 carbon sugar:

arabinose

ribose

Diagram showing the relationship between arabinose and ribose. The word 'arabinose' is above 'ribose'. Blue arrows point from 'arabinose' to 'ribose', with a blue 'X' over the 'a' in 'arabinose'. A blue bracket under 'ribose' is connected to a blue bracket under 'arabinose'.

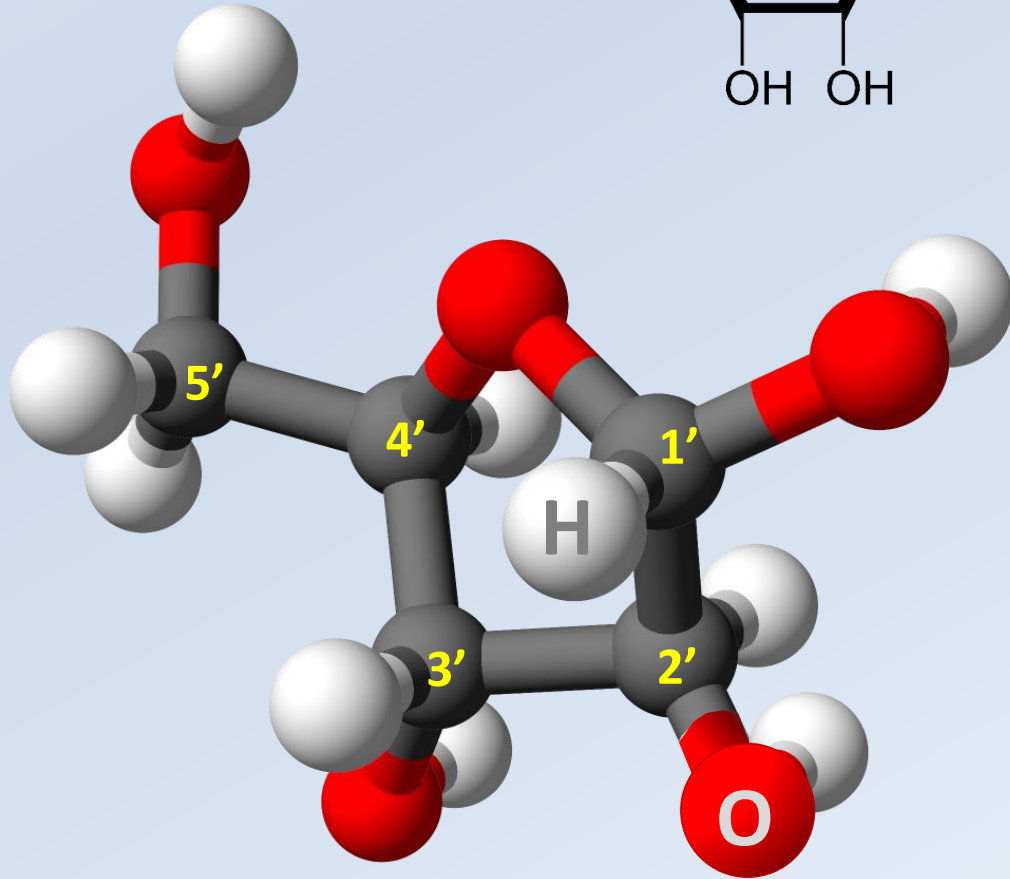
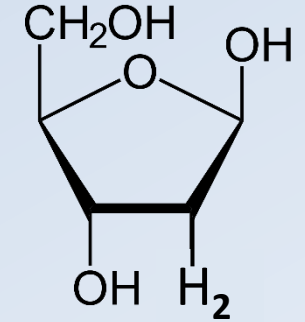
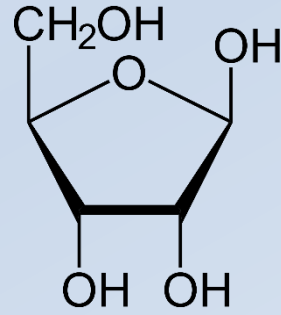


Ribose

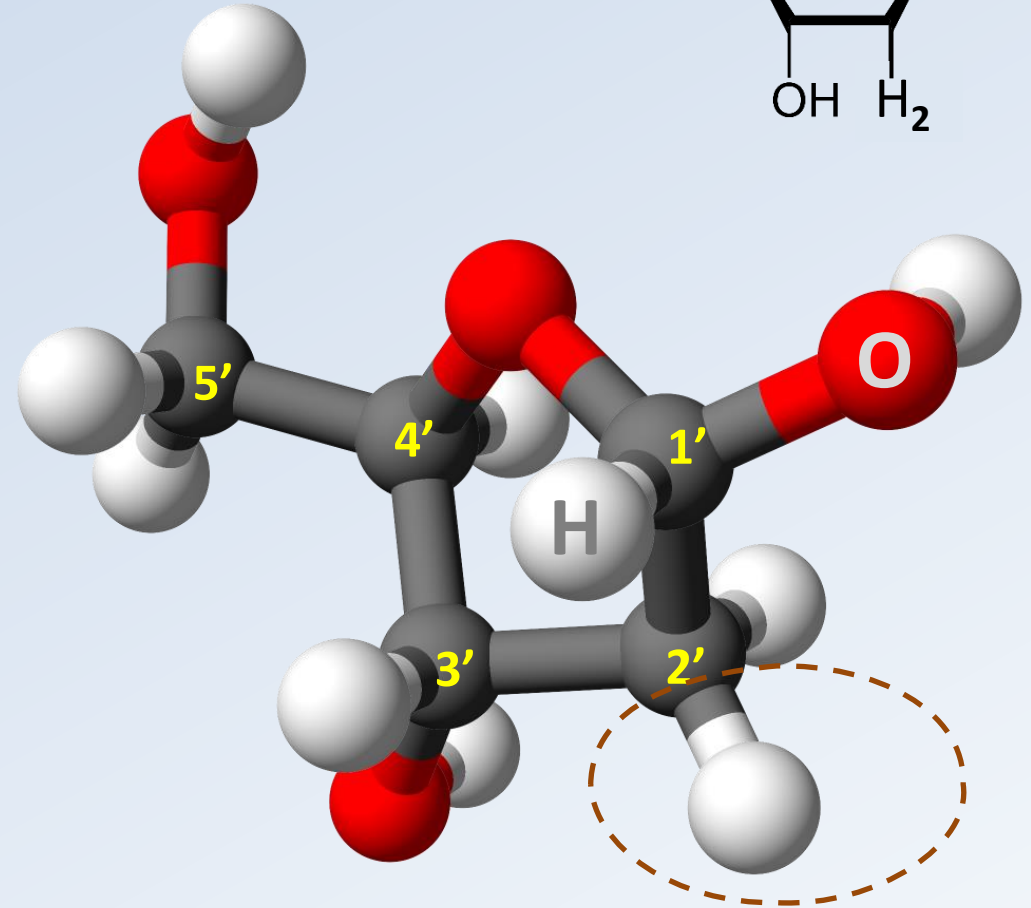
Named after arabinose, in turn named after the Gum Arabic in which it was found.



Nucleic Acids



Ribose

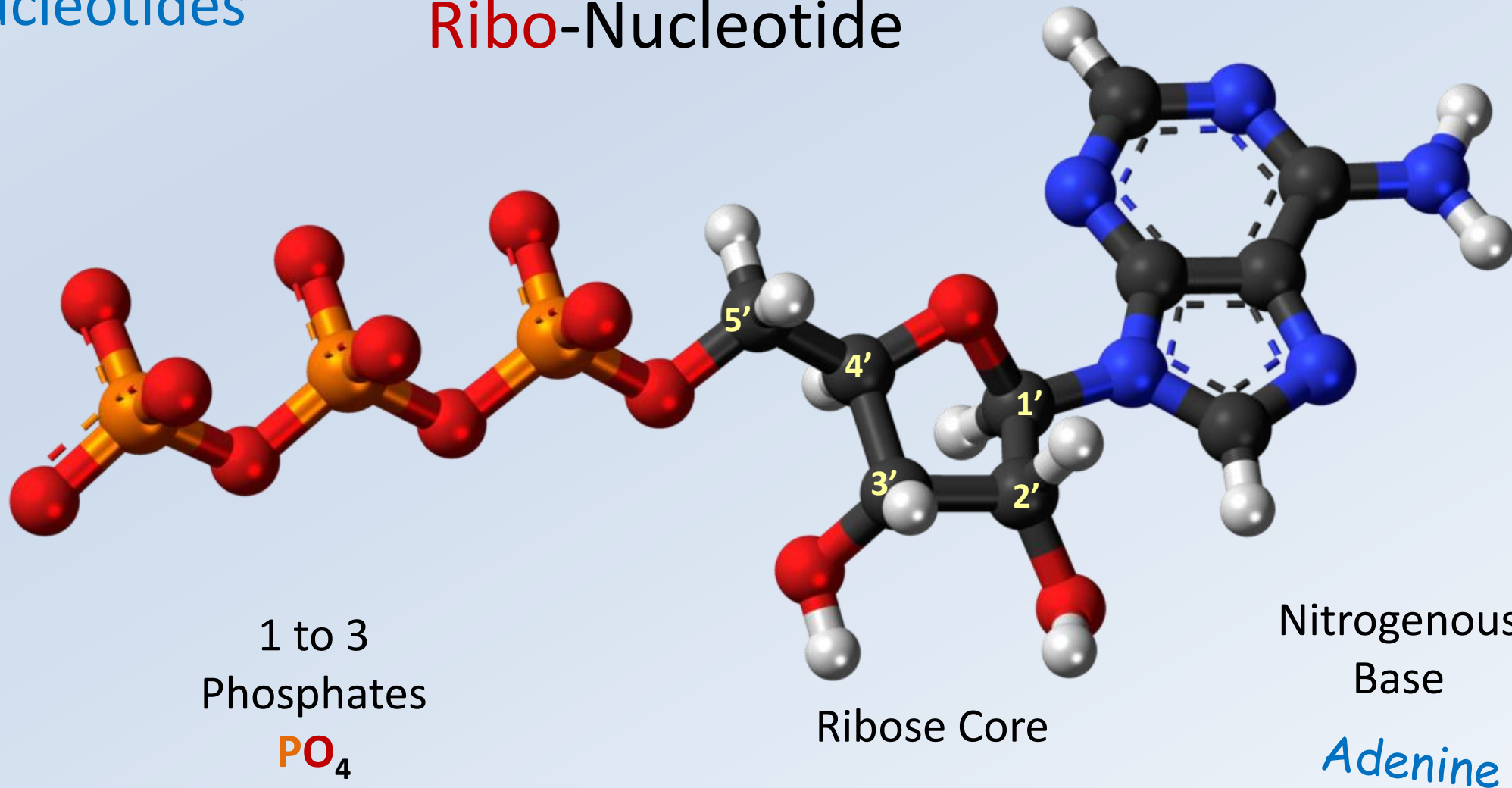


DeOxy-Ribose



Nucleotides

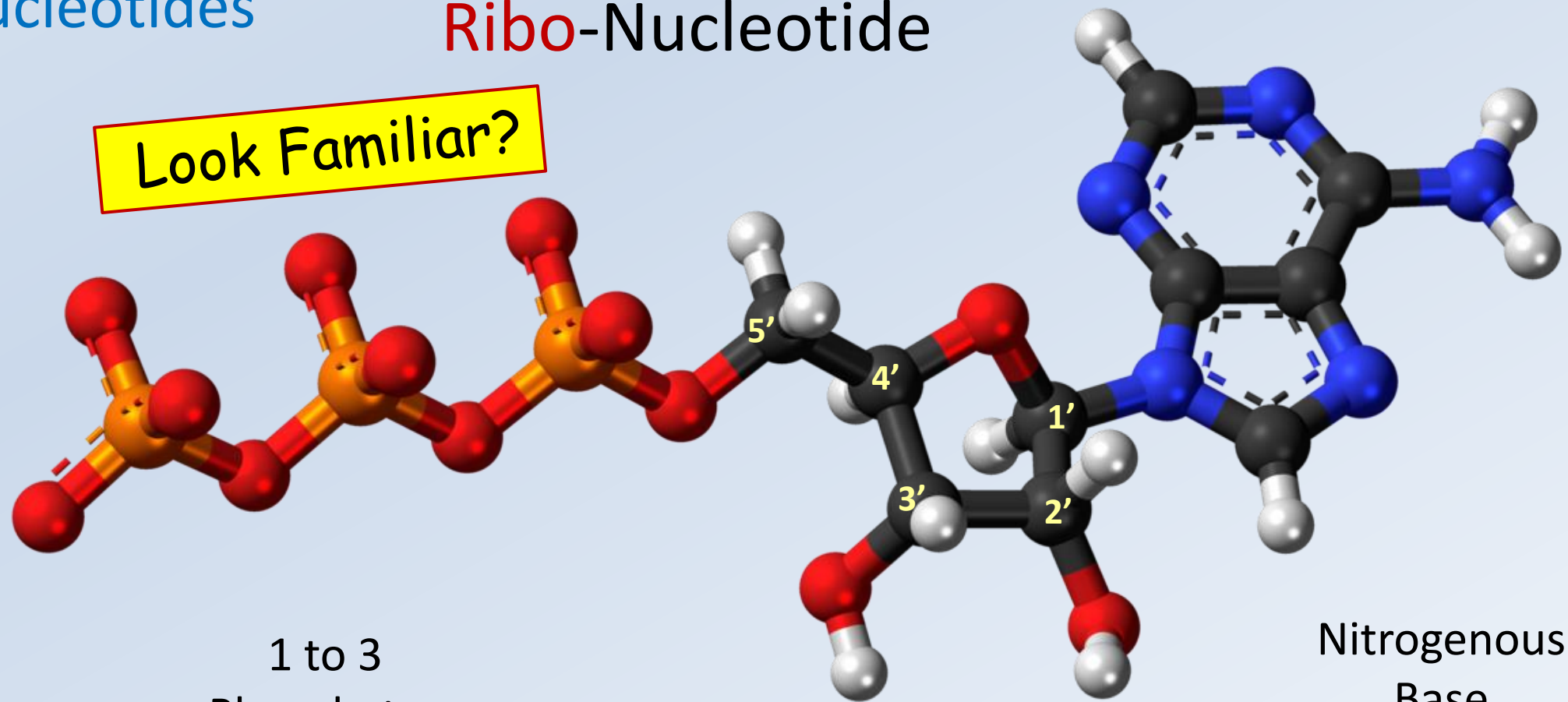
Ribo-Nucleotide



Nucleotides

Ribo-Nucleotide

Look Familiar?



1 to 3
Phosphates



Ribose Core

Nitrogenous
Base

Adenine

RNA
Component

“Adenosine Tri-Phosphate”

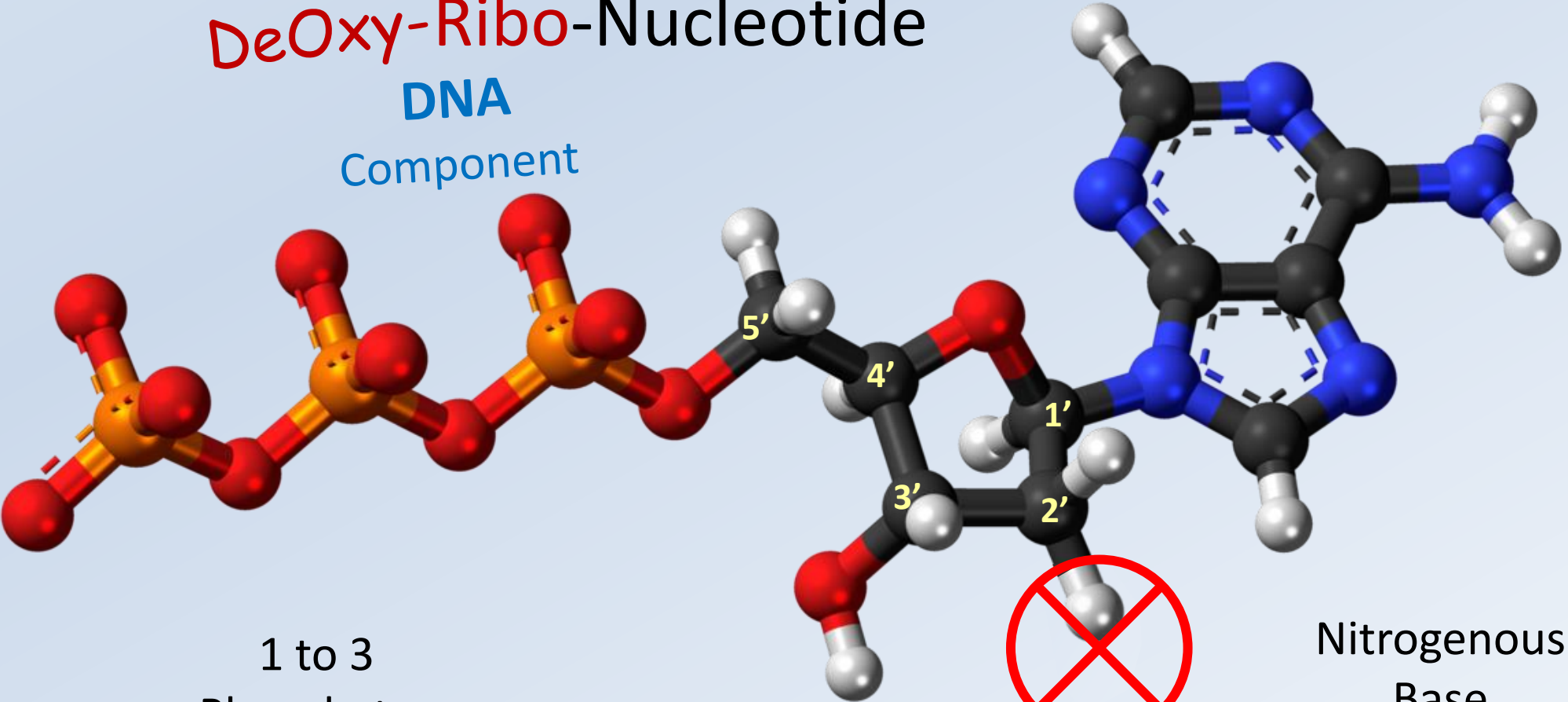
ATP



Nucleotides

DeOxy-Ribo-Nucleotide

DNA
Component



1 to 3
Phosphates
 PO_4

DeOxy-Ribose Core

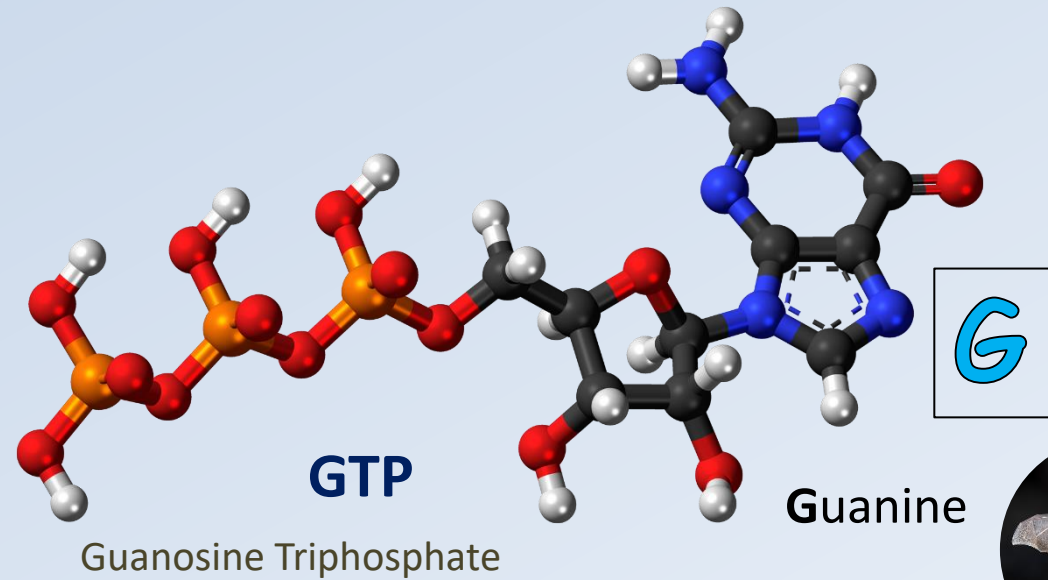
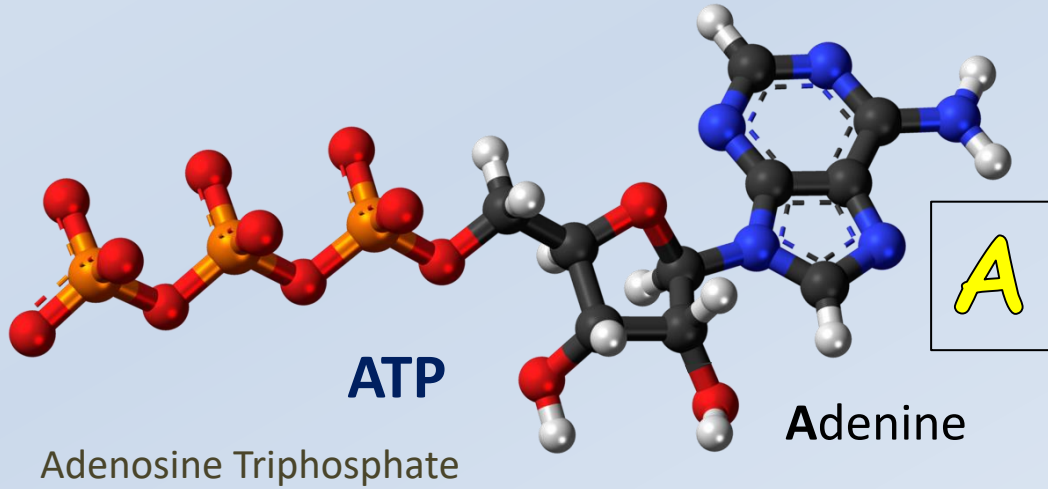
Nitrogenous
Base

Adenine

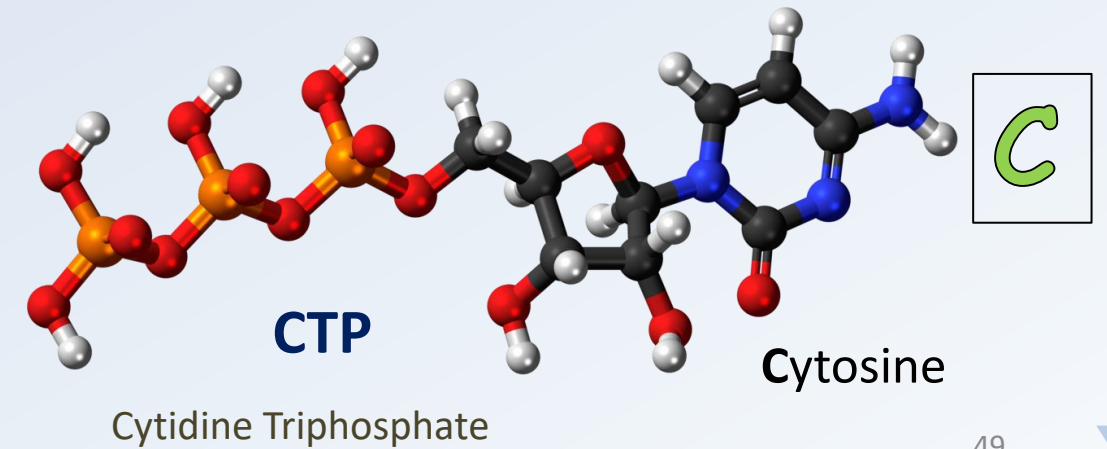
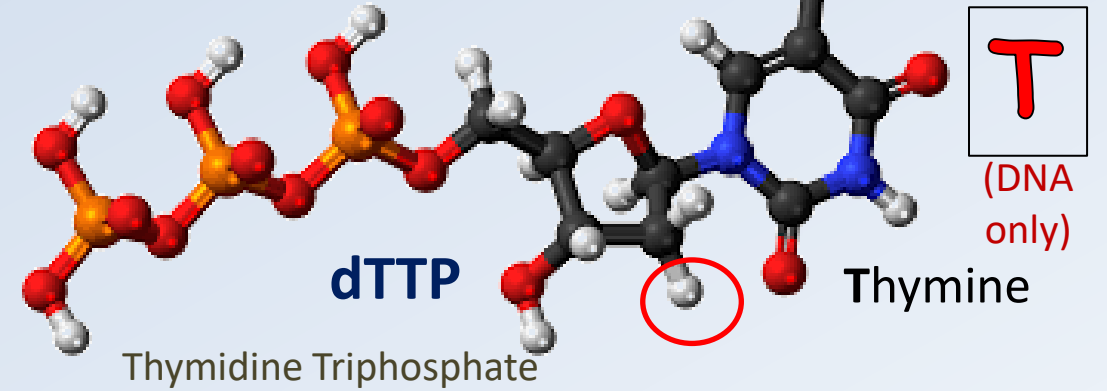
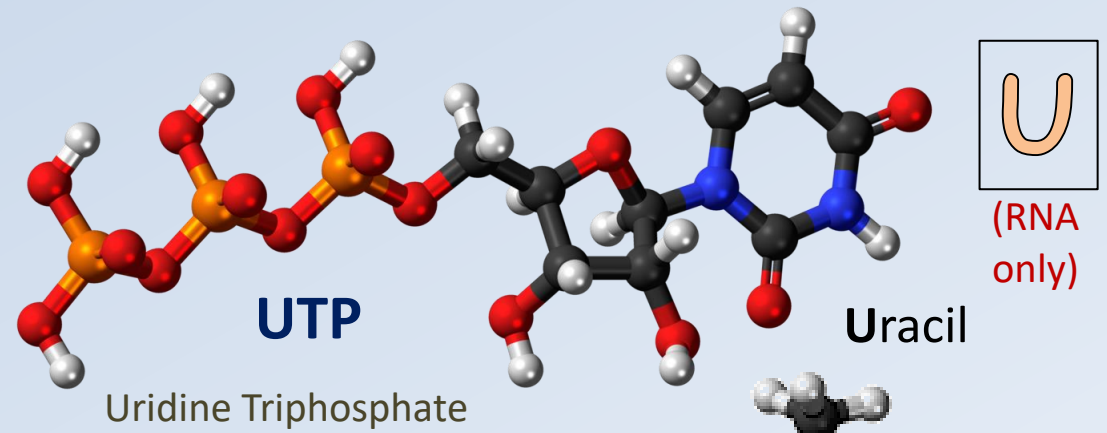
dATP



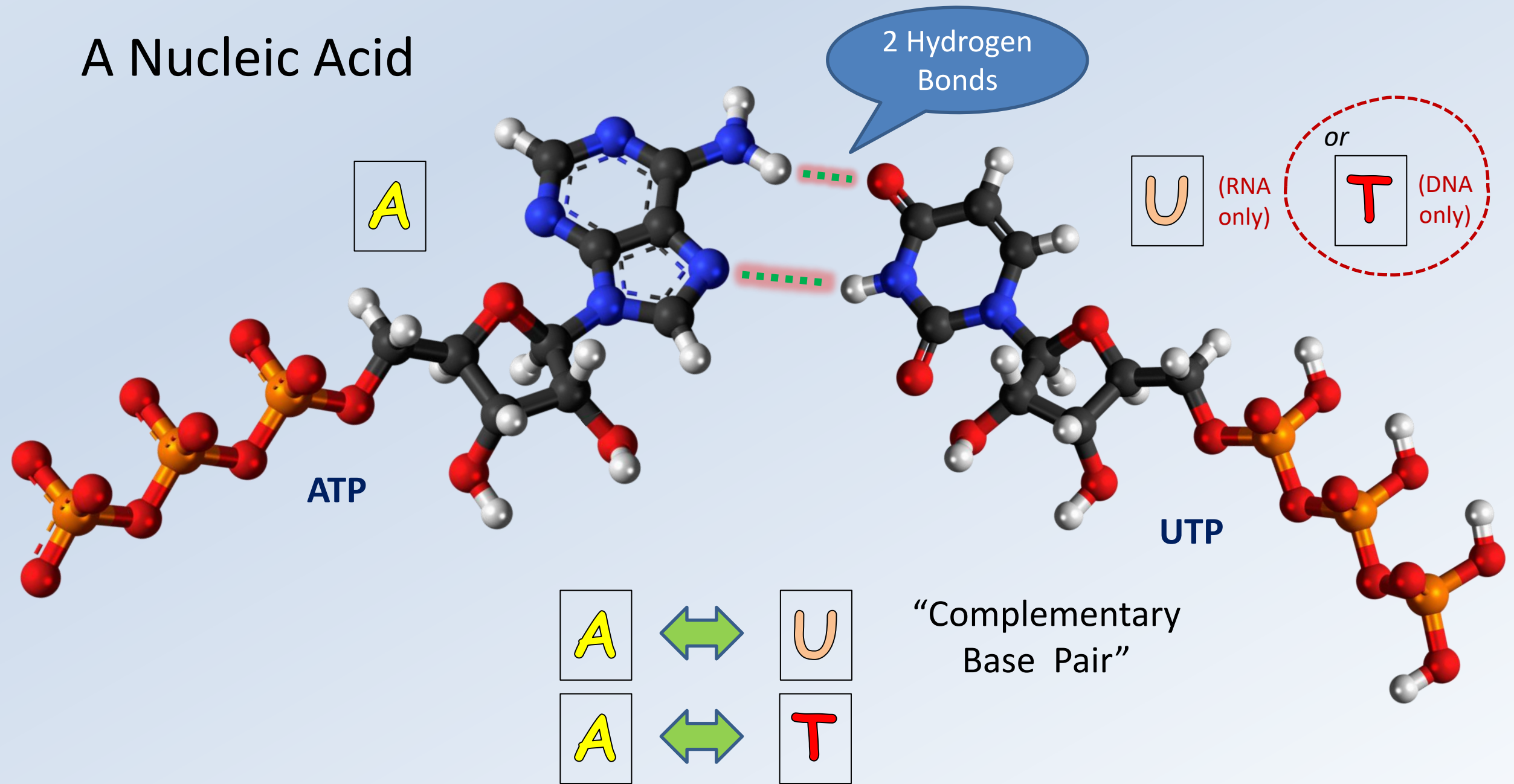
5 4 Nucleotides



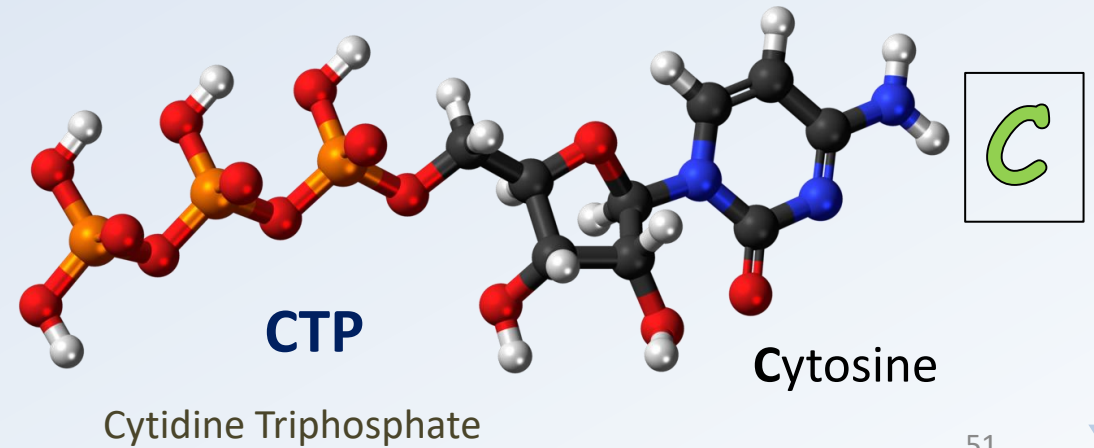
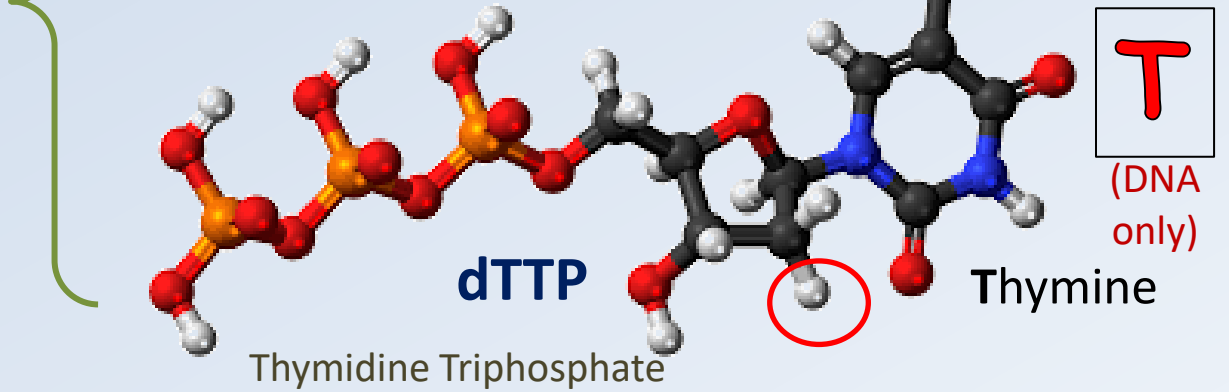
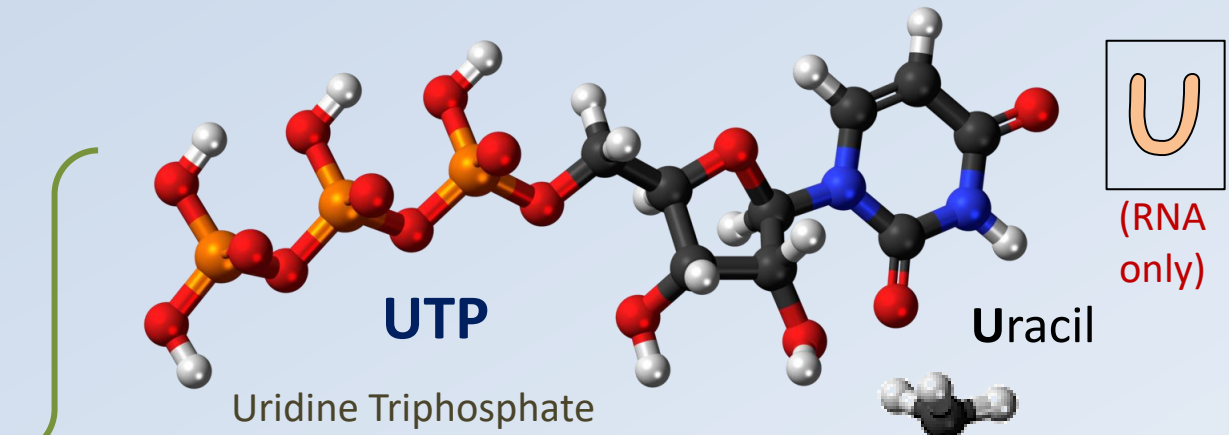
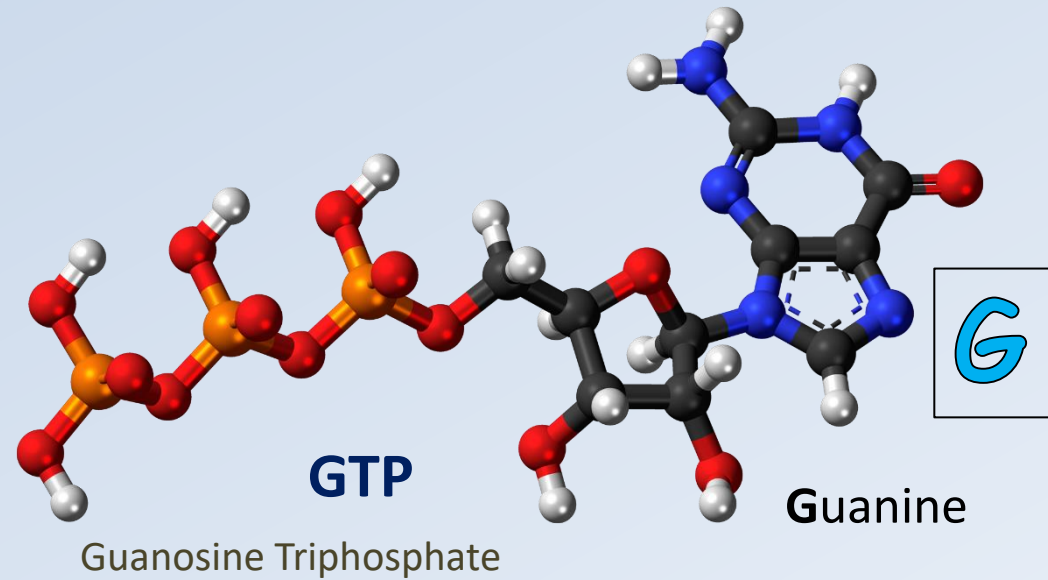
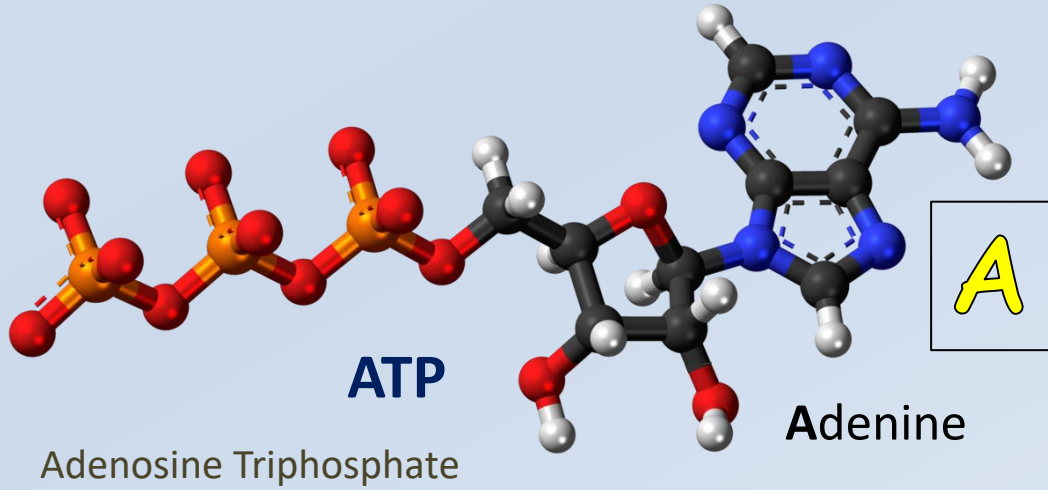
chines 4



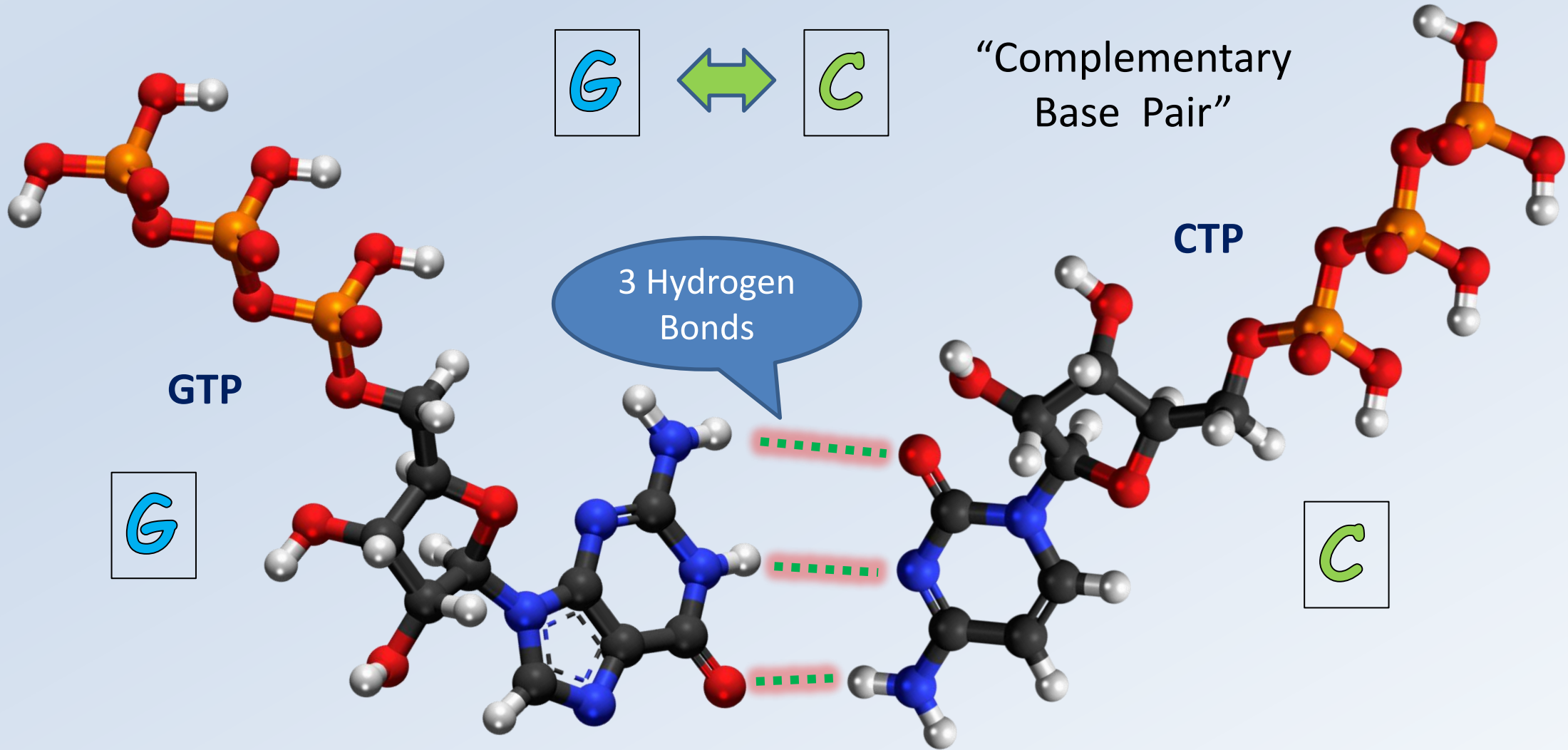
A Nucleic Acid



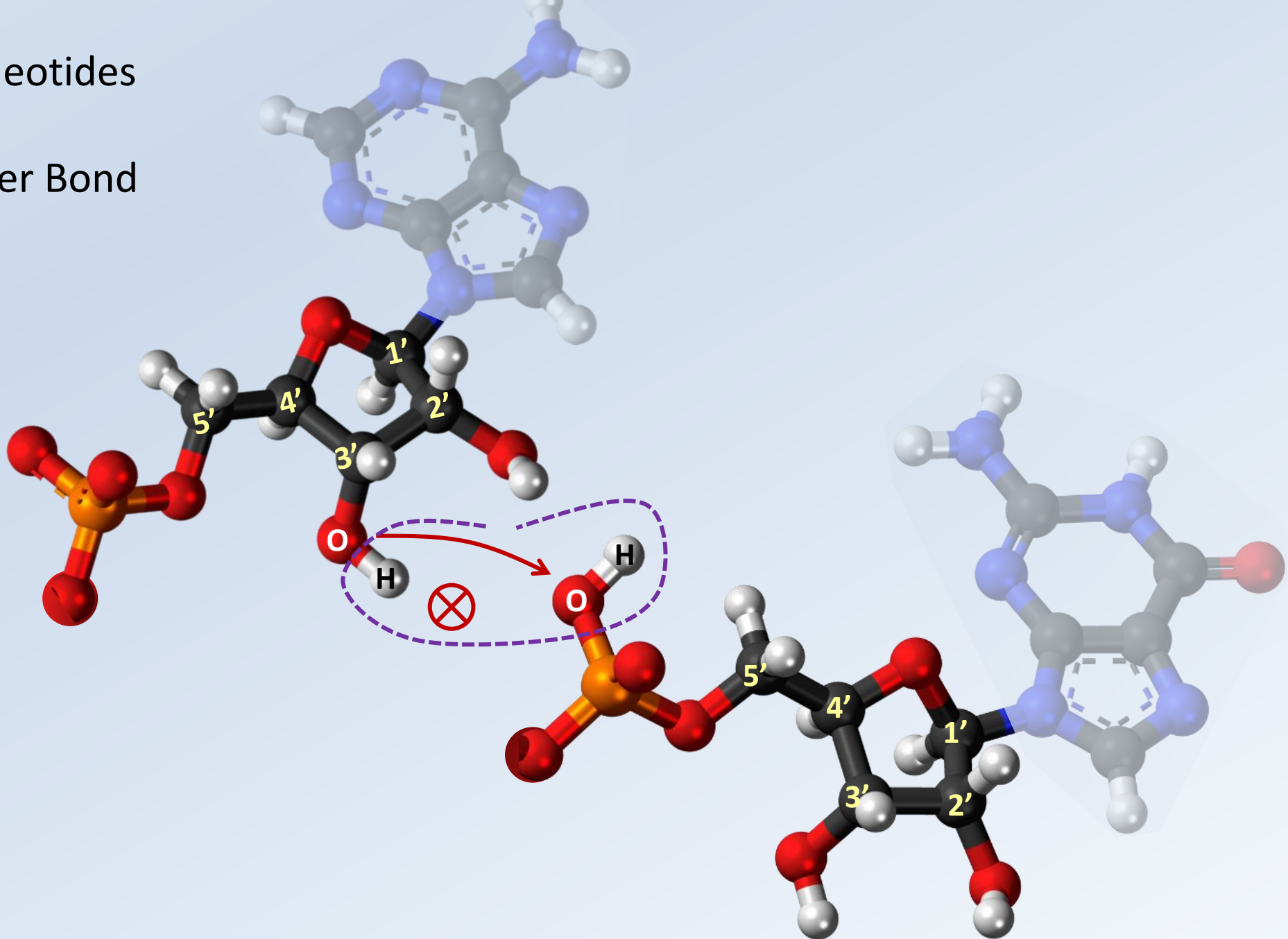
5 4 Nucleotides



Another Nucleic Acid



Joining 2 Nucleotides with a Phosphodiester Bond



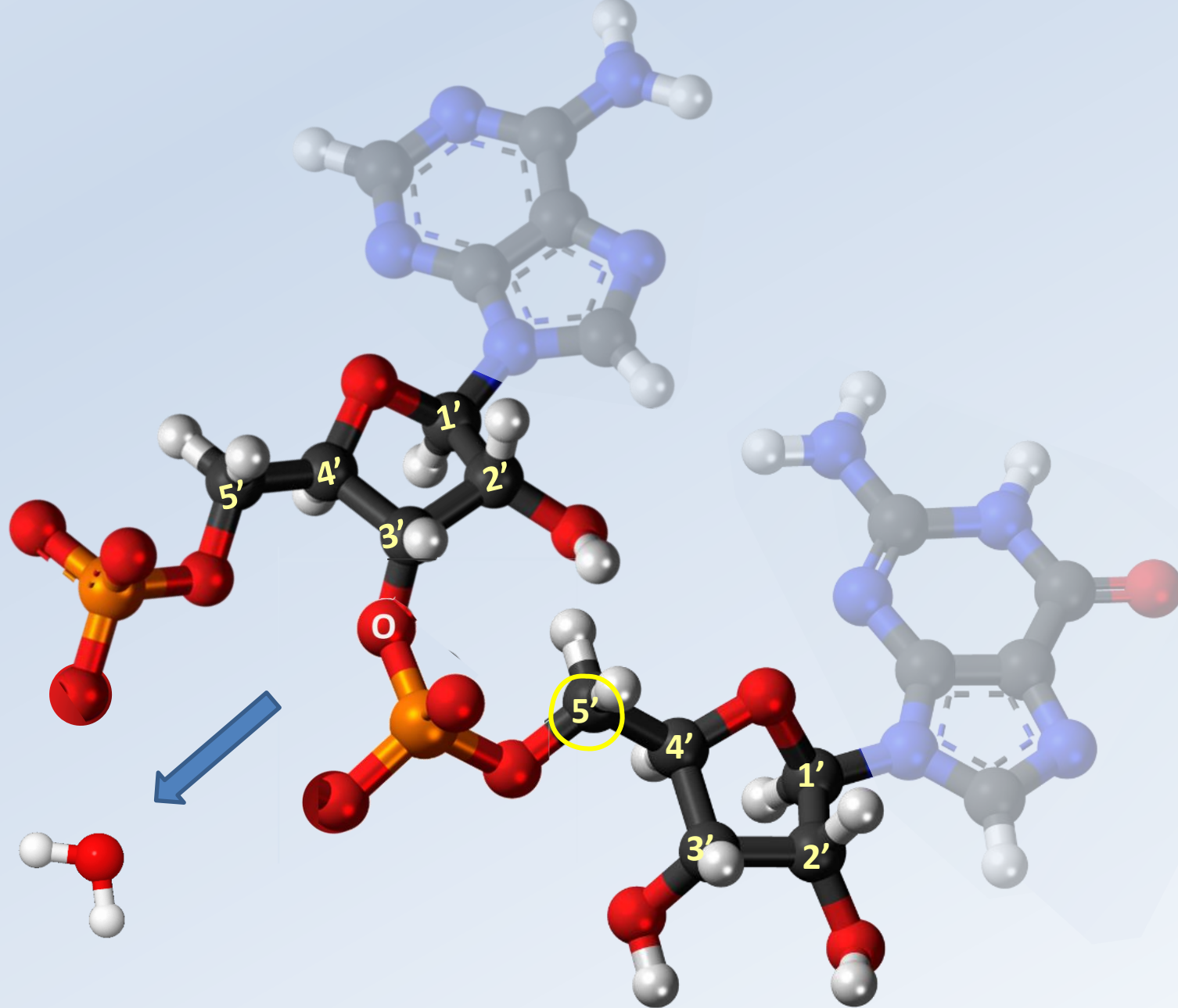
Joining 2 Nucleotides with a Phosphodiester Bond

Polymerization

Note:


3' end of the
first nucleotide
attaches to **5'** end
of the other

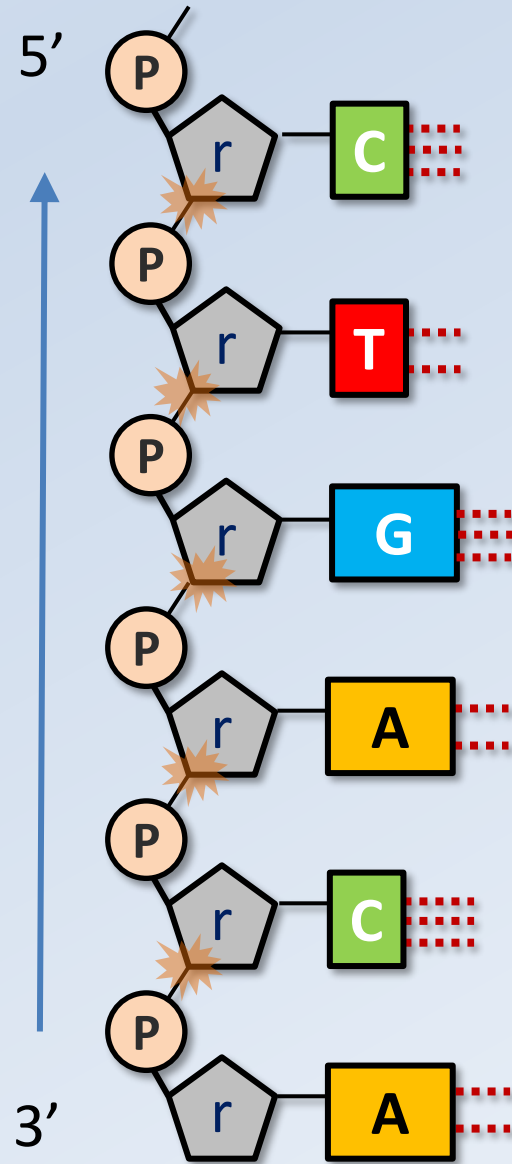
Dehydration
Reaction



Nucleotide Polymerization

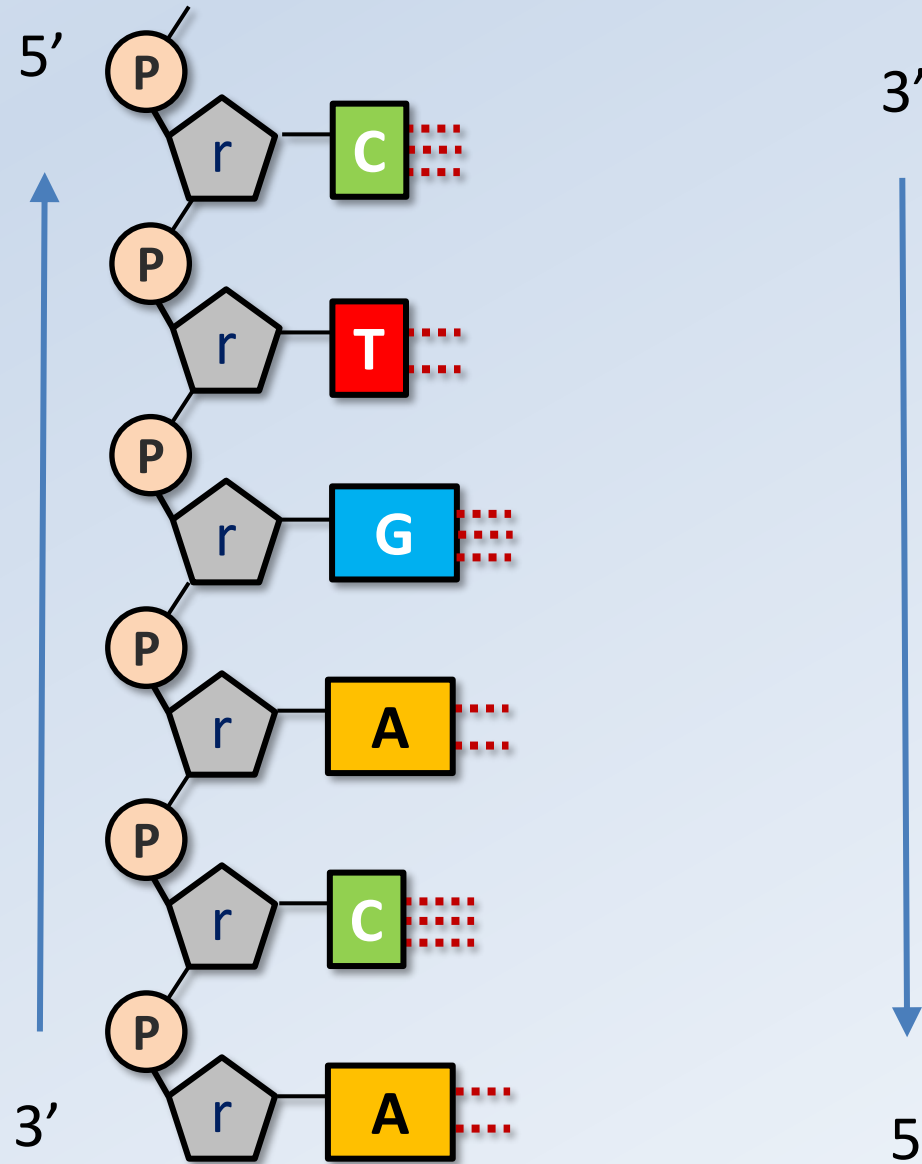
Single Strand
of RNA or DNA

Phosphodiester
Bonds 



Nucleotide Polymerization

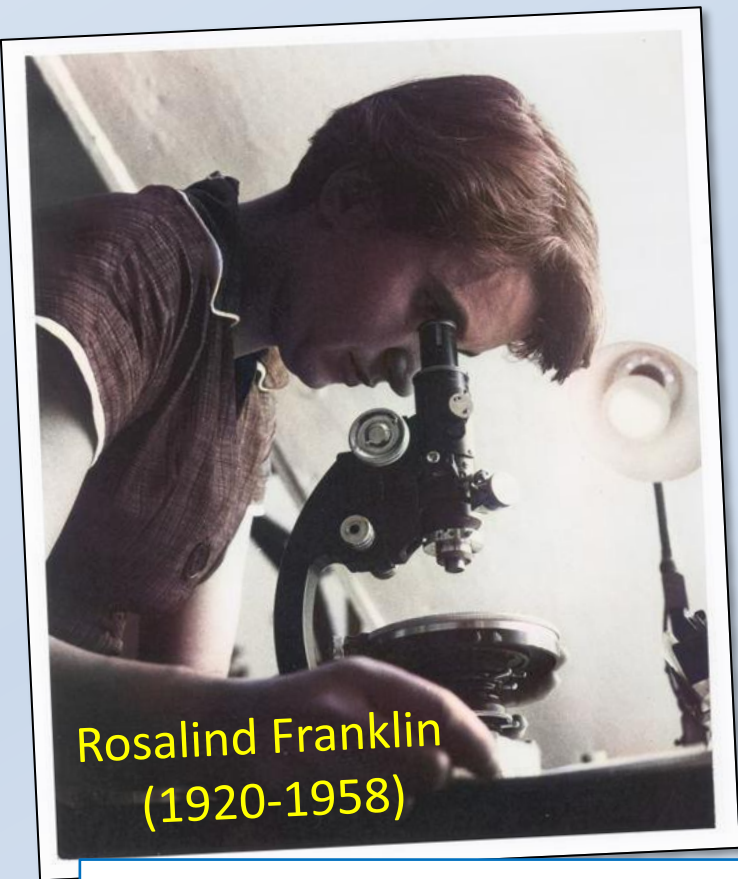
Single Strand
of RNA or DNA



An Exact
Complementary
Strand will
spontaneously
find its position
and stick!

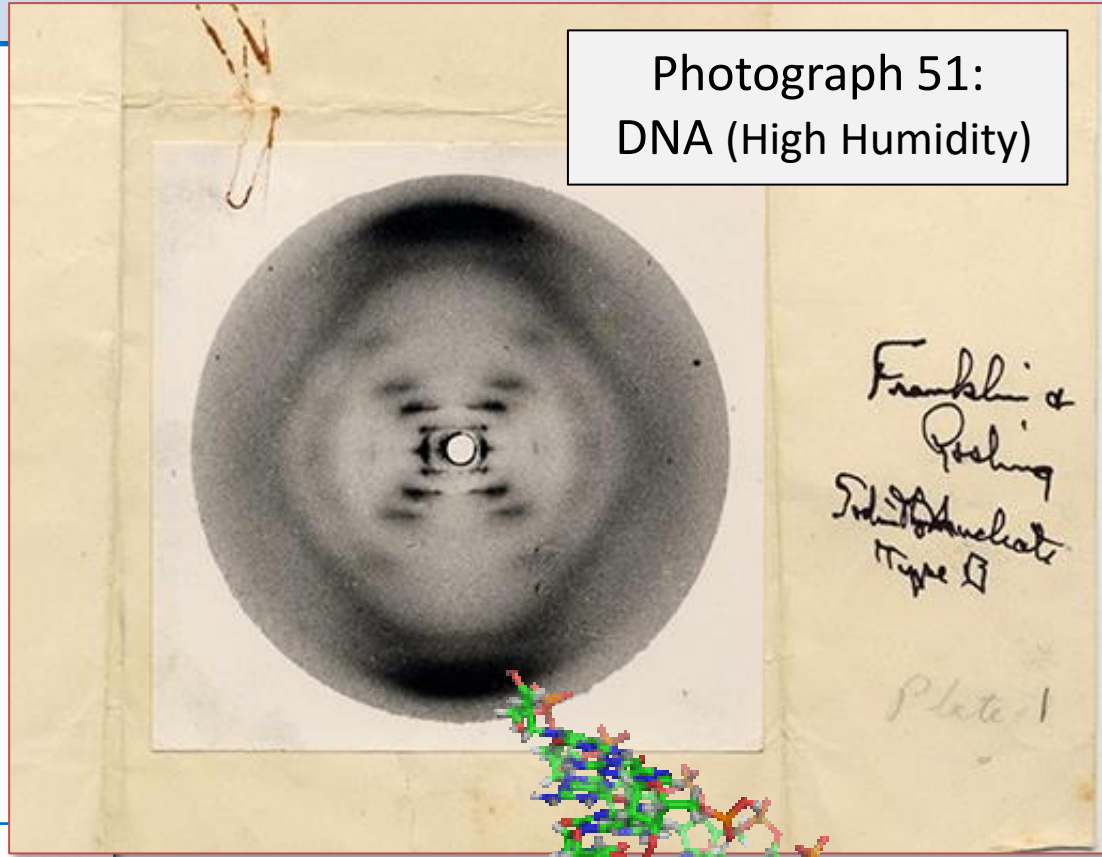
We now have
Double
Stranded DNA
or RNA



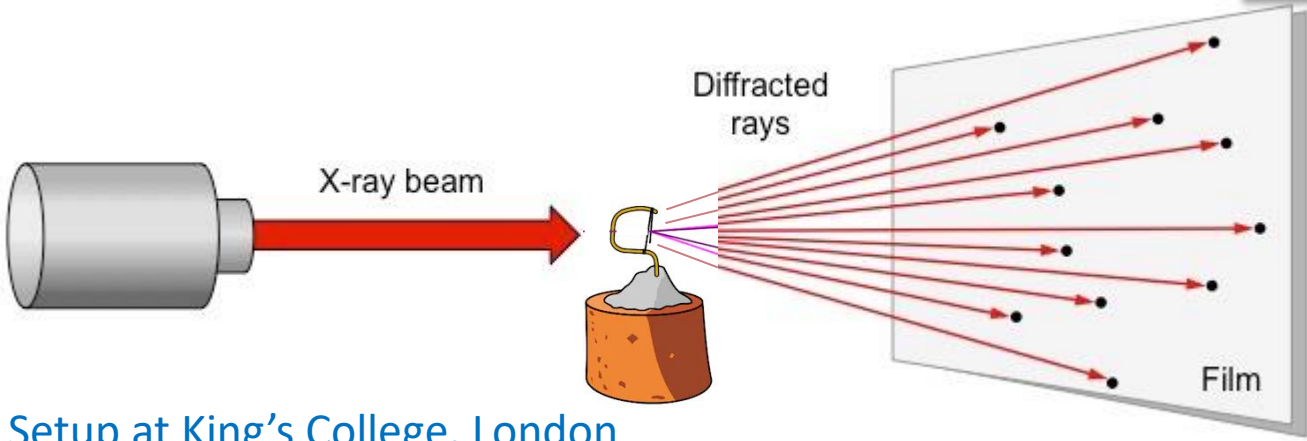


Rosalind Franklin
(1920-1958)

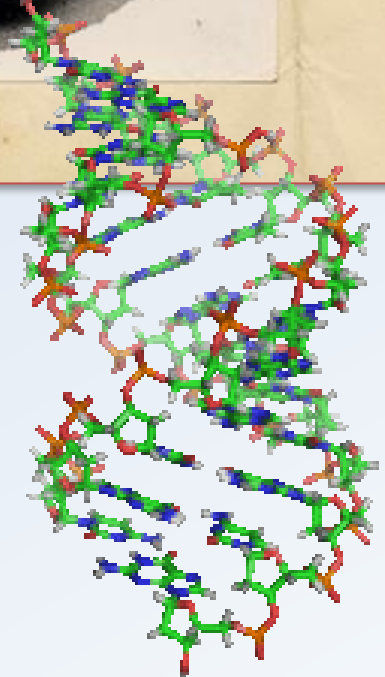
Key to the Discovery of the Structure of DNA May 1952

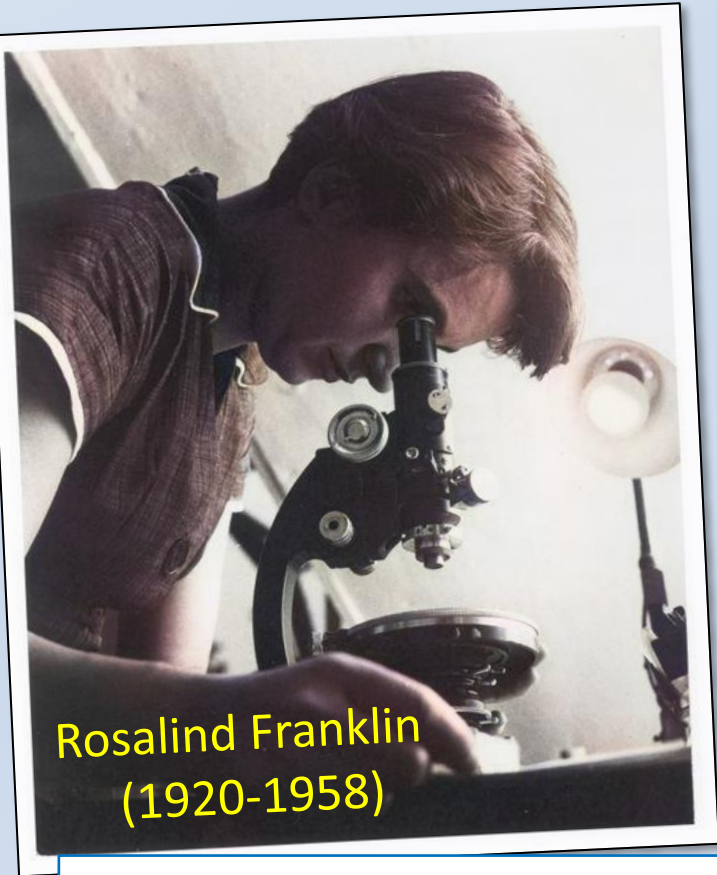


Photograph 51:
DNA (High Humidity)

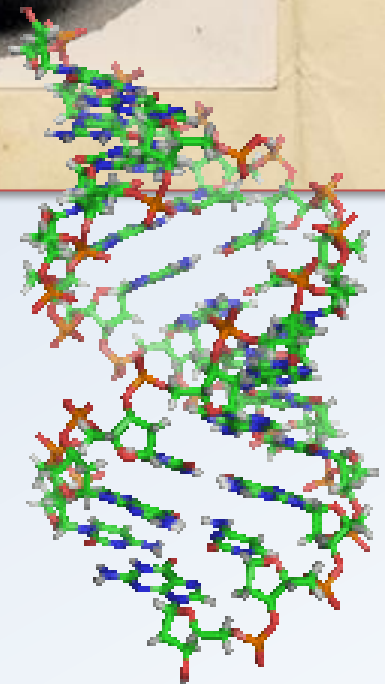
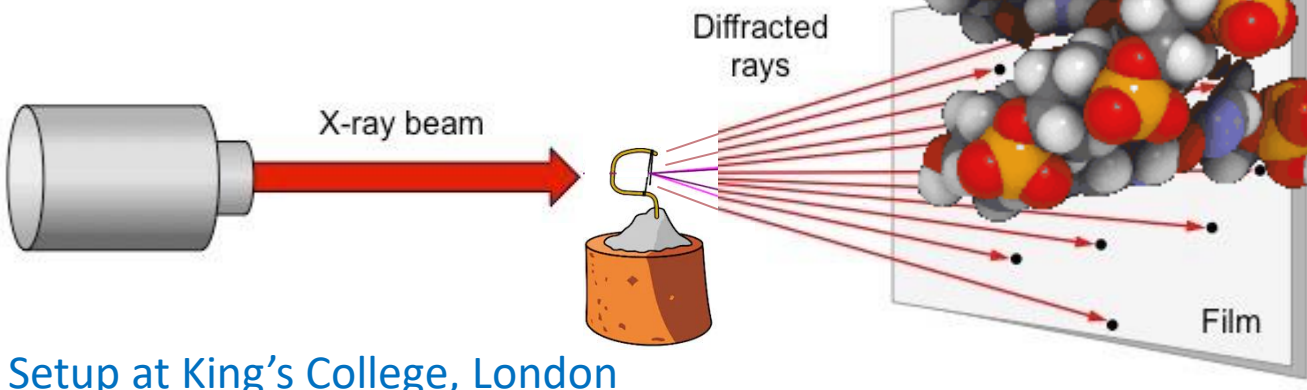
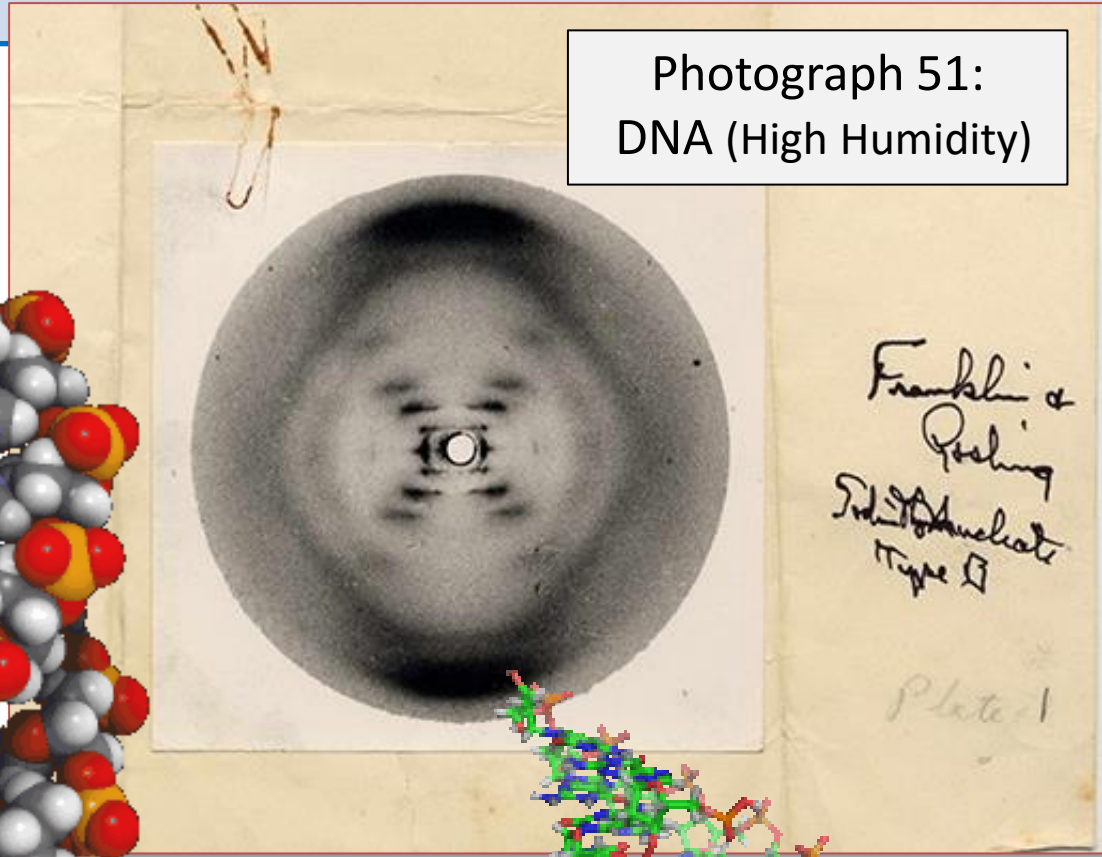
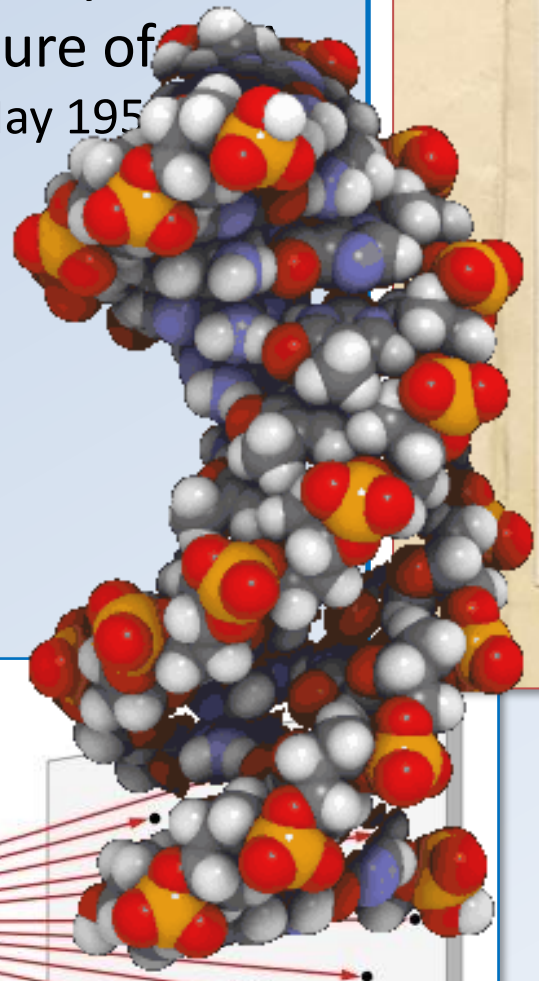


Setup at King's College, London

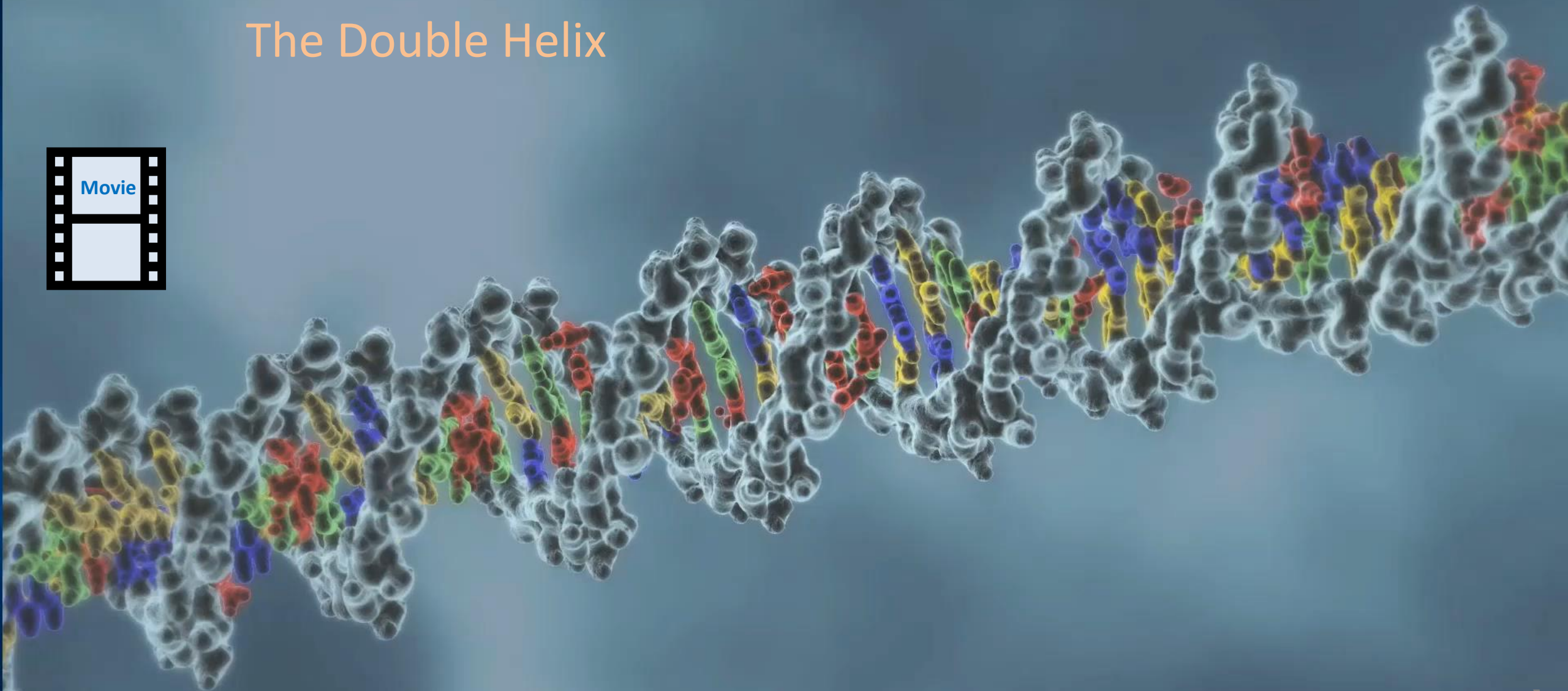
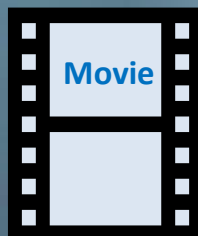


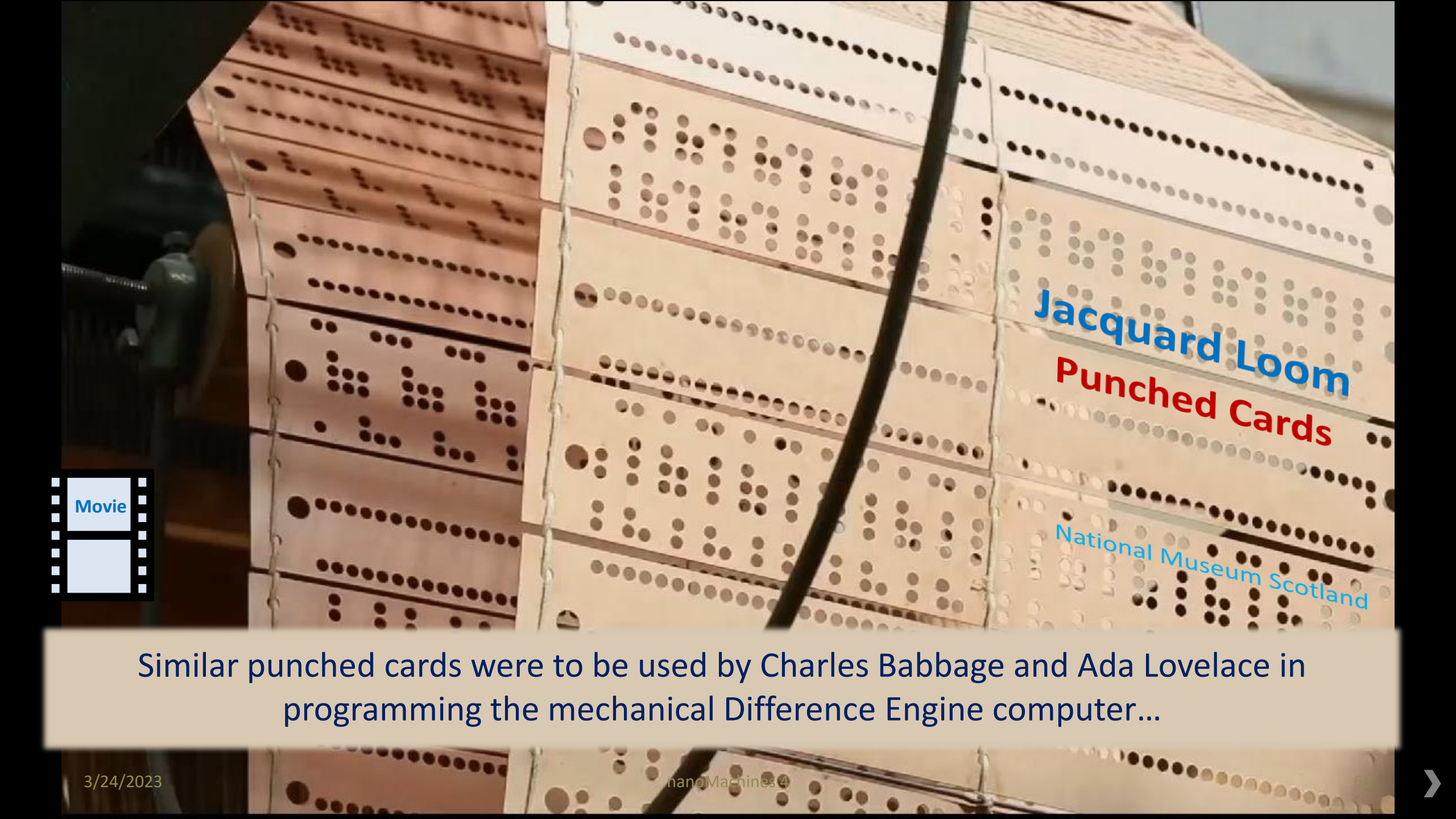


Key to the
Discovery of the
Structure of
May 195



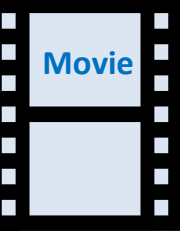
The Double Helix





Jacquard Loom
Punched Cards

National Museum Scotland



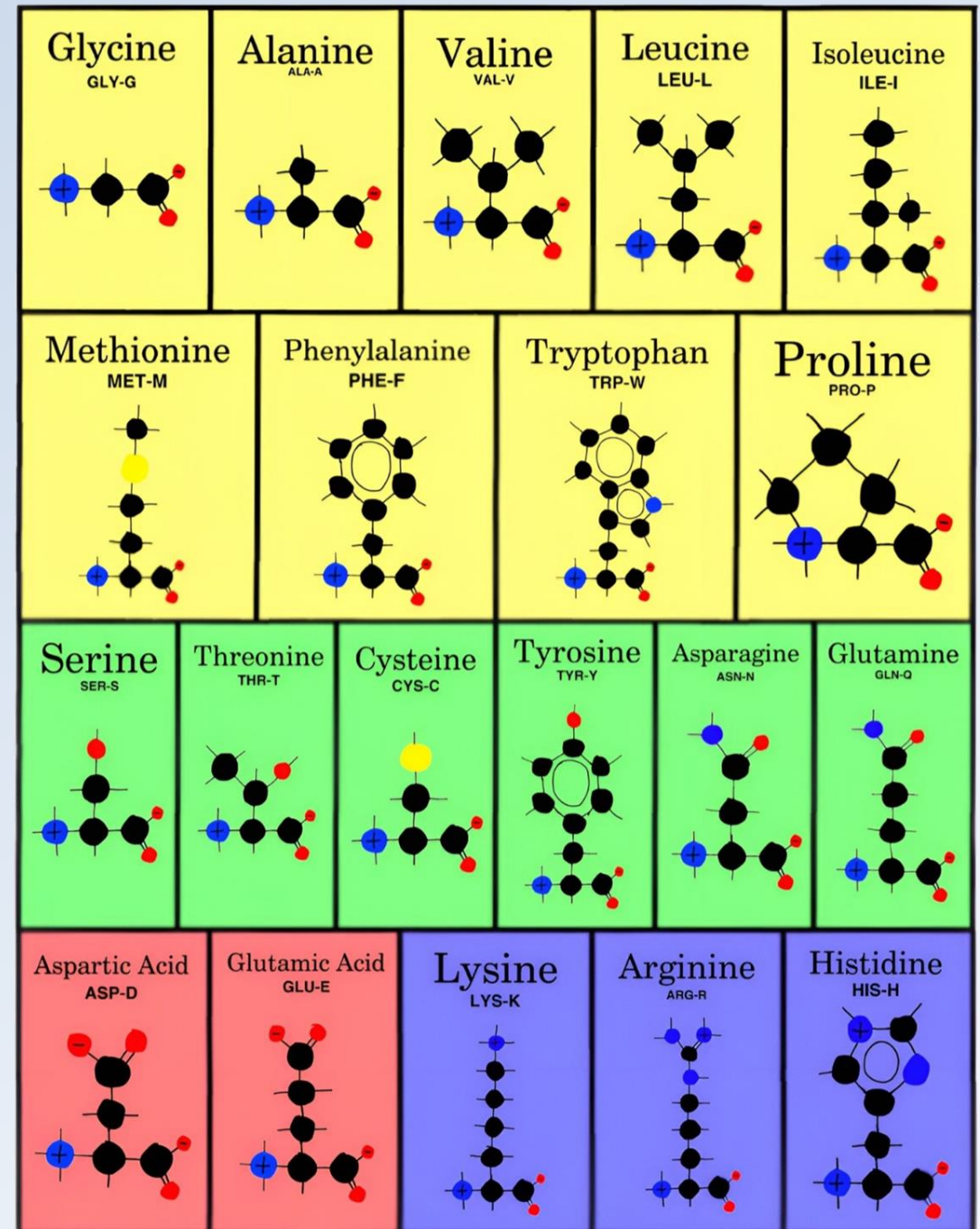
Similar punched cards were to be used by Charles Babbage and Ada Lovelace in programming the mechanical Difference Engine computer...



The 20 Major α -Amino Acids

G	Glycine	Gly	P	Proline	Pro
A	Alanine	Ala	V	Valine	Val
L	Leucine	Leu	I	Isoleucine	Ile
M	Methionine	Met	C	Cysteine	Cys
F	Phenylalanine	Phe	Y	Tyrosine	Tyr
W	Tryptophan	Trp	H	Histidine	His
K	Lysine	Lys	R	Arginine	Arg
Q	Glutamine	Gln	N	Asparagine	Asn
E	Glutamic Acid	Glu	D	Aspartic Acid	Asp
S	Serine	Ser	T	Threonine	Thr

 Non-polar	 Acidic
 Polar	 Basic



The “Universal” Amino Acid Codon Code

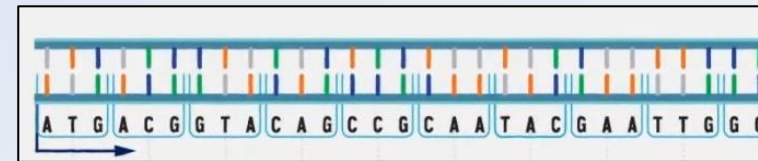
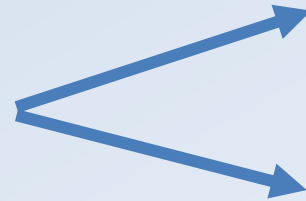
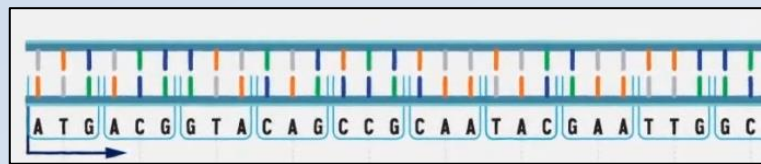
Some Amino Acids coded in 2 to 6 different ways

1st base	2nd base								3rd base
	U = Uracil		C = Cytosine		A = Adenine		G = Guanine		
U	UUU	(Phe/F) Phenylalanine	UCU	(Ser/S) Serine	UAU	(Tyr/Y) Tyrosine	UGU	(Cys/C) Cysteine	U
	UUC	Phenylalanine	UCC		UAC		UGC		C
	UUA	6 Start Code	UCA		UAA	Stop	UGA	Stop	A
	UUG		UCG		UAG		UGG	(Trp/W) Tryptophan	G
C	CUU	(Leu/L) Leucine	CCU	(Pro/P) Proline	CAU	(His/H) Histidine	CGU	(Arg/R) Arginine	U
	CUC		CCC		CAC		CGC		C
	CUA		CCA		CAA	CGA	A		
	CUG		CCG		CAG	CGG	G		
A	AUU	(Ile/I) Isoleucine	ACU	(Thr/T) Threonine	AAU	(Asn/N) Asparagine	AGU	(Ser/S) Serine	U
	AUC		ACC		AAC		AGC		C
	AUA		ACA		AAA	AGA	A		
	AUG		ACG		AAG	AGG	G		
G	GUU	(Val/V) Valine	GCU	(Ala/A) Alanine	GAU	(Asp/D) Aspartic acid	GGU	(Gly/G) Glycine	U
	GUC		GCC		GAC		GGC		C
	GUA		GCA		GAA	GGA	A		
	GUG		GCG		GAG	GGG	G		



Questions?

① DNA Replication



2
Identical
Copies

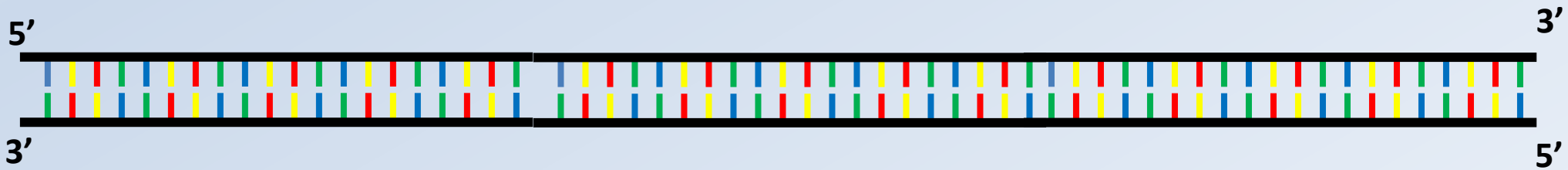


① DNA Replication



Should be easy...

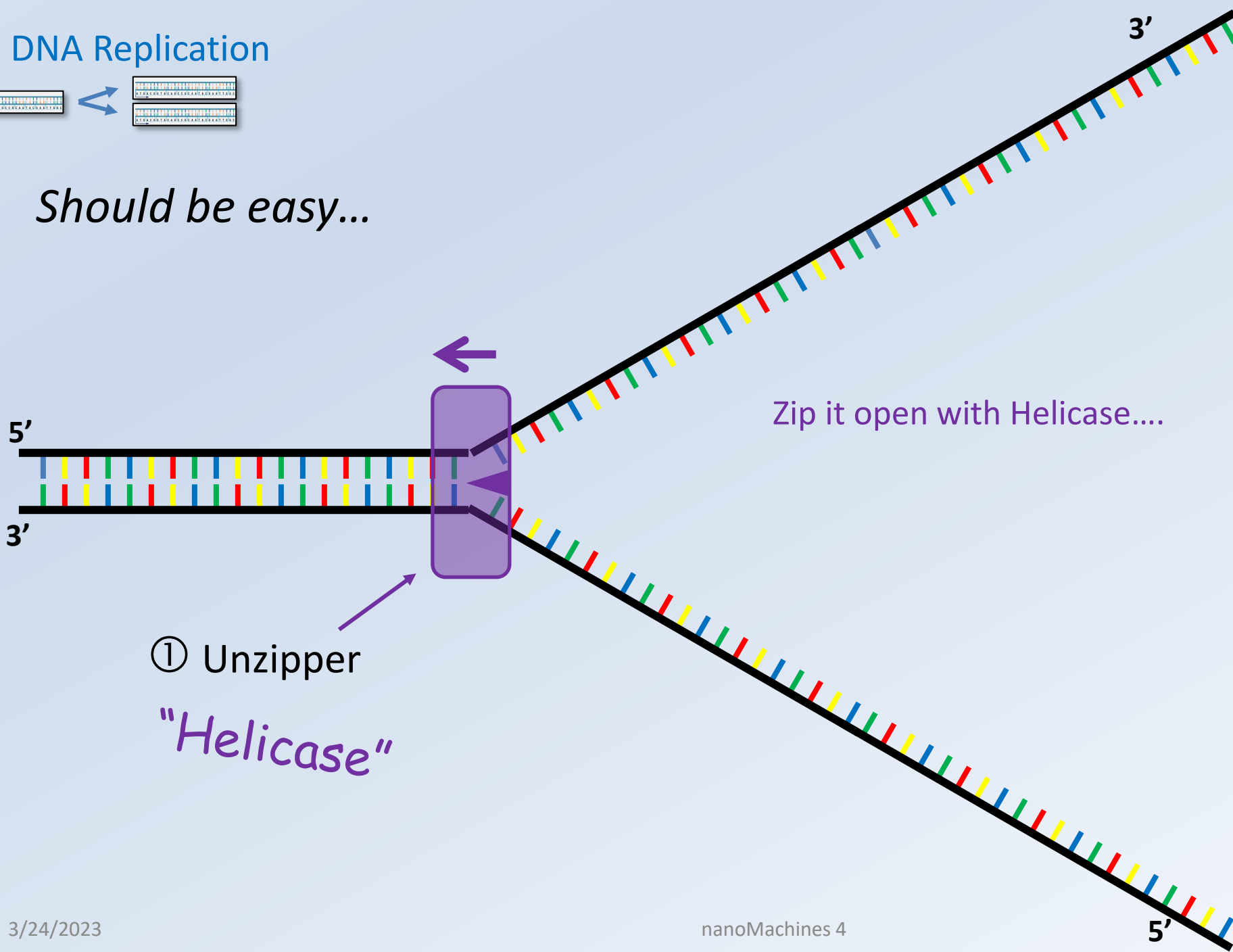
Start with double stranded DNA



① DNA Replication



Should be easy...



① Unzipper
"Helicase"

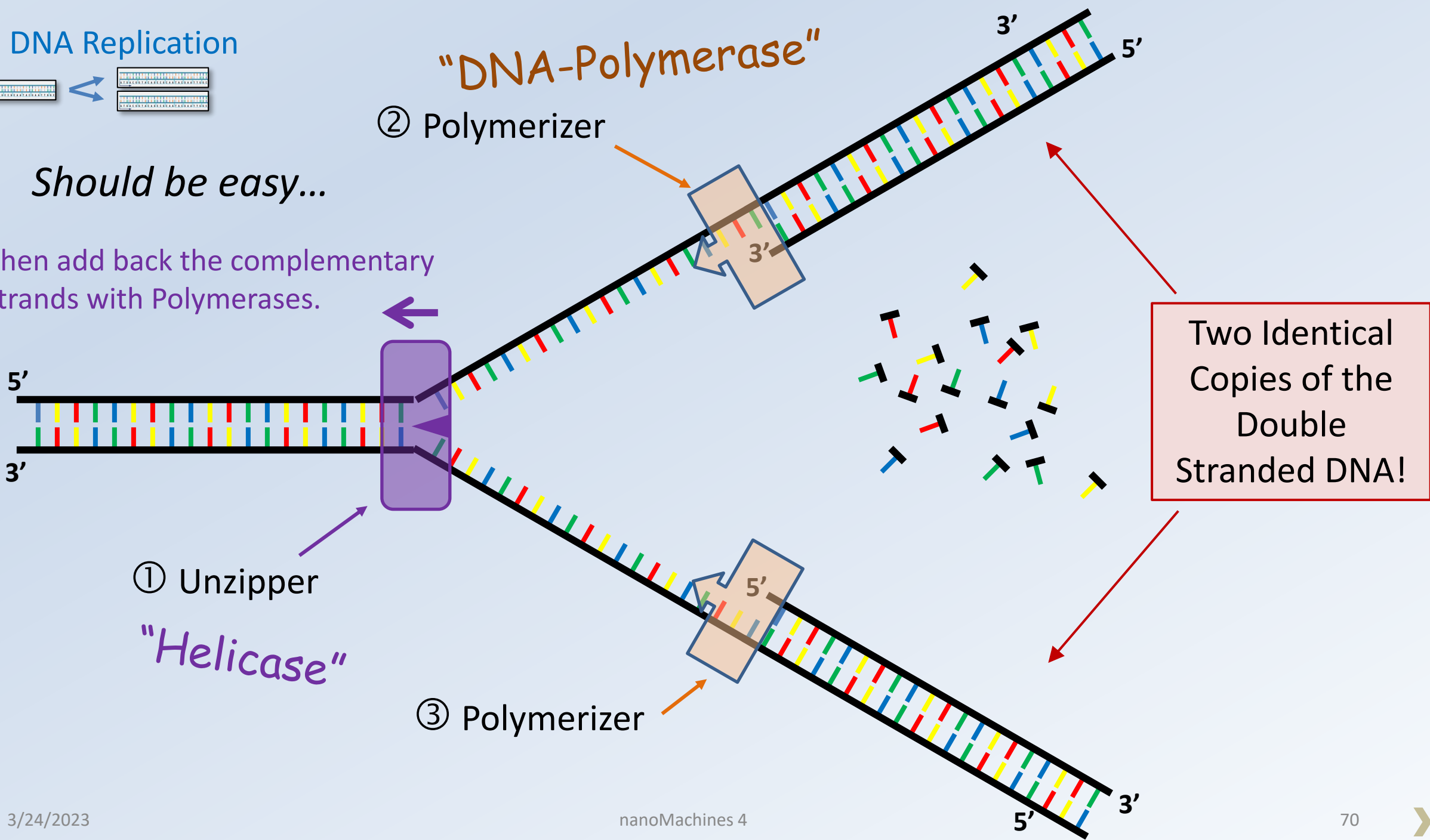


① DNA Replication



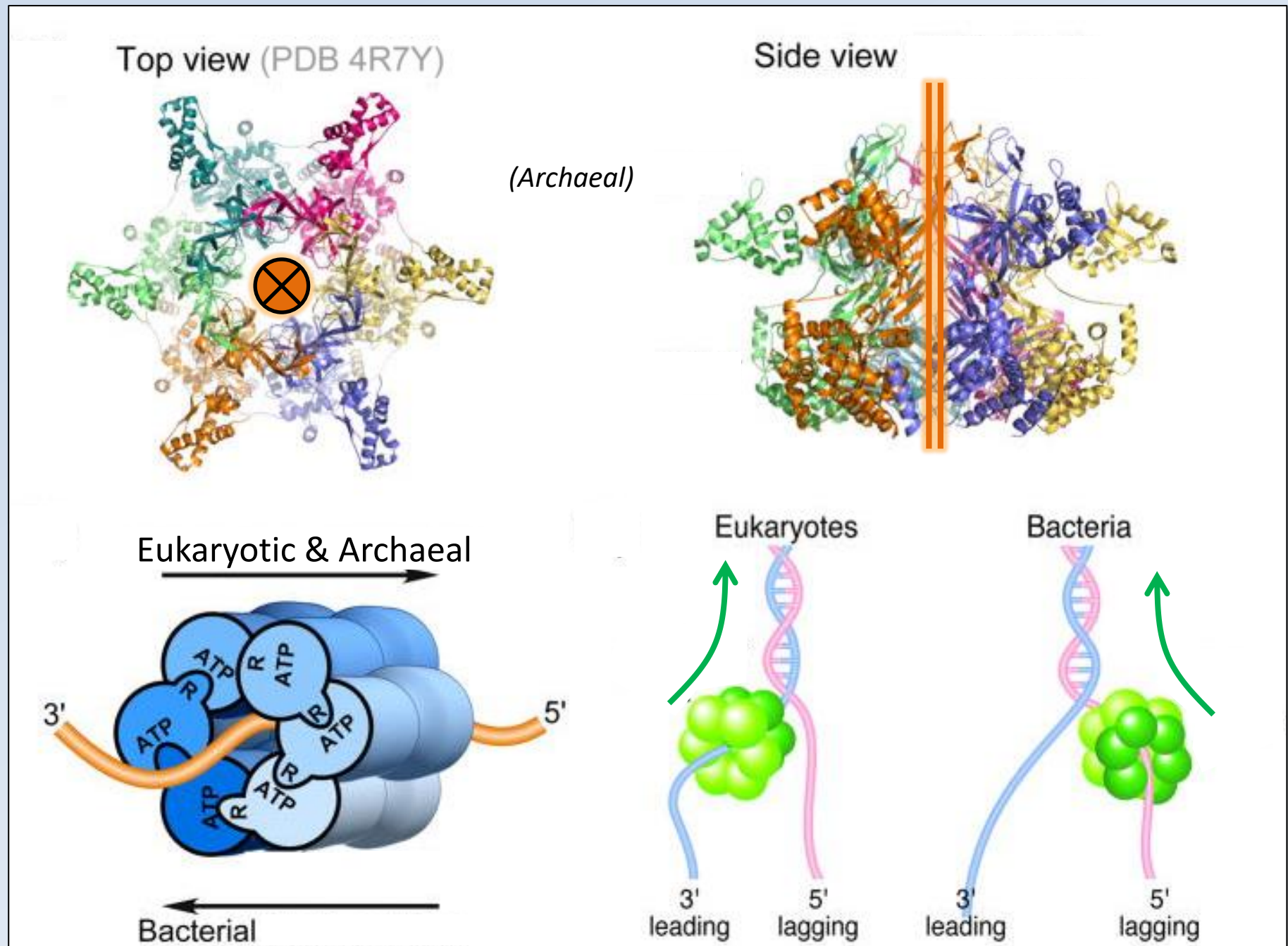
Should be easy...

Then add back the complementary strands with Polymerases.

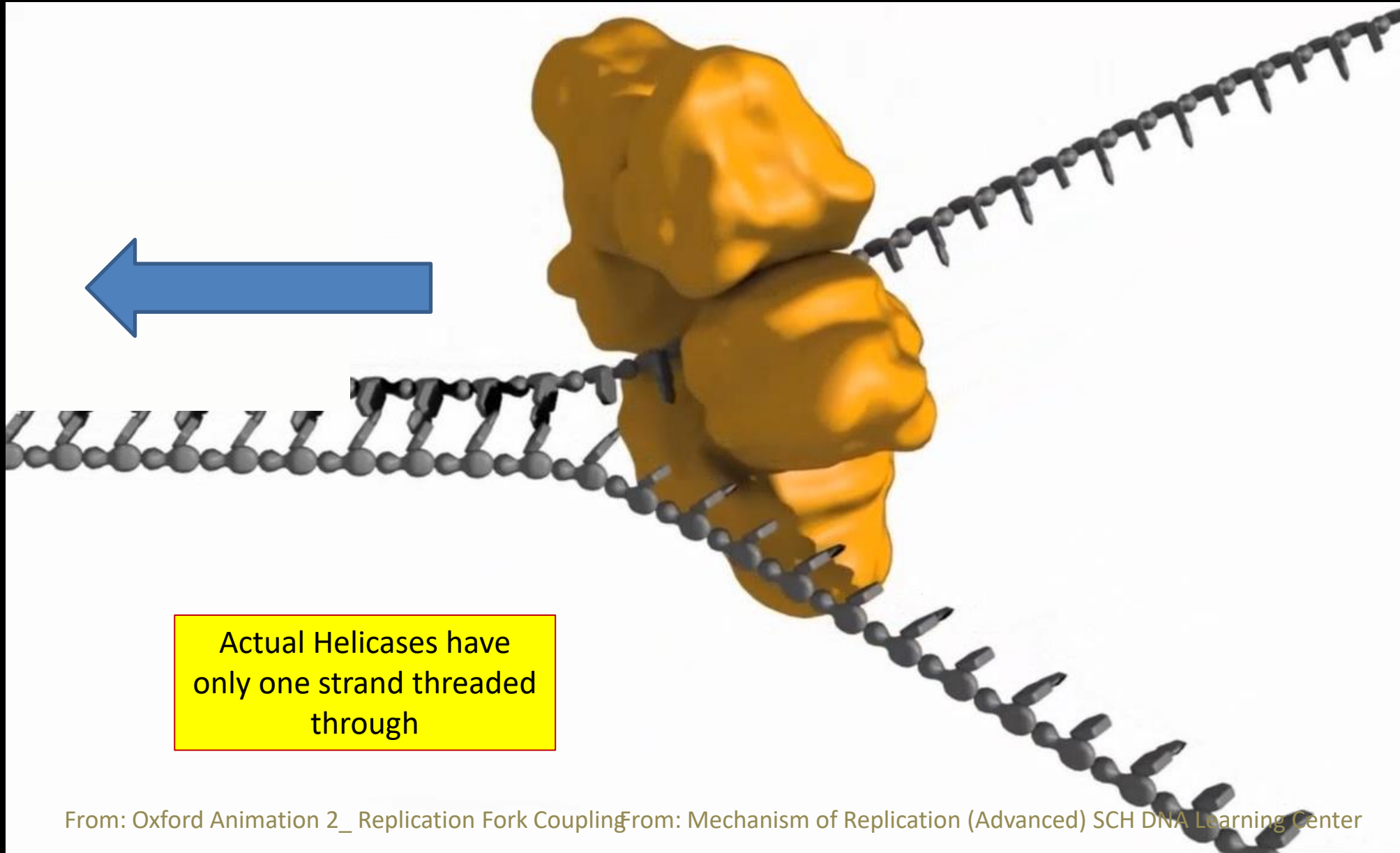


Unzipper
Helicases are completely different in Bacteria vs. Eukaryotes (and Archaea)

- Hexamers
- Burn ATP
- Surround only *one* DNA strand

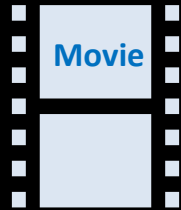


Helicase Animations

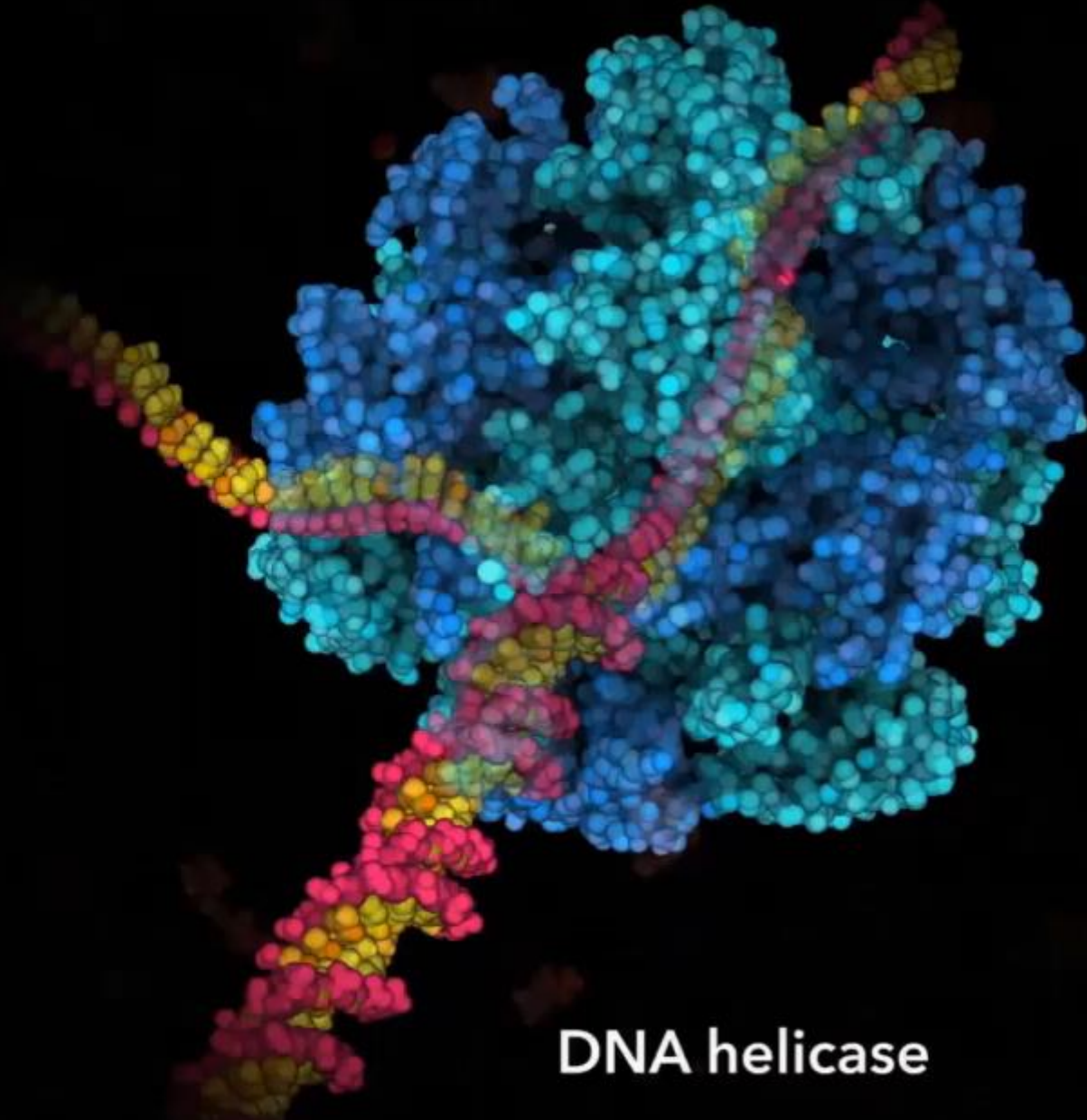


Actual Helicases have only one strand threaded through

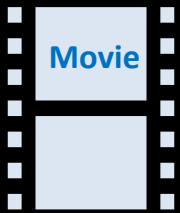
From: Oxford Animation 2_ Replication Fork Coupling From: Mechanism of Replication (Advanced) SCH DNA Learning Center



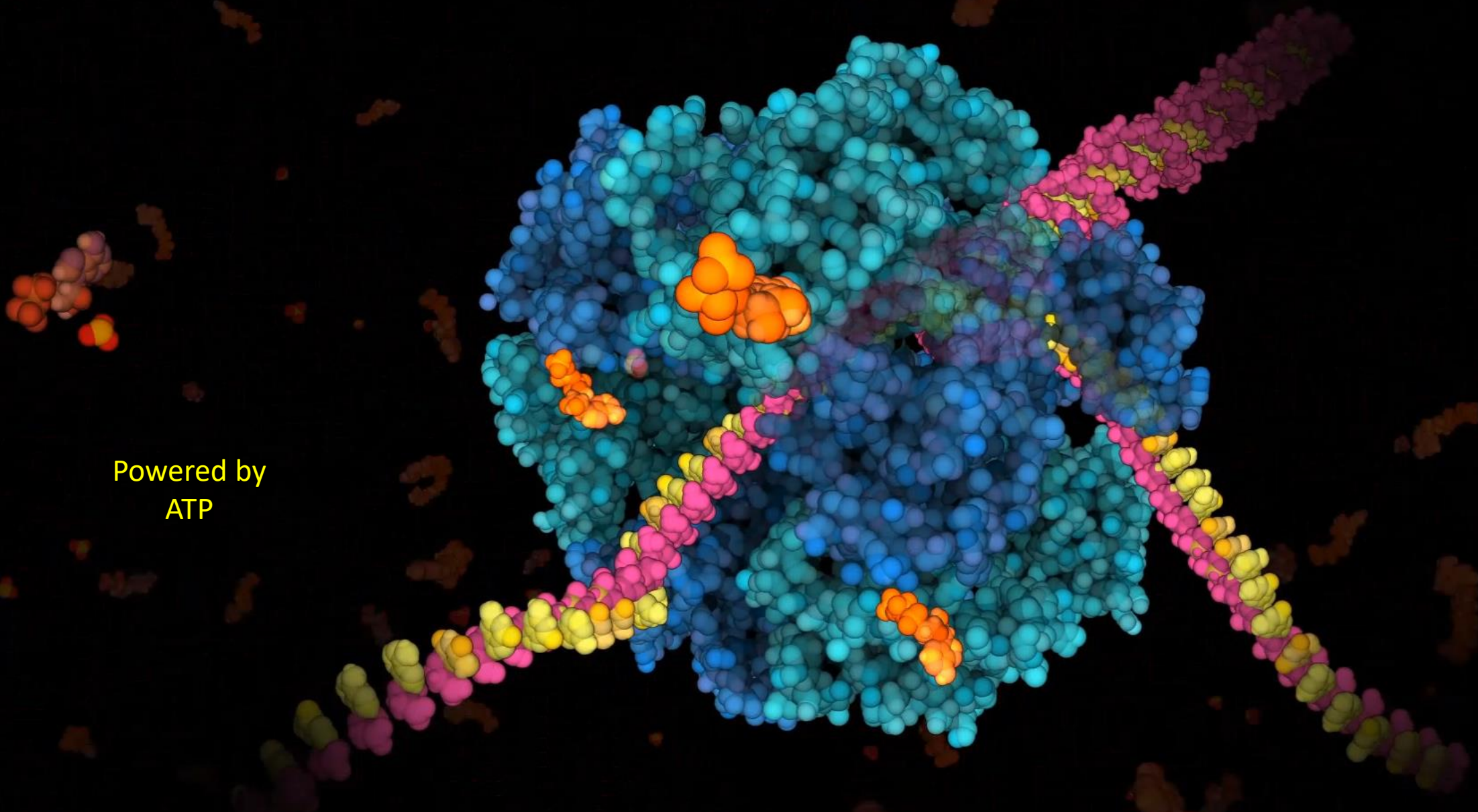
Helicase Animations



DNA helicase

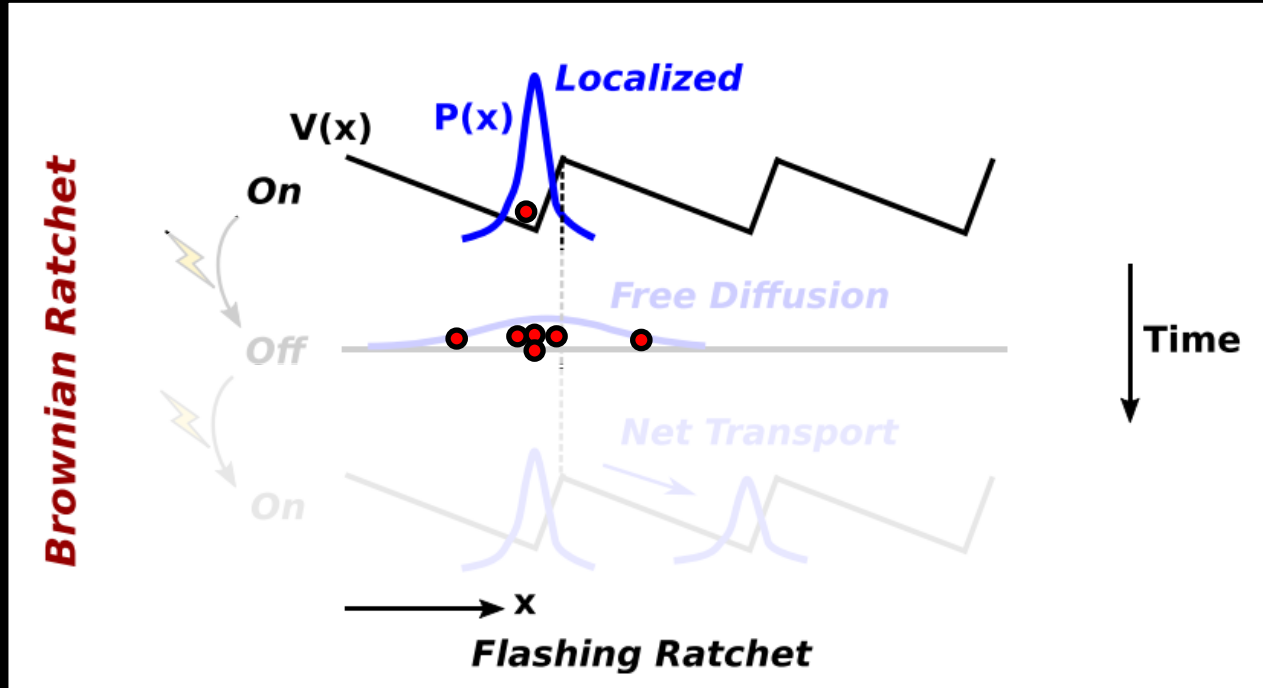
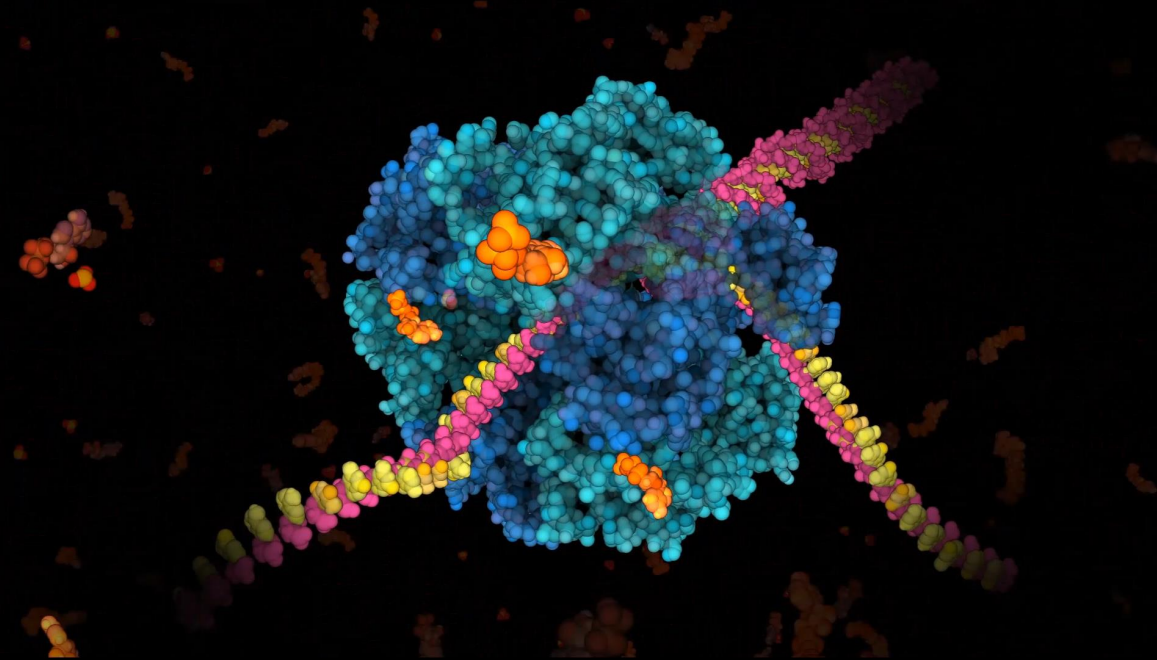


Powered by
ATP



Helicase steps are very small –
0.34 nm per base.

Recent work suggests that
Eukaryotic Helicases are likely
close to being....



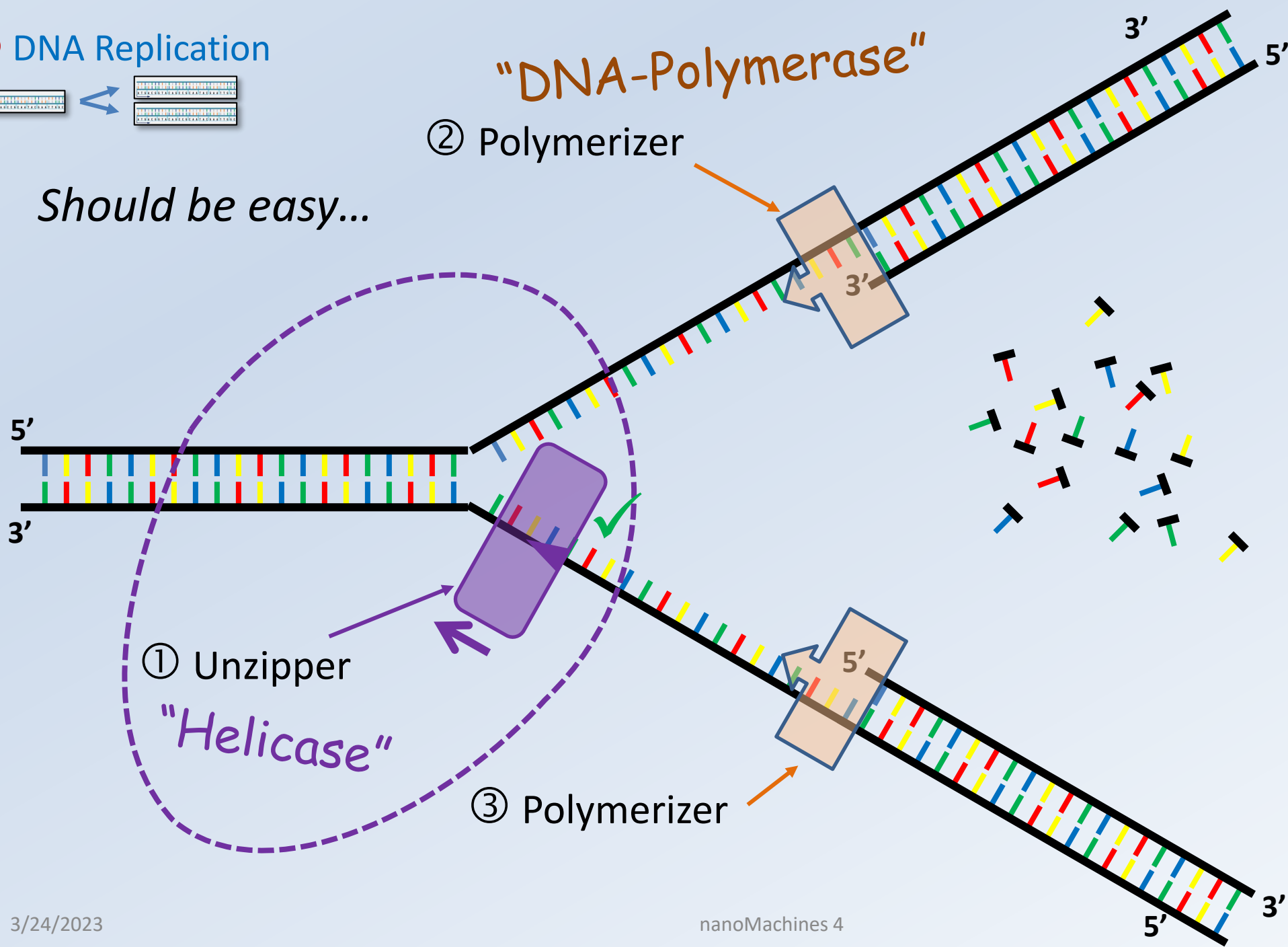
← “Brownian Ratchets”

Sometimes observed to
take a step backwards

① DNA Replication



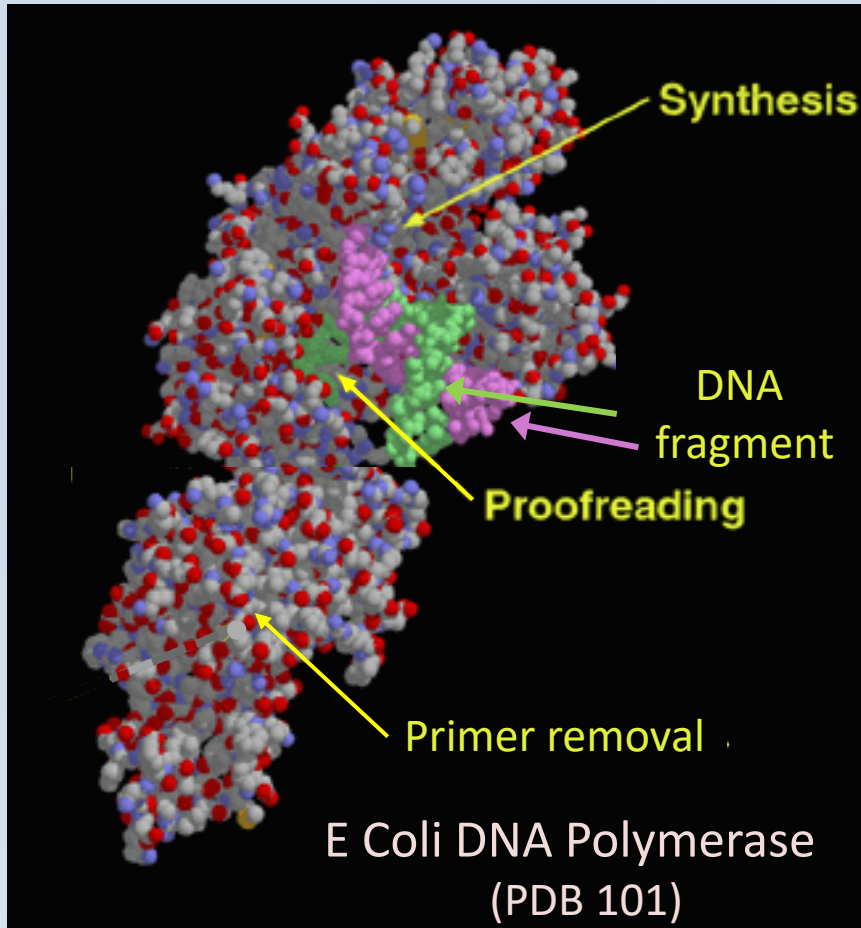
Should be easy...



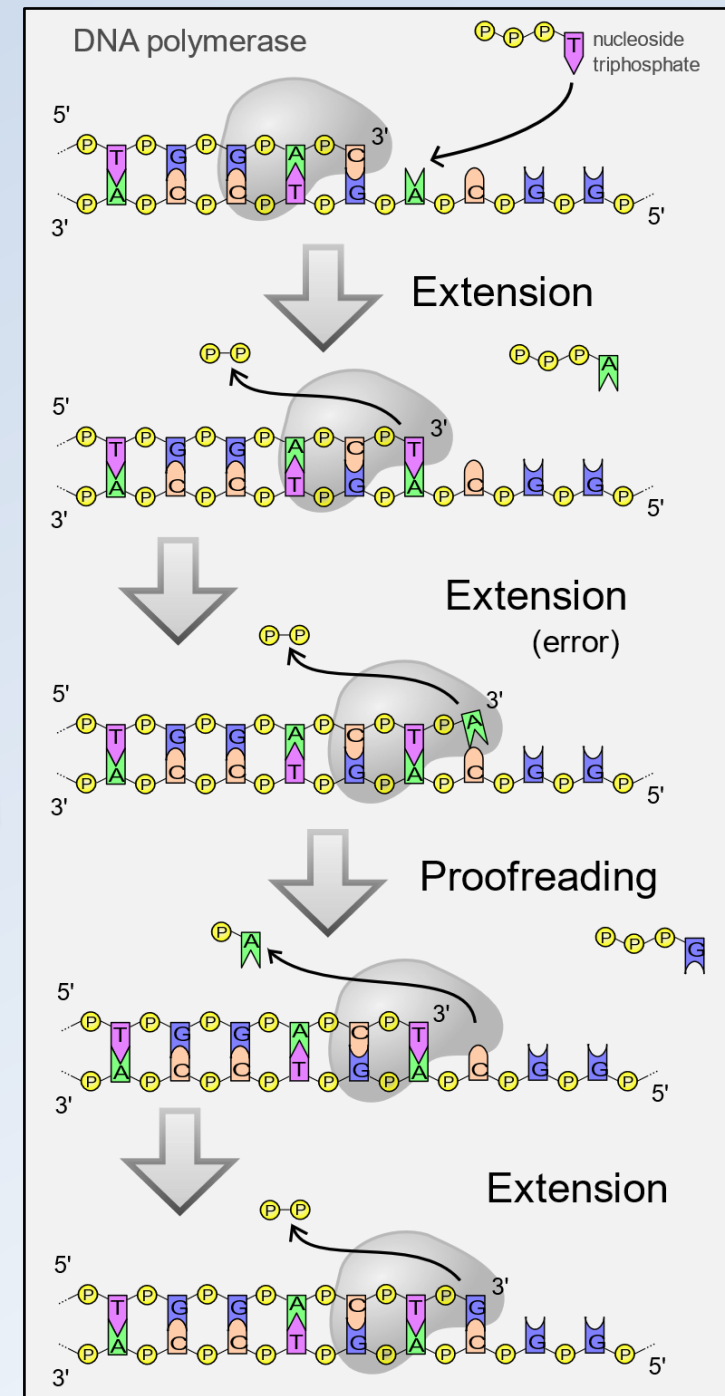
① DNA Replication



DNA-Polymerase



- Active sites are conserved across all life forms
- With **Proofreading**, Error Rates as low as 1 per Billion bases



DNA Polymerase 2010

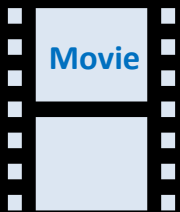
Drew Berry

webinar
DrewBerry

Single DNA
Strand Input

Double DNA
Strand Output

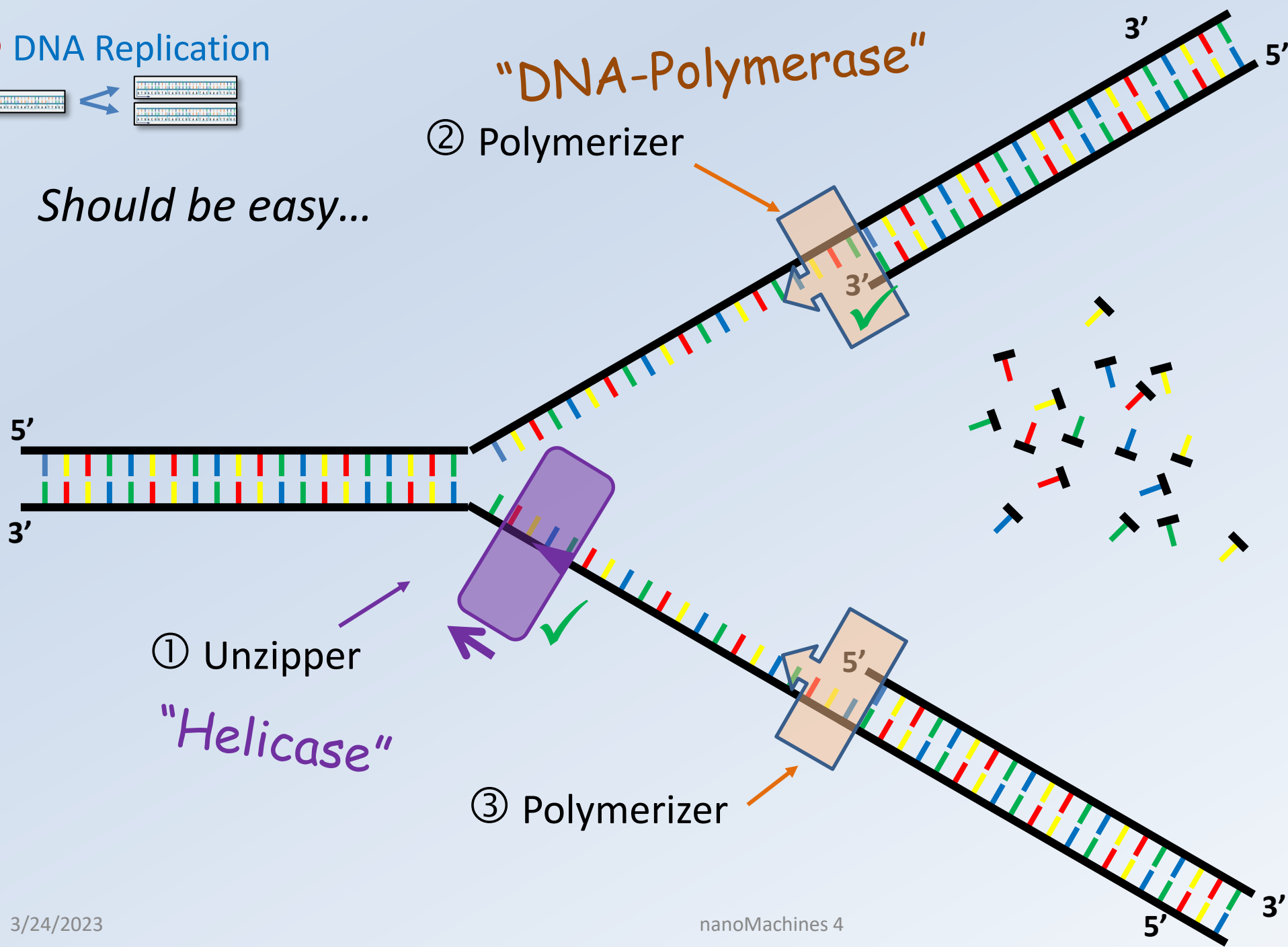
DNA Polymerase




① DNA Replication

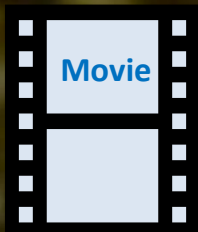


Should be easy...





Supercoiling is a
result of Helicase
unwinding of DNA

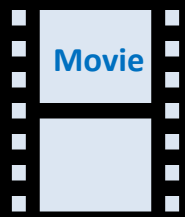


Untangler of Knots

Discovery Science (2022)



Untangler of Knots: Topoisomerases



Discovery Science (2022)

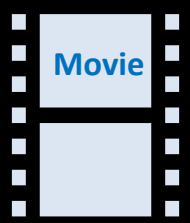
3/24/2023

nanoMachines 4

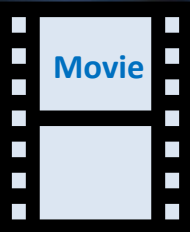
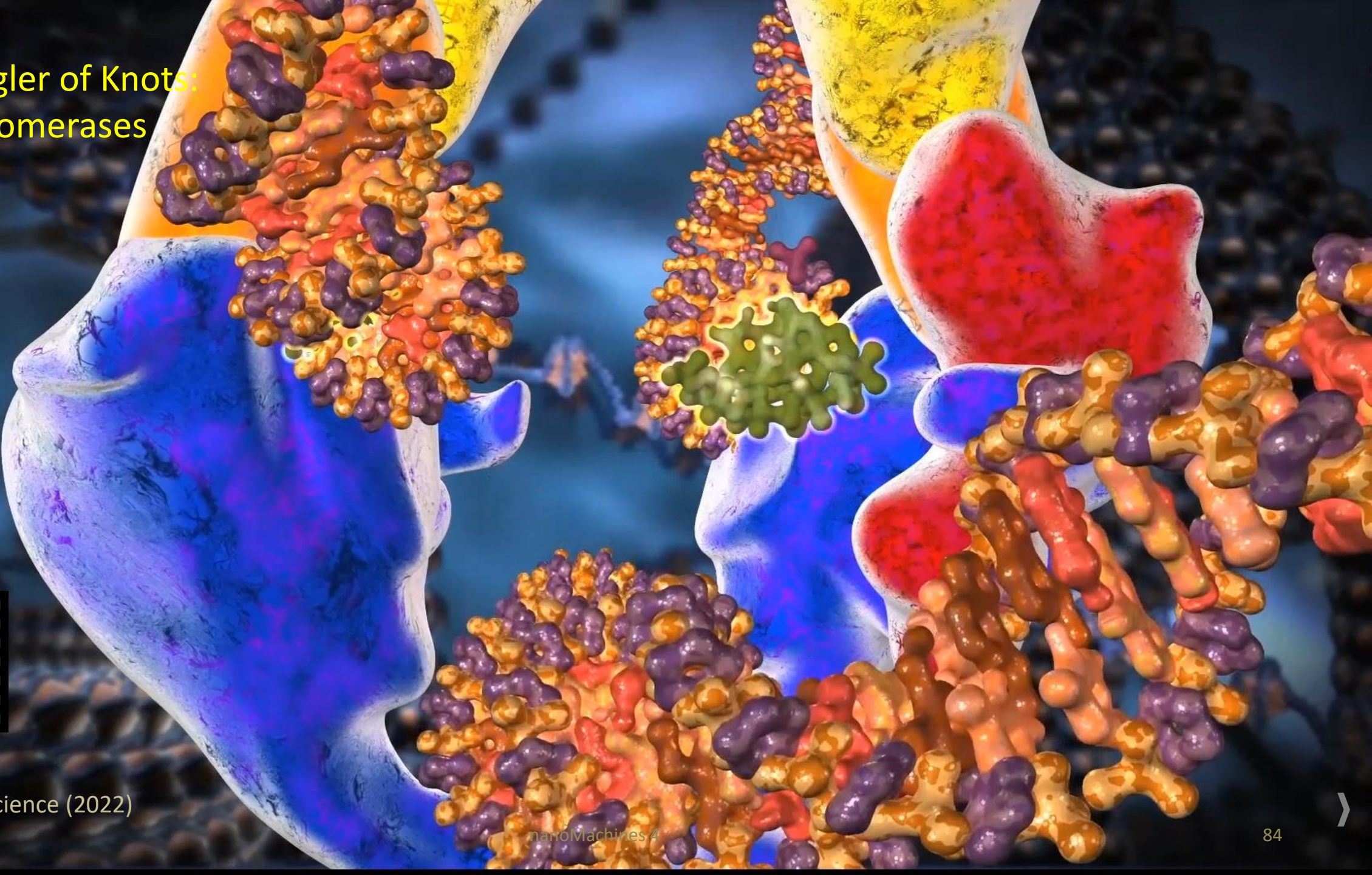


Untangler of Knots: Topoisomerases

ATP ADP



Untangler of Knots: Topoisomerases

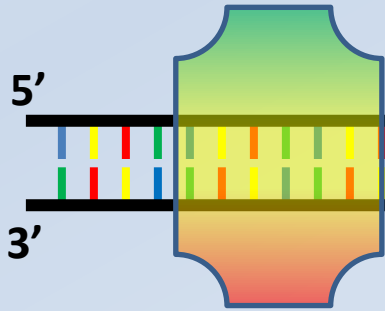


① DNA Replication



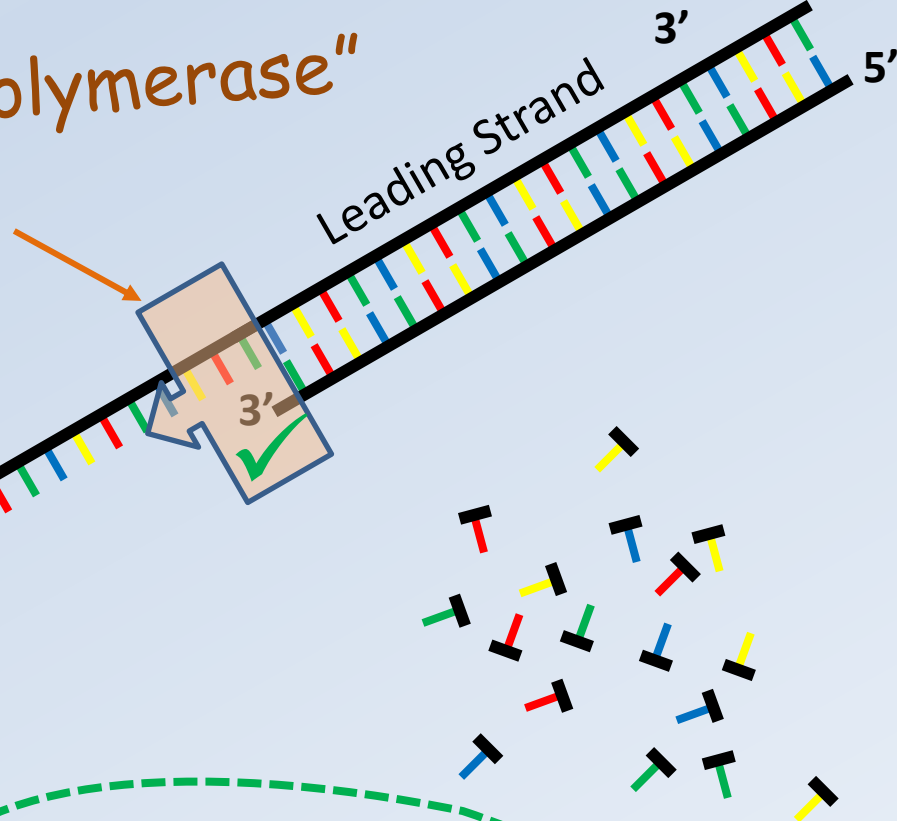
Maybe not so
~~Should be easy...~~

"Topo-
Isomerase"



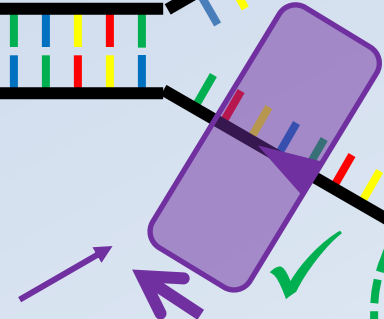
"DNA-Polymerase"

② Polymerizer

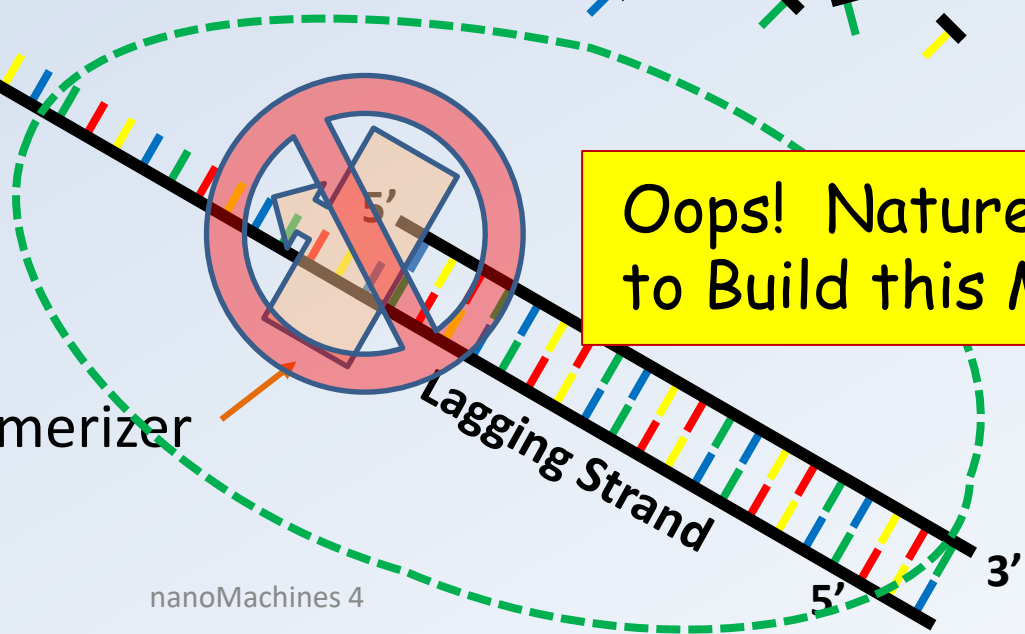


① Unzipper

"Helicase"



③ Polymerizer



Oops! Nature Forgot
to Build this Machine

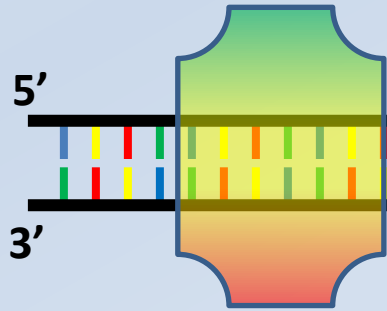


① DNA Replication



Maybe not so
~~Should be easy...~~

"Topo-
Isomerase"



① Unzipper

"Helicase"

"DNA-Polymerase"

② Polymerizer

③ Polymerizer

Leading Strand 3' 5'

Okazaki fragment

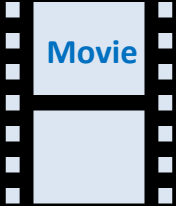
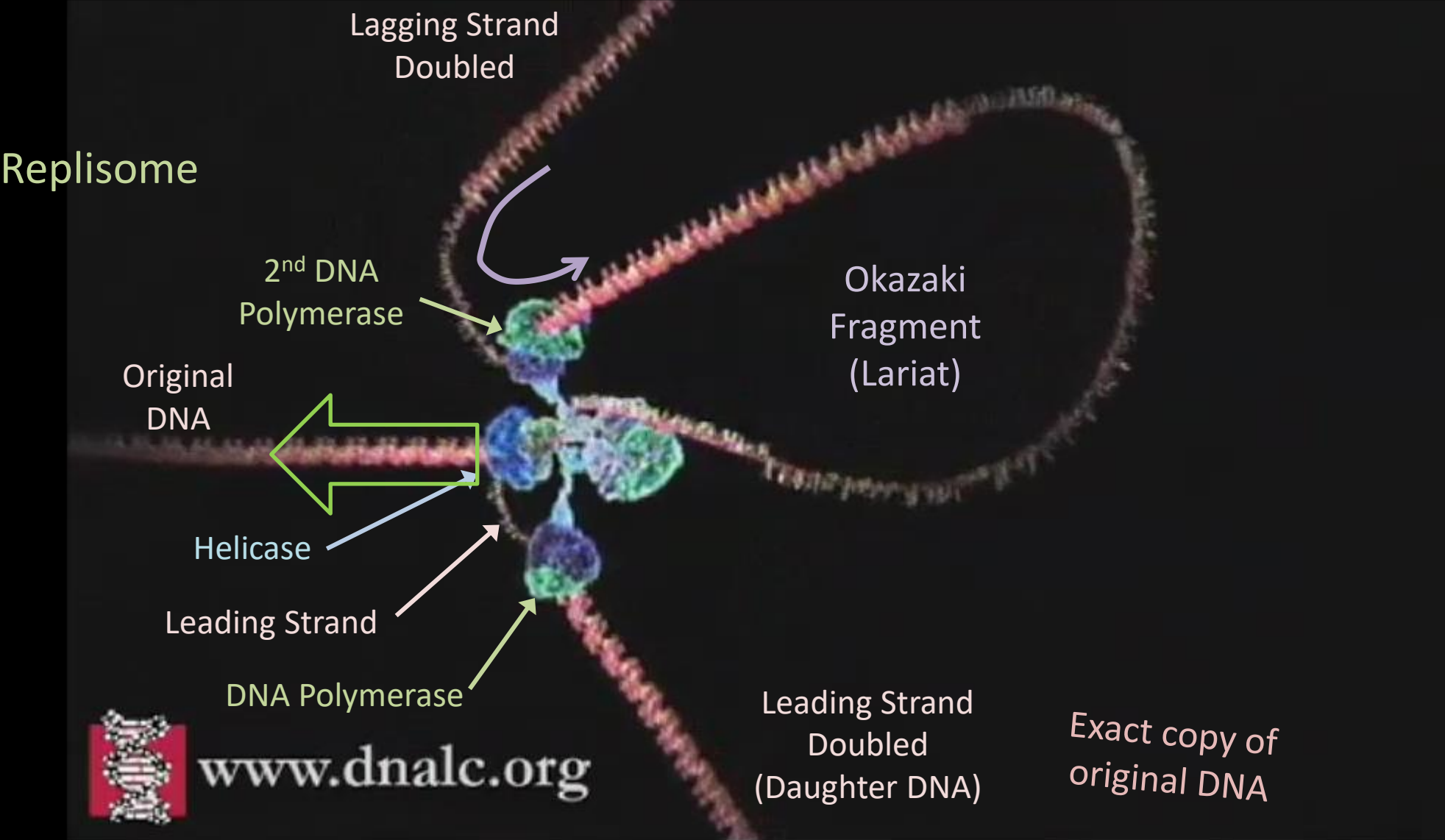
Lagging Strand 5' 3'

Rube Goldberg solution is to use regular 3'-5' DNA Polymerase backwards in short Okazaki fragments.



Mechanism of Replication

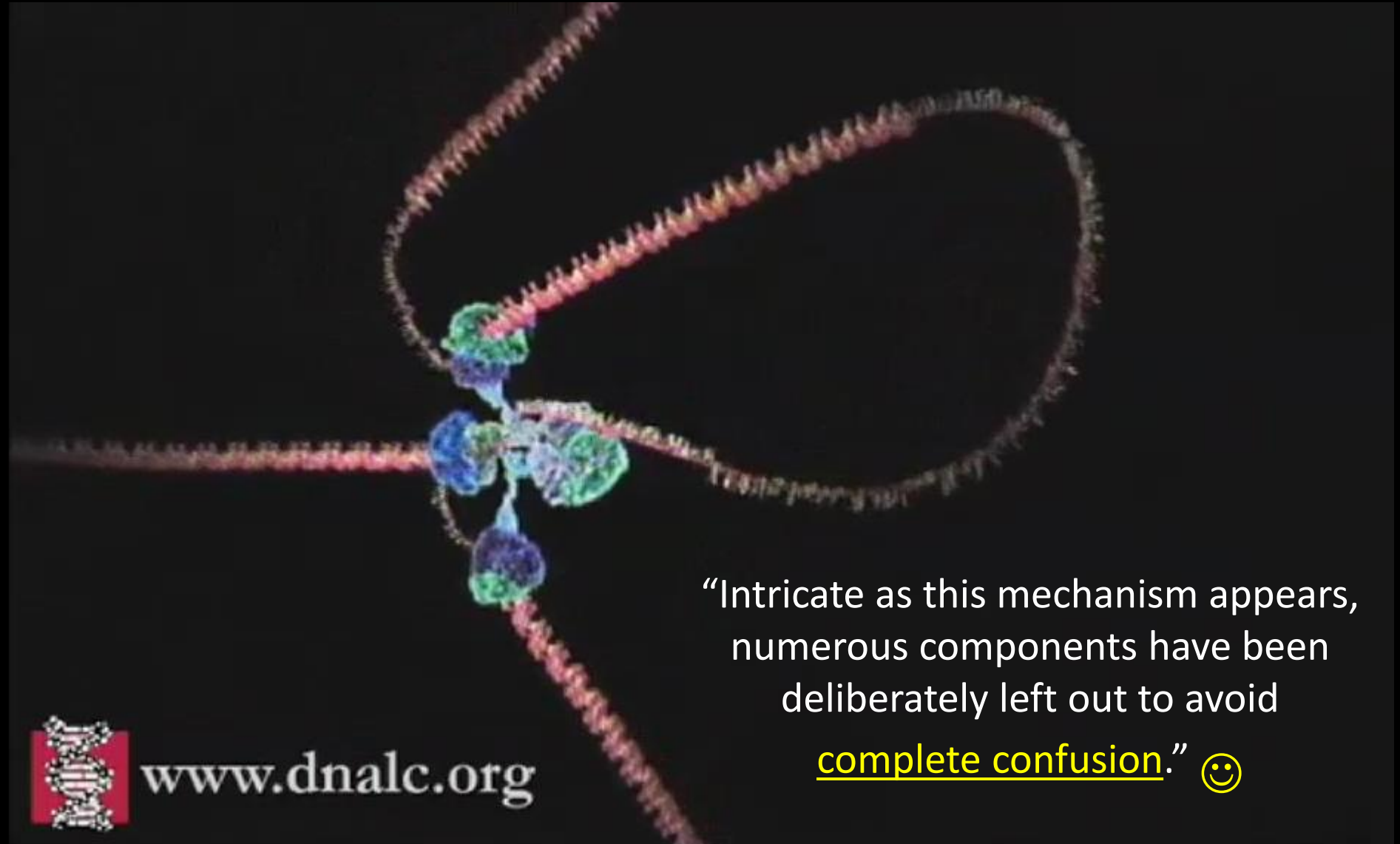
(DNA Learning Center)



Mechanism of Replication

(DNA Learning Center)

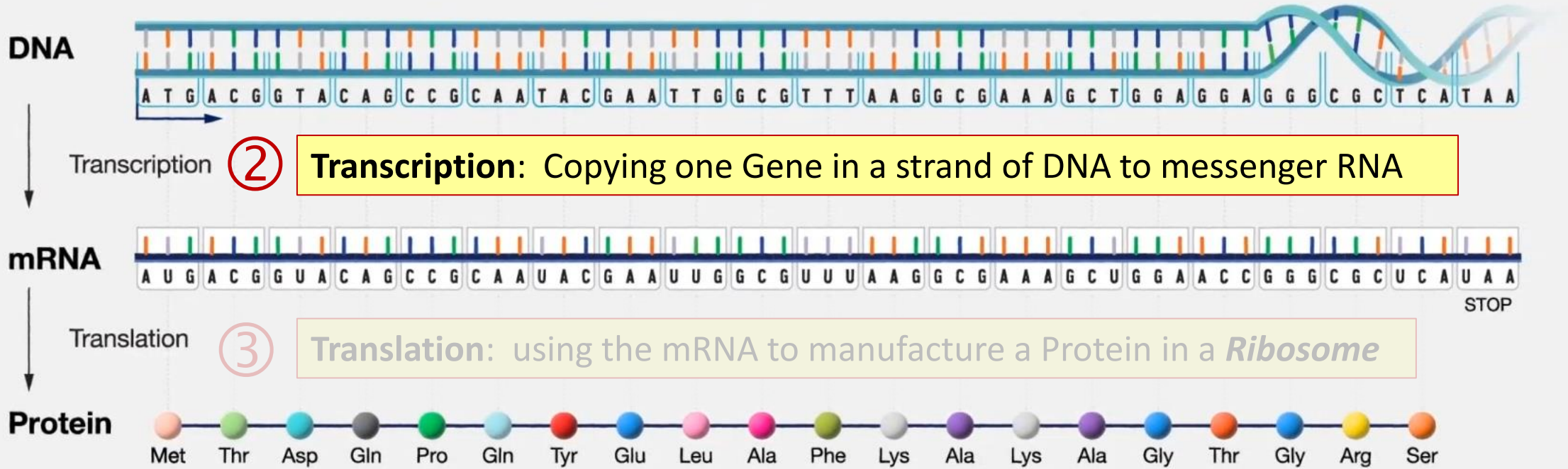
Replisome



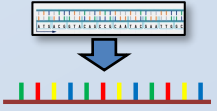
Questions?

Central Dogma

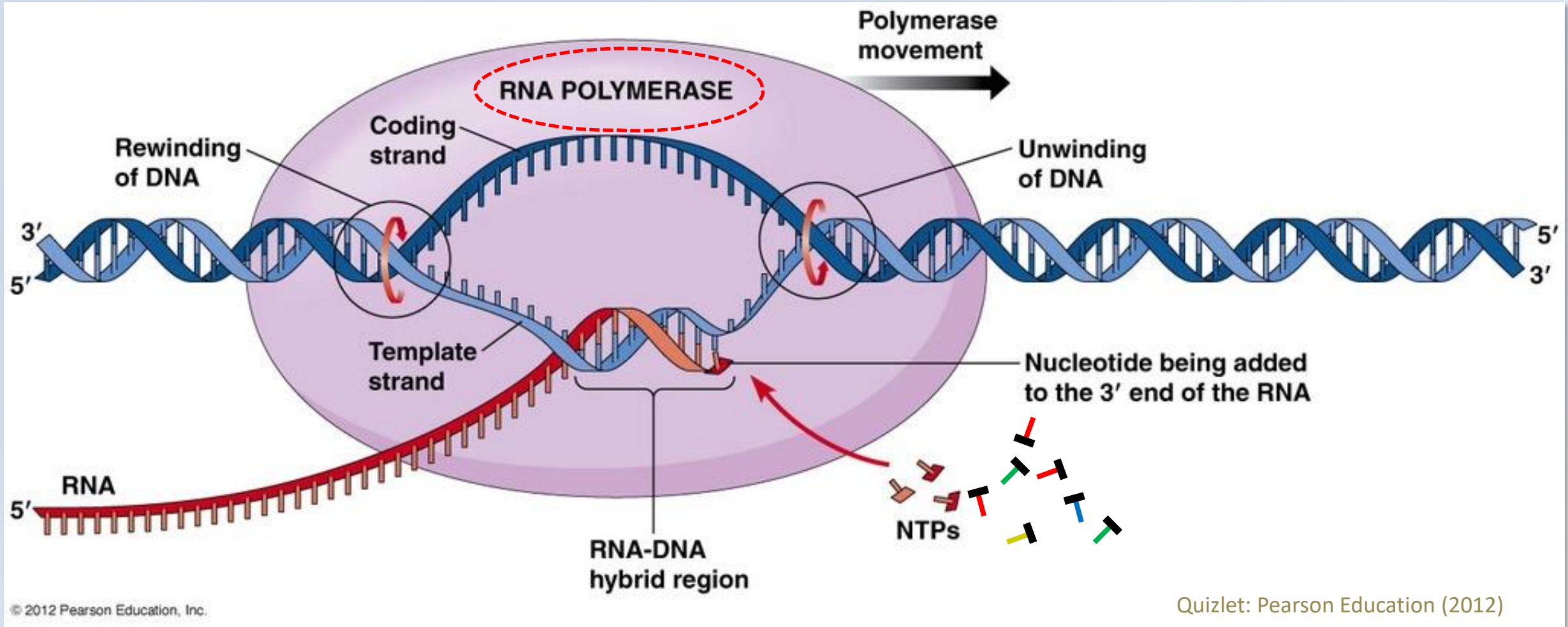
① **Replication:** Duplicating DNA when cells divide.



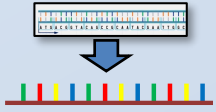
② RNA Transcription



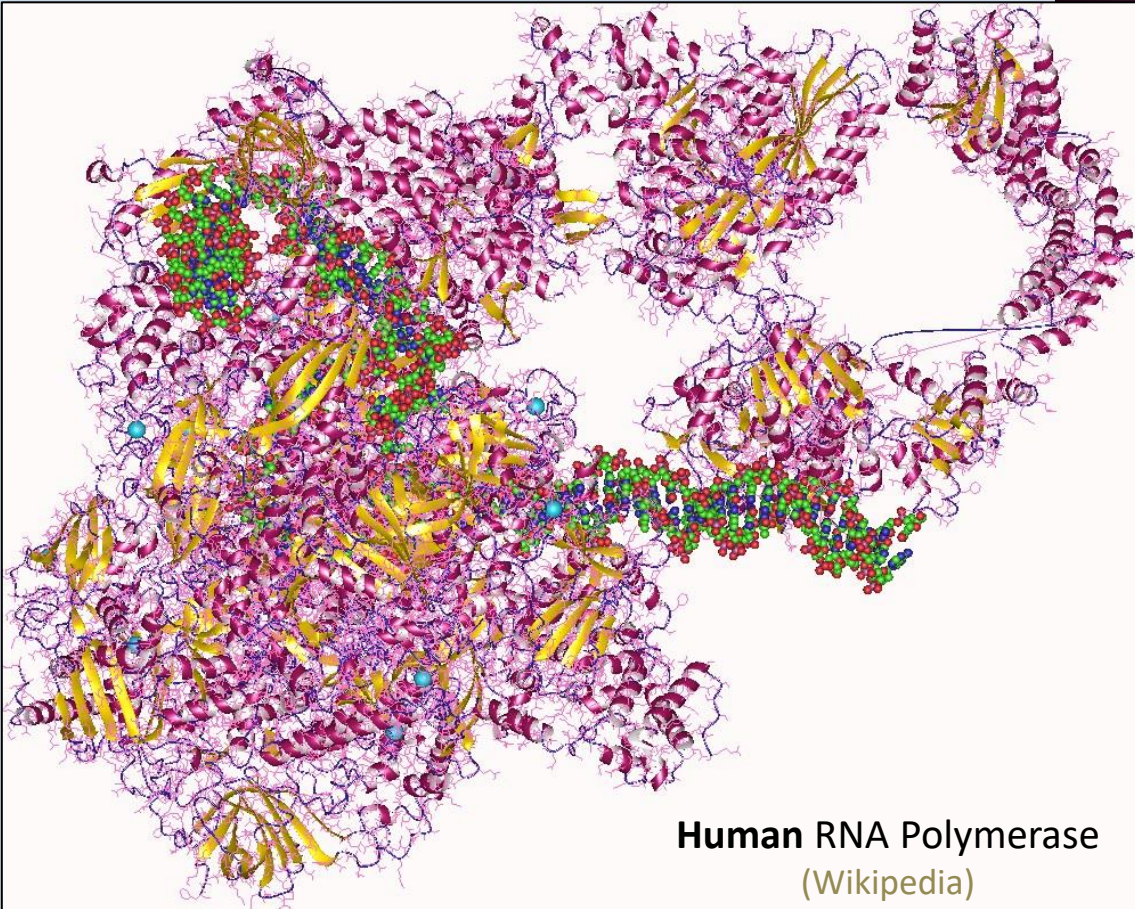
Building Messenger RNA (mRNA)



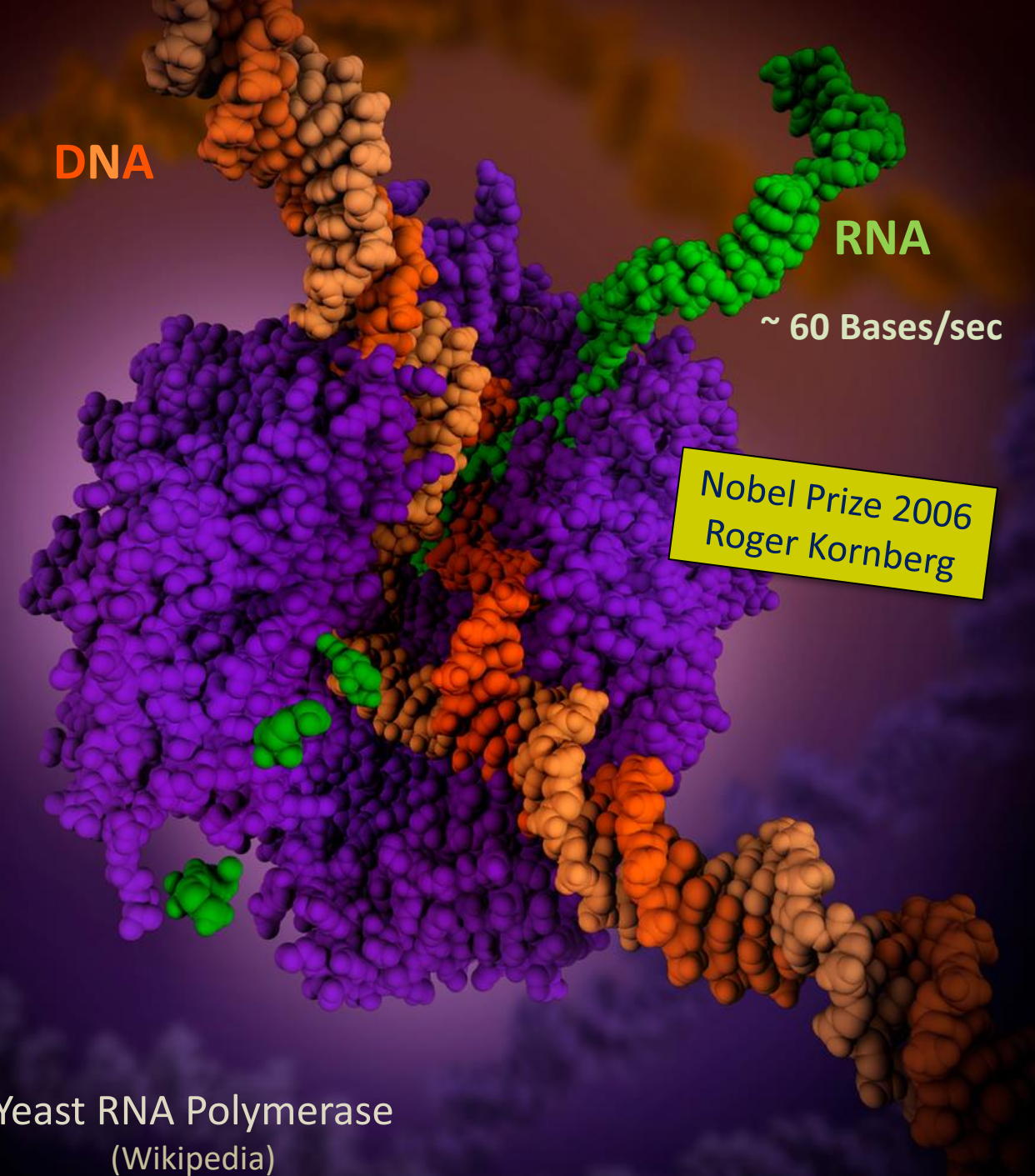
② RNA Transcription



RNA Polymerase



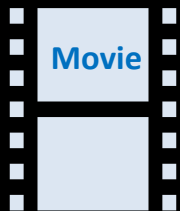
3/24/2023



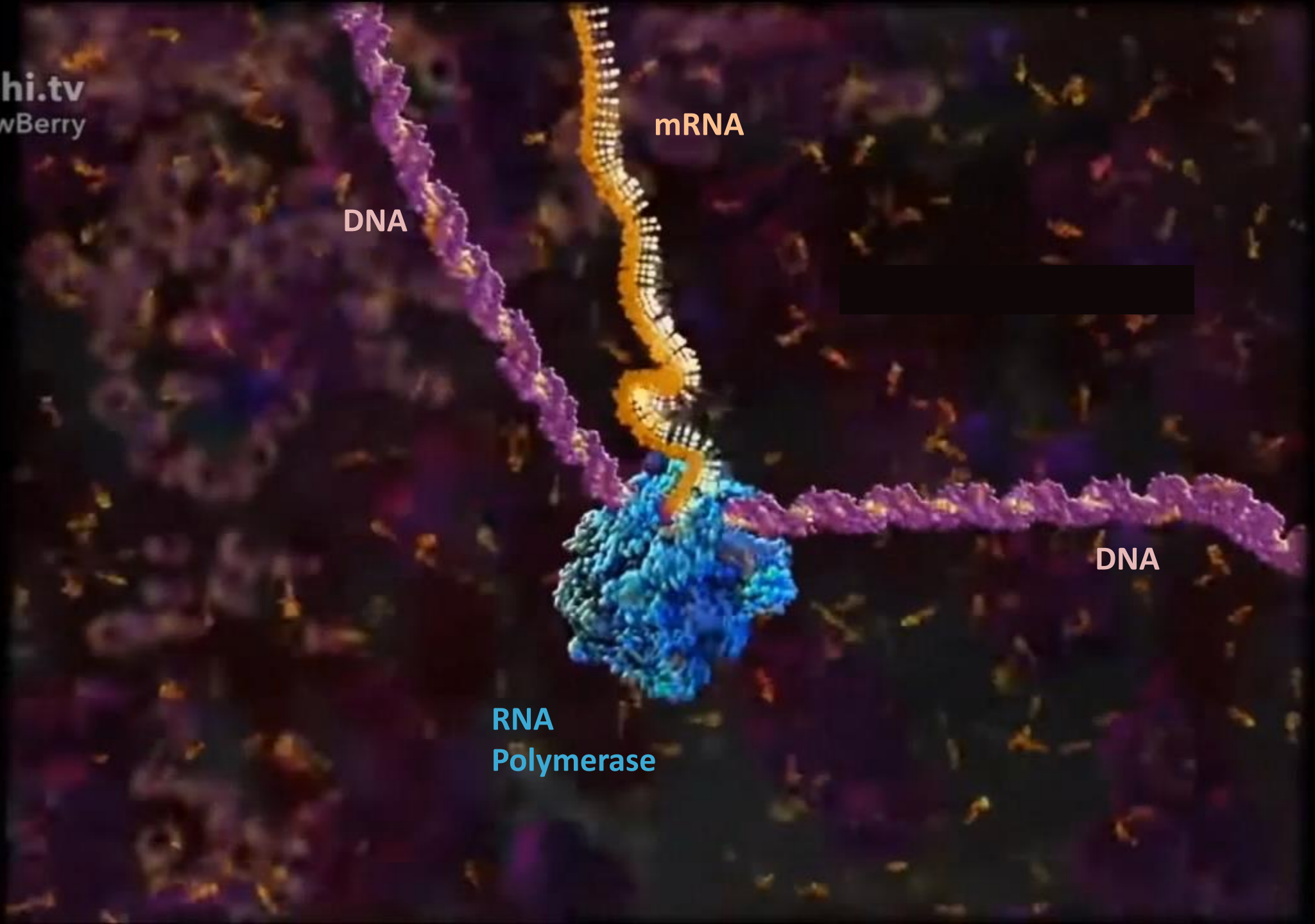
wehi.tv
DrewBerry

Gene
Transcription
RNA Polymerase

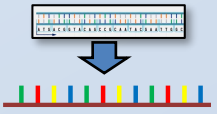
from:
DNA Animation
(2002-2014)
Drew Berry



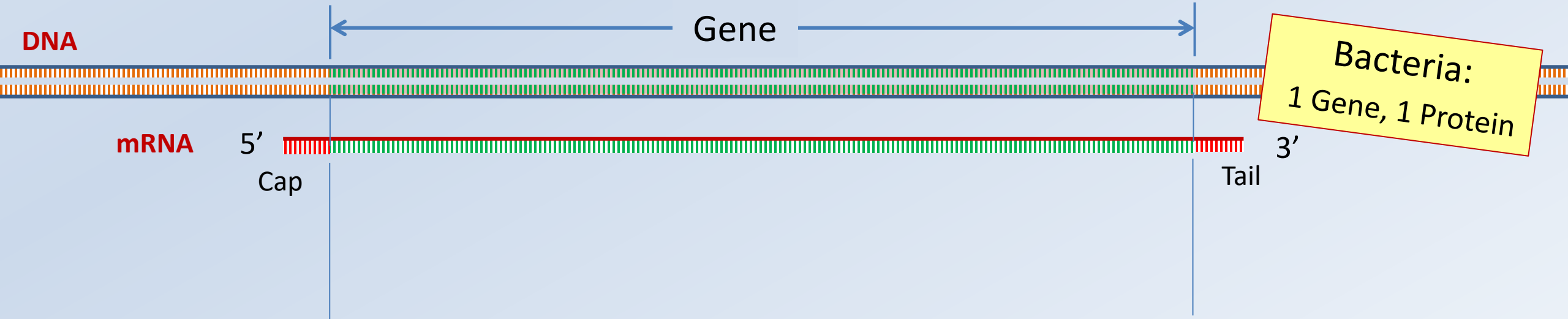
3/24/2023



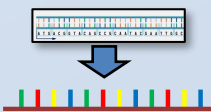
② RNA Transcription



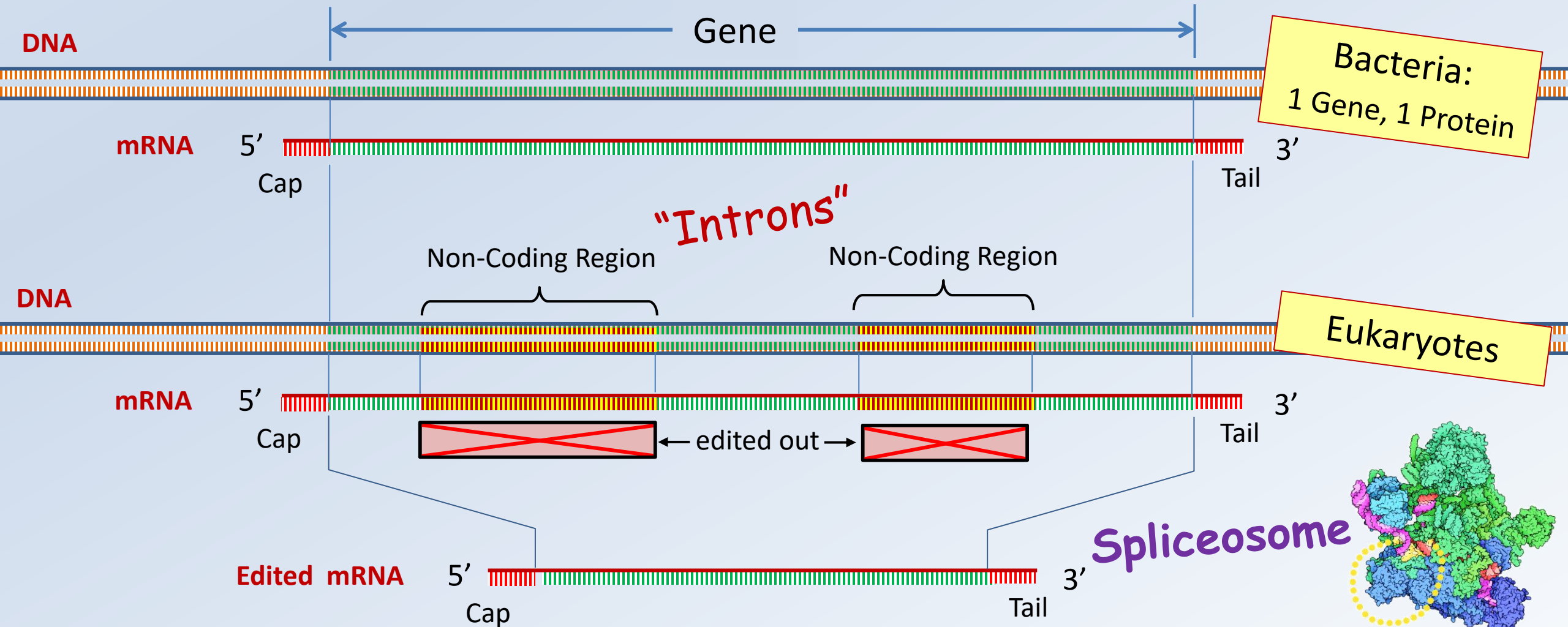
Introns



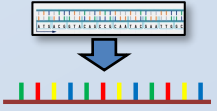
② RNA Transcription



Introns



② RNA Transcription



Yeast Spliceosome (at a particular stage)

Protein subunits



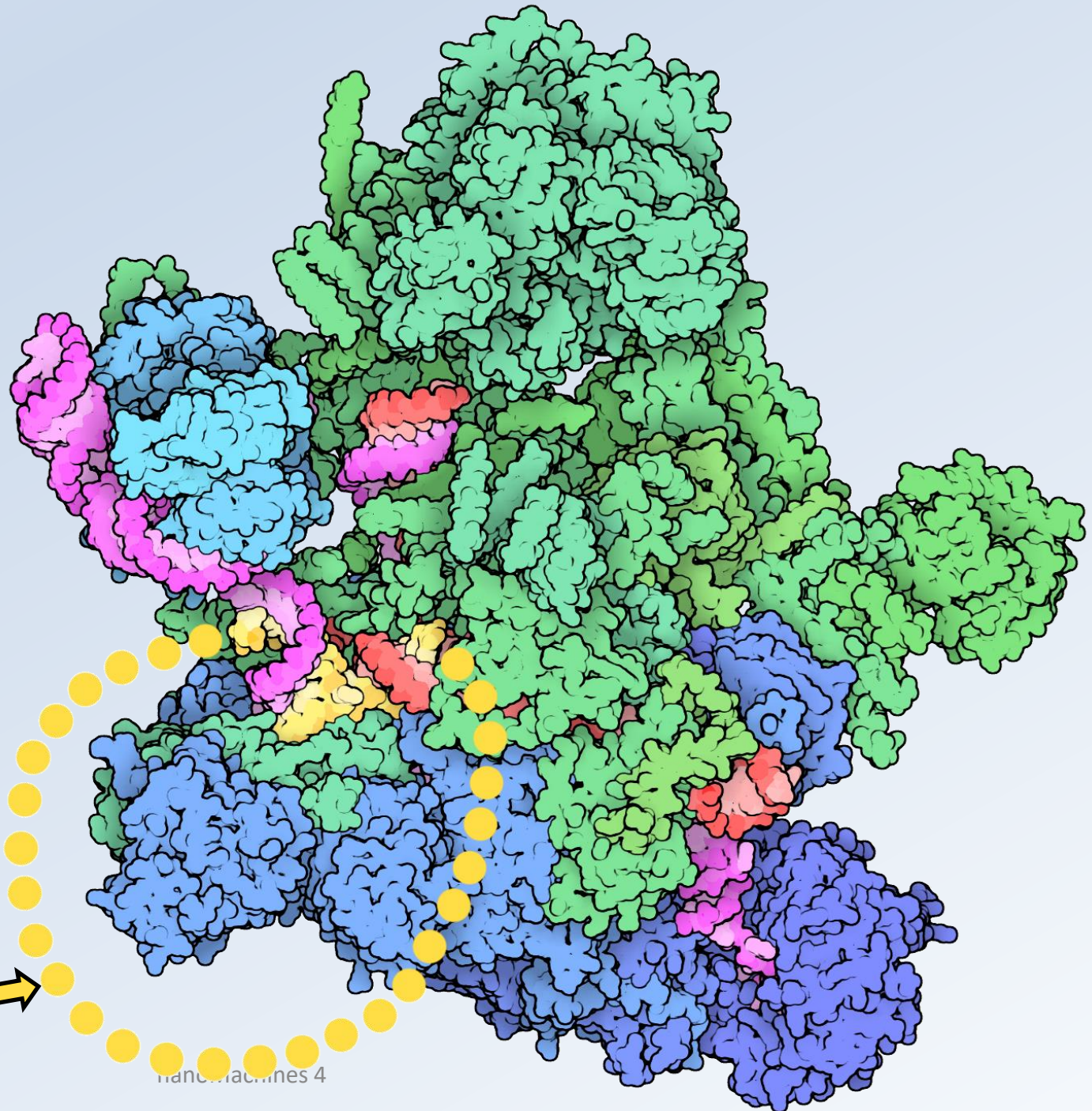
RNA structural subunits



Intron to be excised

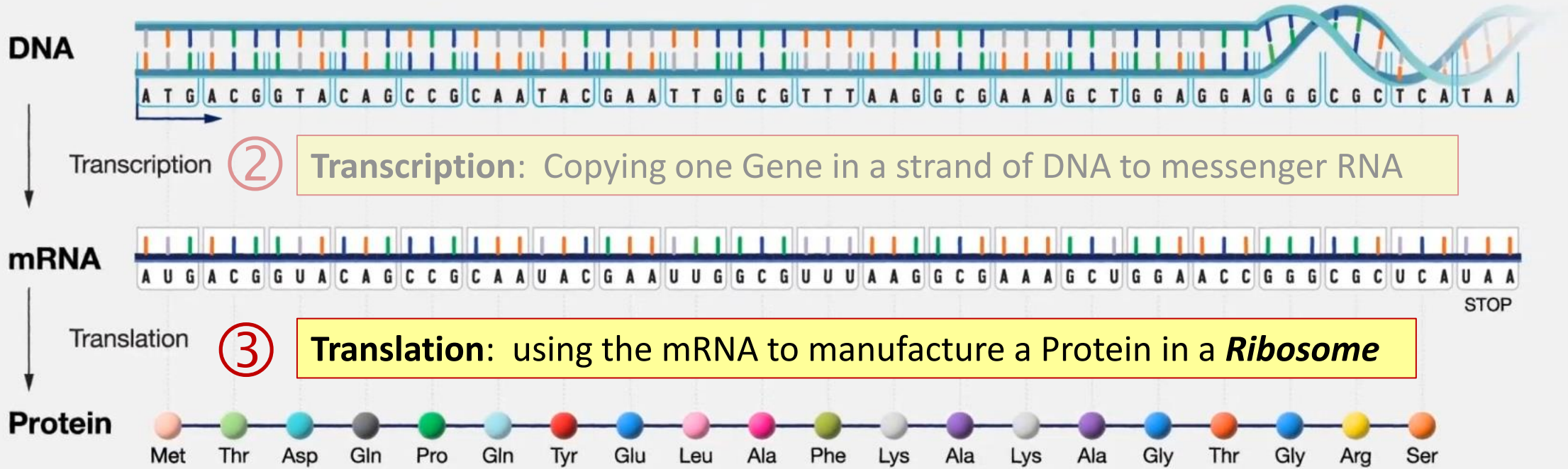


mRNA
Intron
"Lariat"

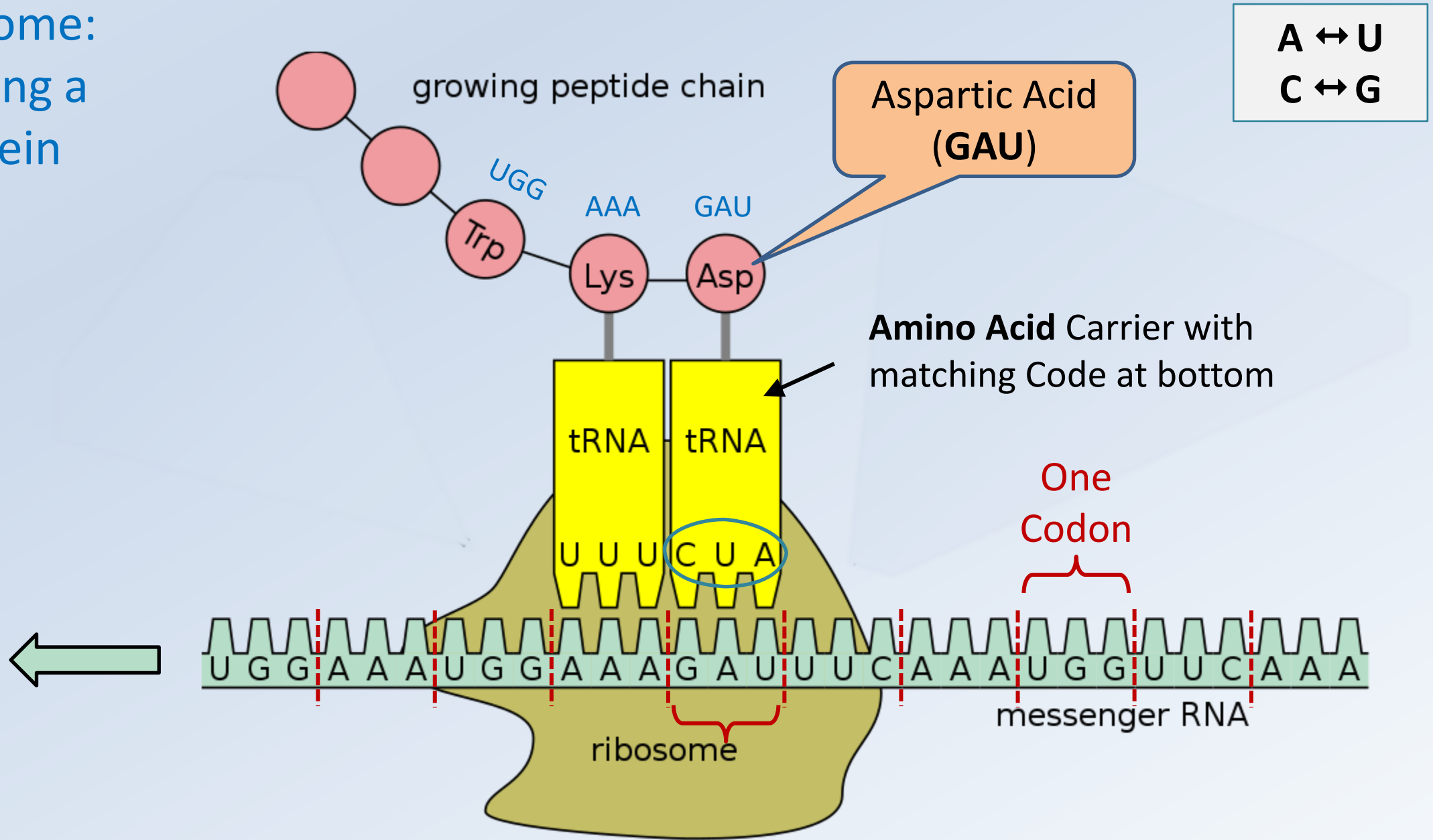


Central Dogma

① **Replication:** Duplicating DNA when cells divide.



Ribosome: Building a Protein

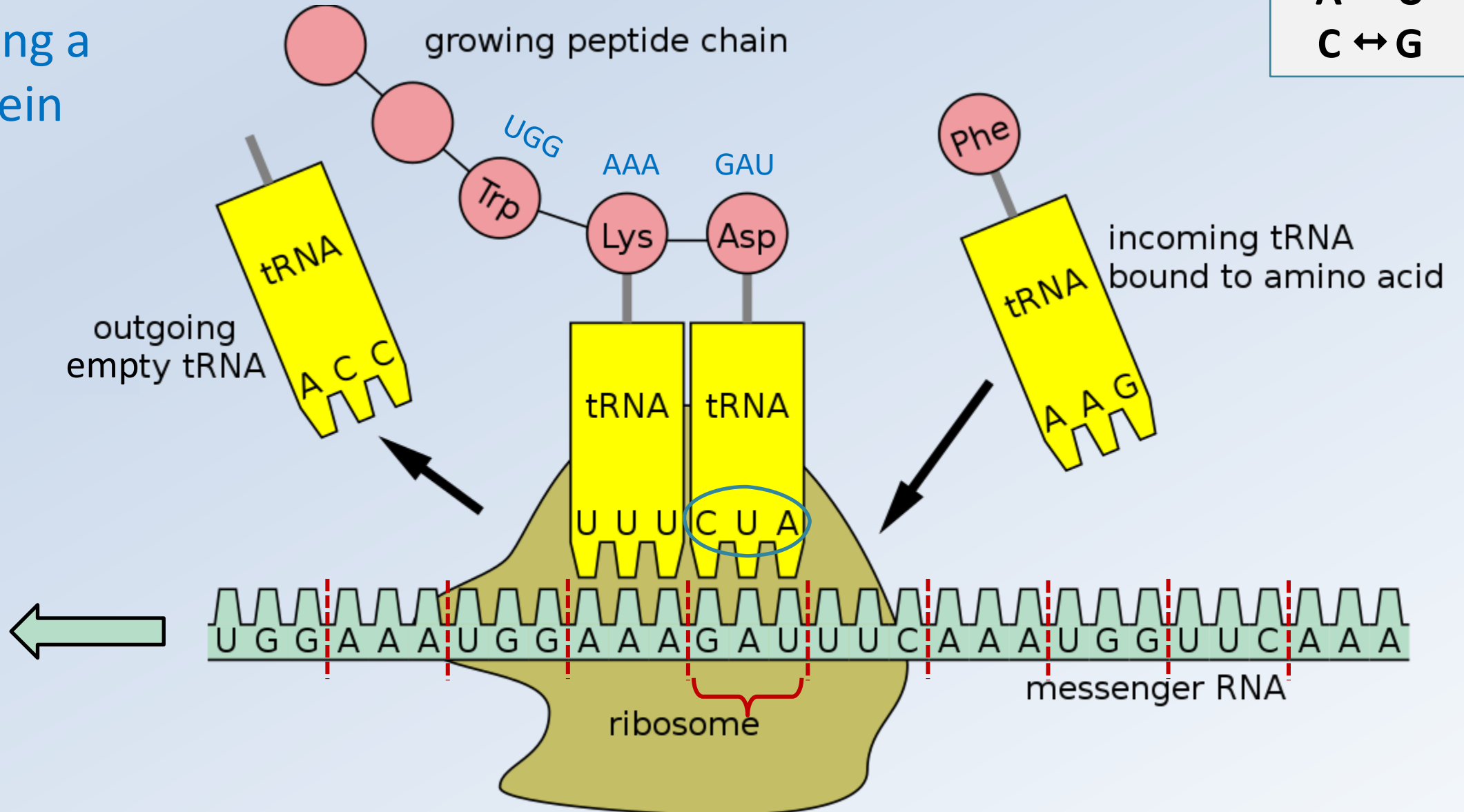


A ↔ U
C ↔ G



Ribosome: Building a Protein

A ↔ U
C ↔ G

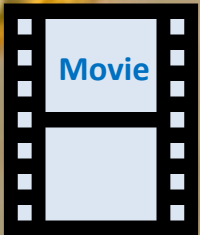
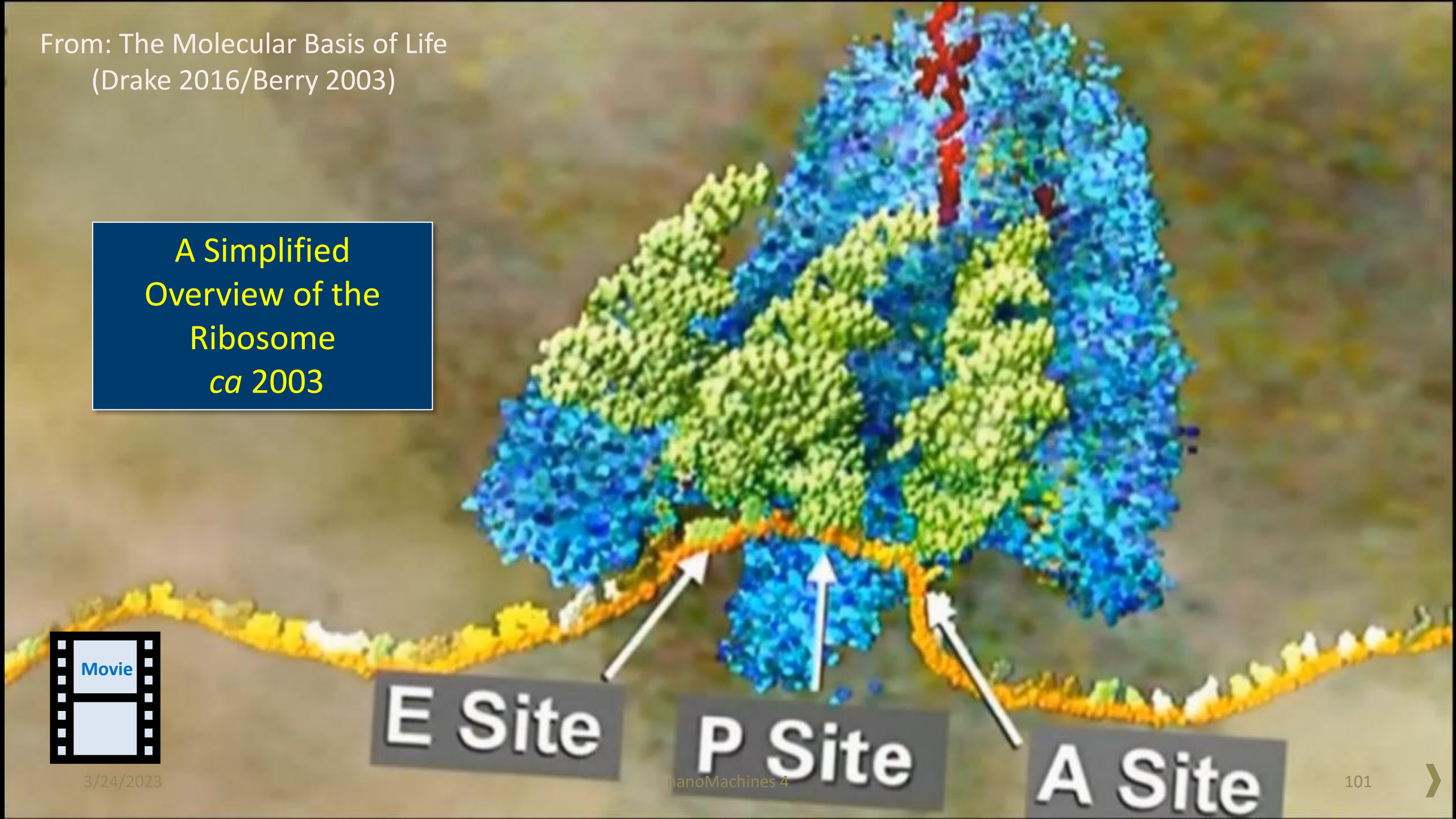


Peptide Synthesis



From: The Molecular Basis of Life
(Drake 2016/Berry 2003)

A Simplified
Overview of the
Ribosome
ca 2003



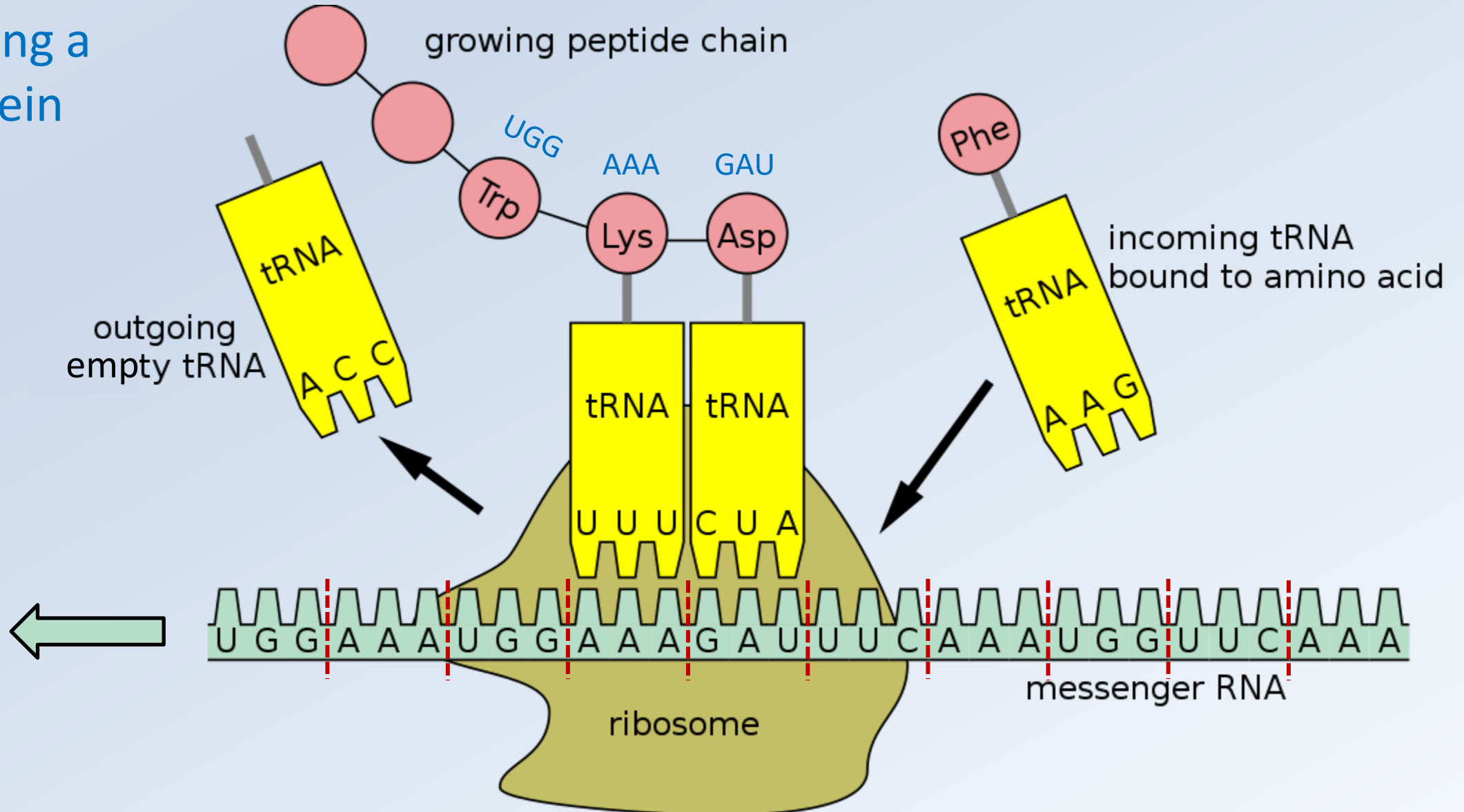
3/24/2023

nanoMachines 4

101



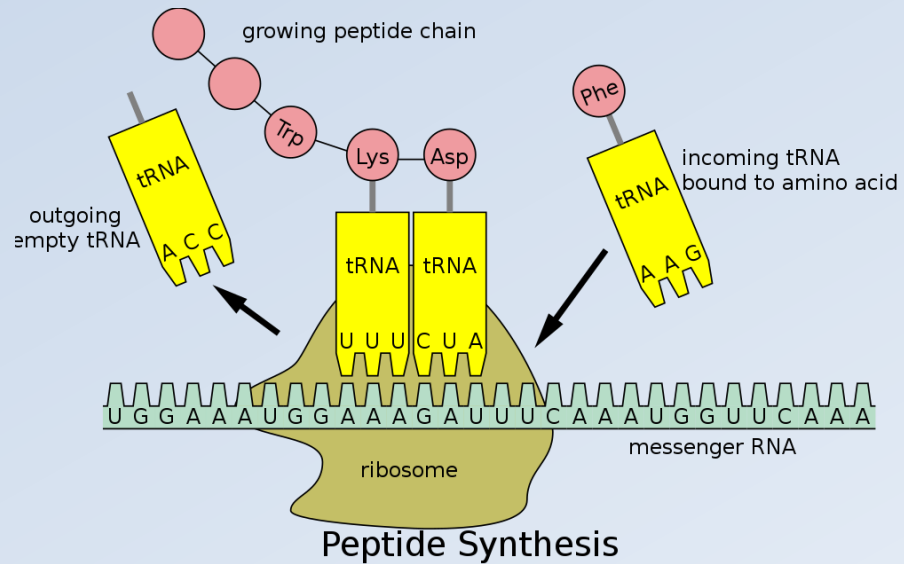
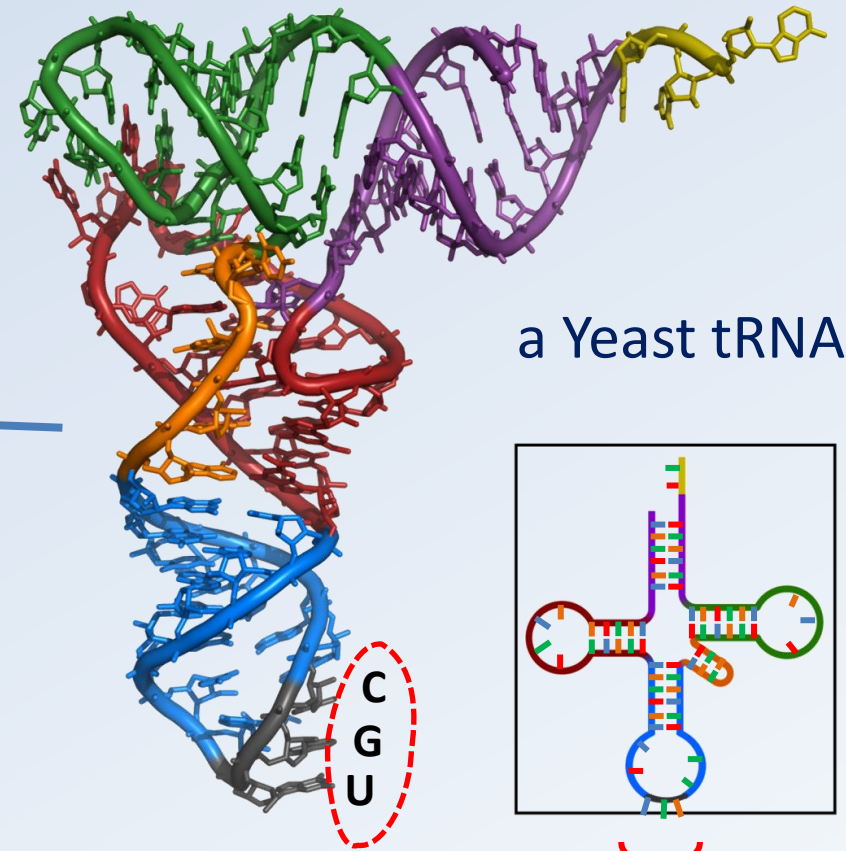
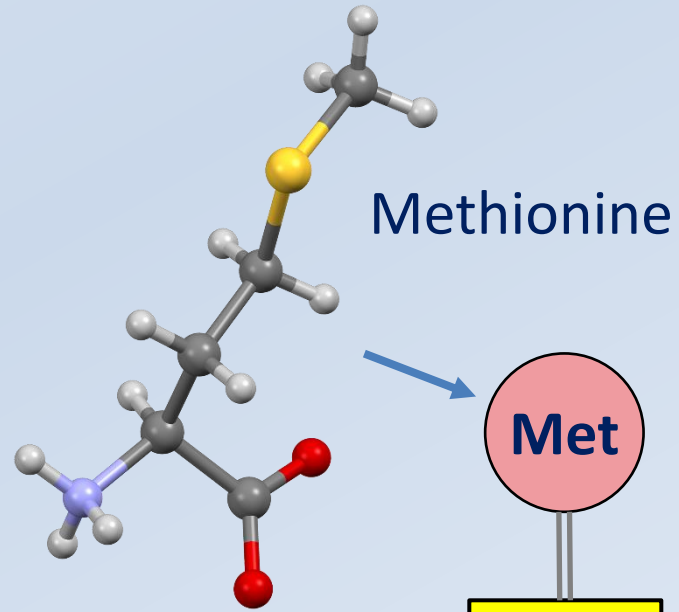
Ribosome: Building a Protein



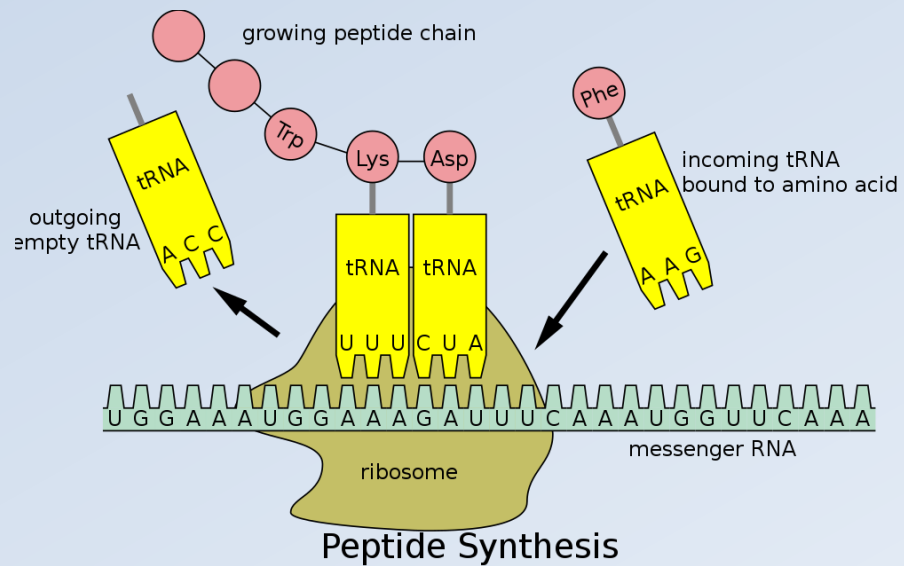
Peptide Synthesis



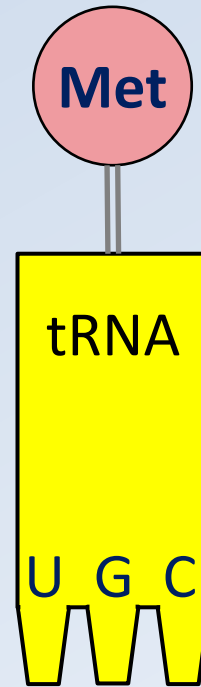
Ribosome: Building a Protein



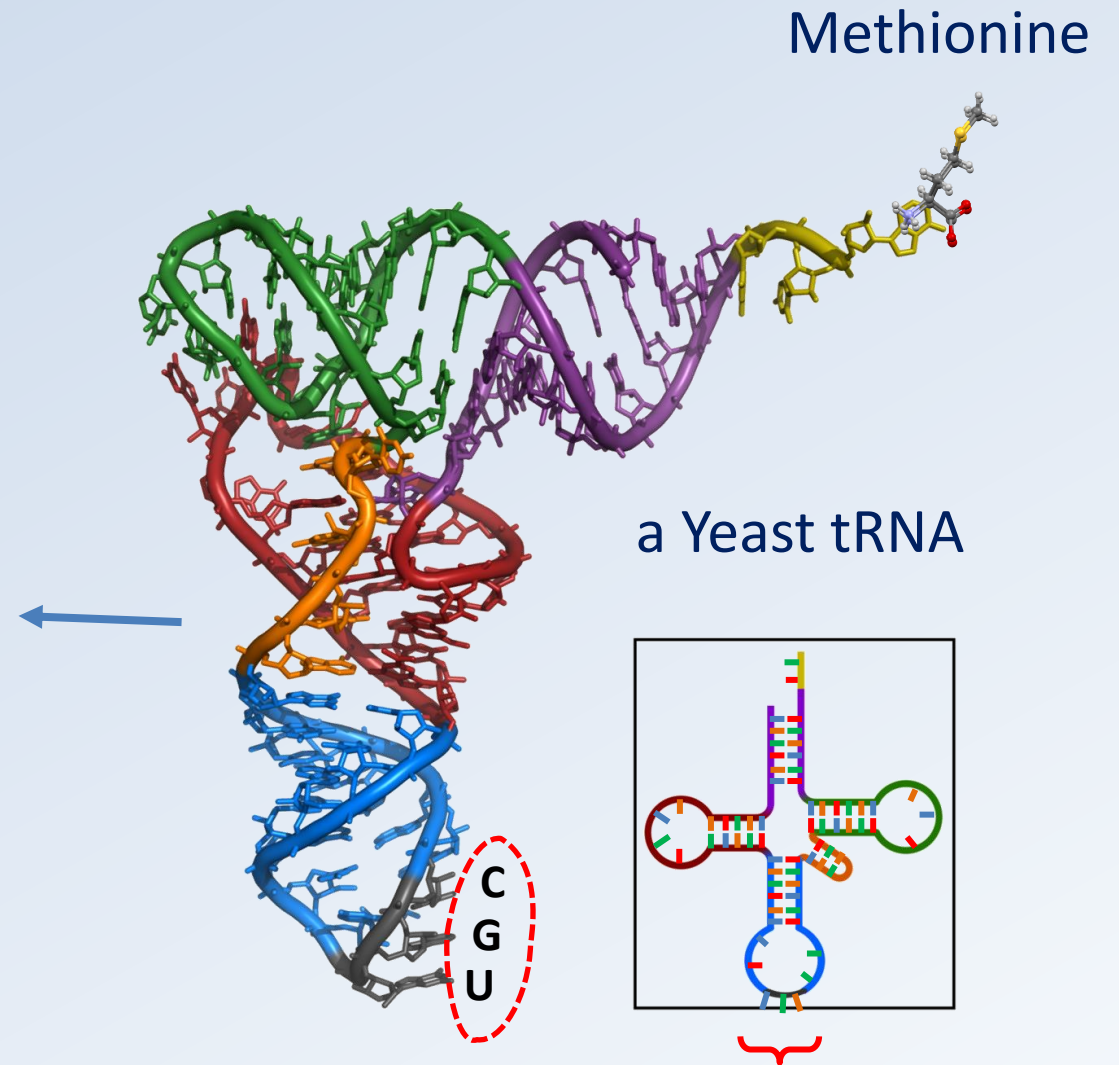
Ribosome: Building a Protein



3/24/2023



nanoMachines 4

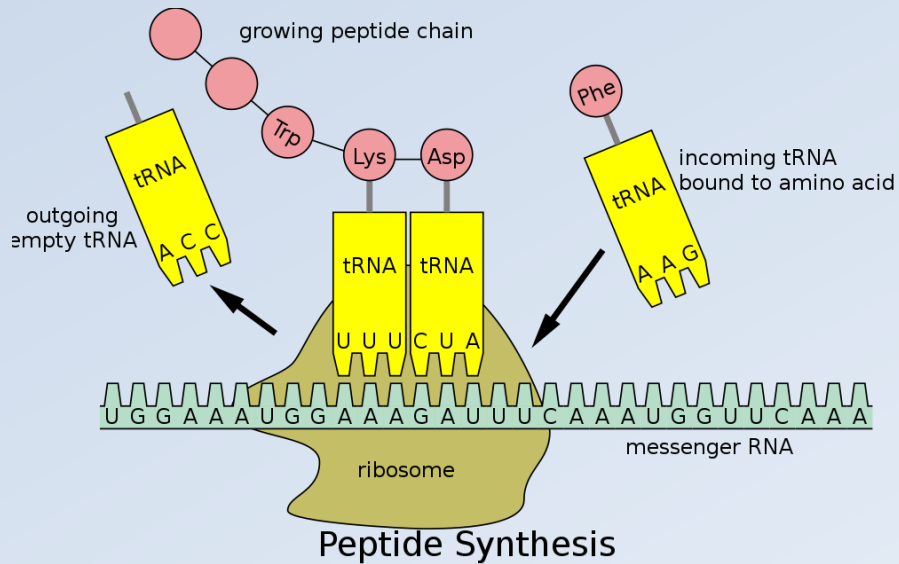
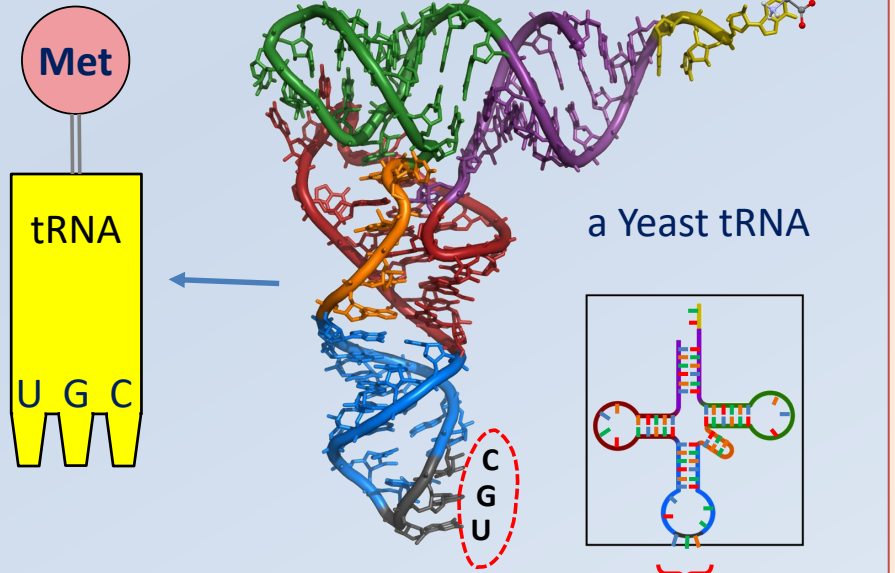


Wikipedia

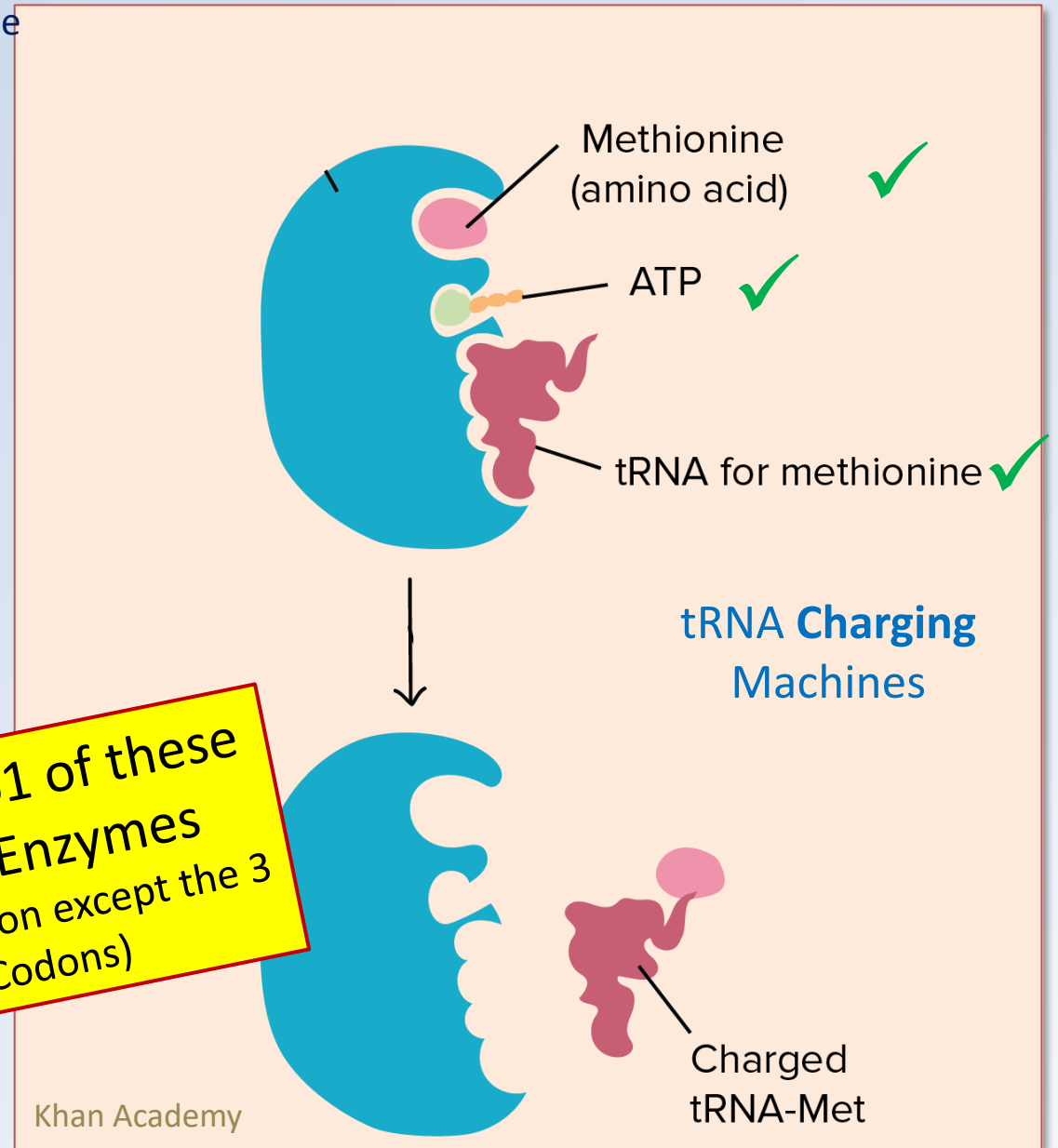
104



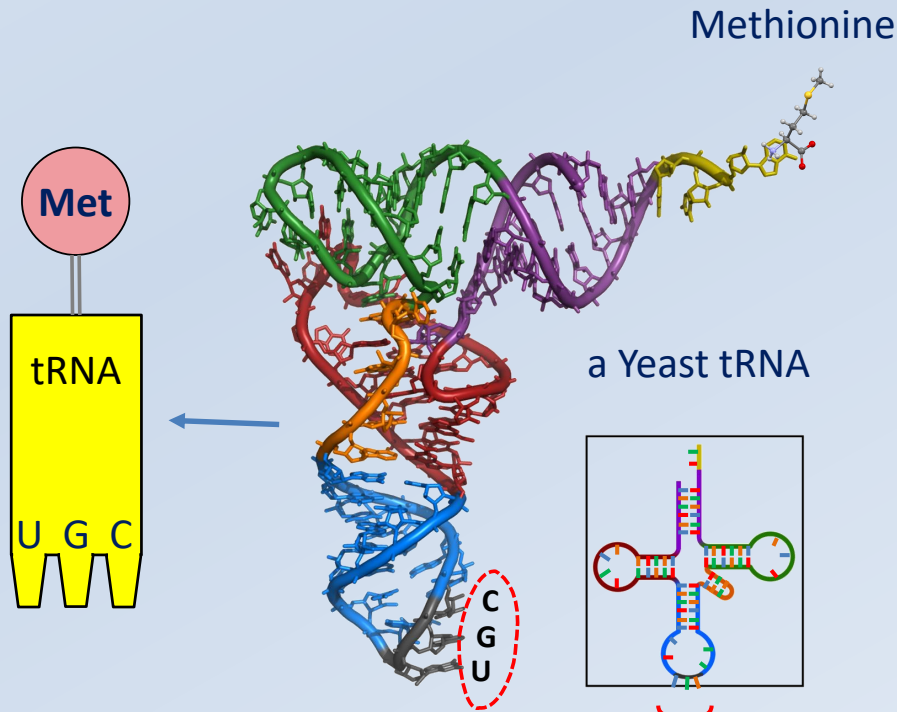
Ribosome: Building a Protein



We'll need 61 of these Charging Enzymes (1 for each Codon except the 3 STOP Codons)



Ribosome: Building a Protein

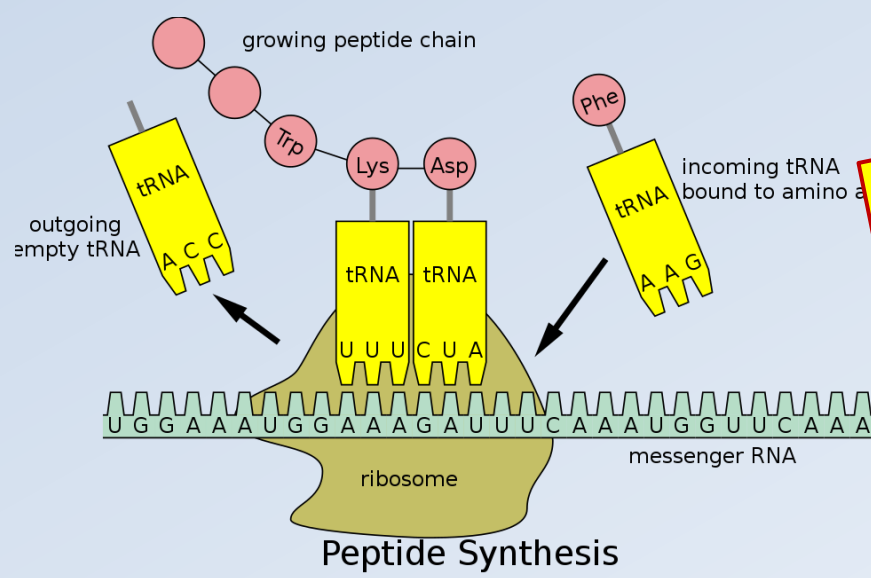
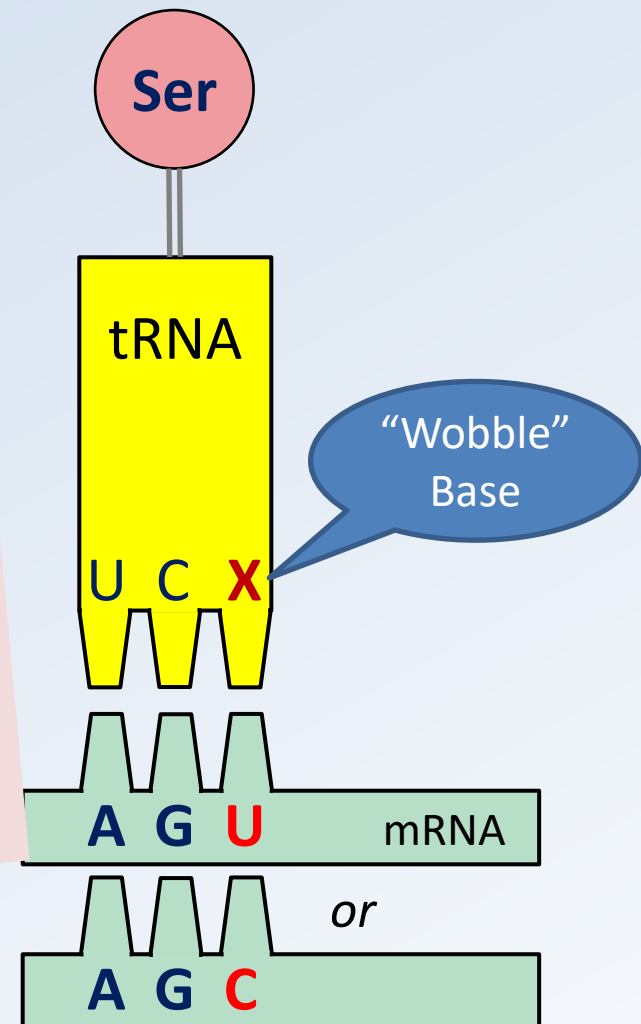


Example: **Serine** } 2 codes
AGU
AGC

A ↔ U
 C ↔ G

Nature's
 Hack

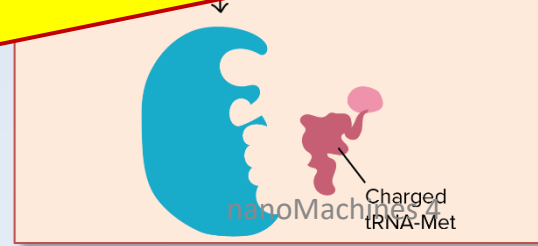
Using special
 "Wobble" bases,
 some tRNA's can
 cover several
 Codons for
 same Amino
 Acid



Aminoacyl-tRNA synthetase for methionine

40 or 50

We'll need ~~61~~ of these
 Charging Enzymes

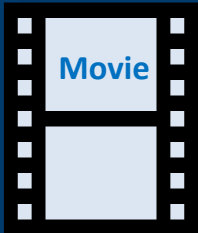


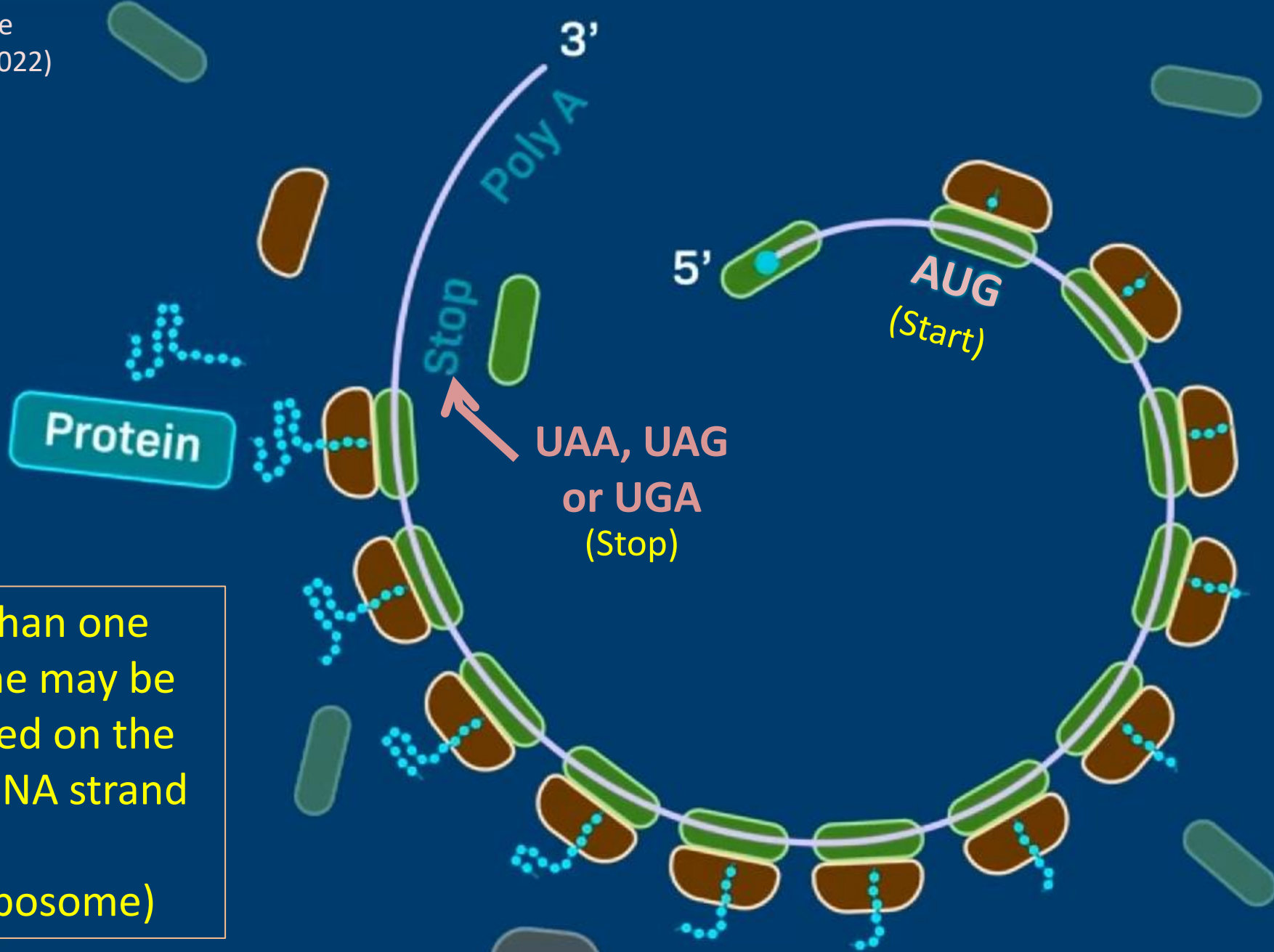
Polyribosome
(LabExchange 2022)

UAA, UAG
or UGA

More than one
Ribosome may be
assembled on the
same mRNA strand

(PolyRibosome)



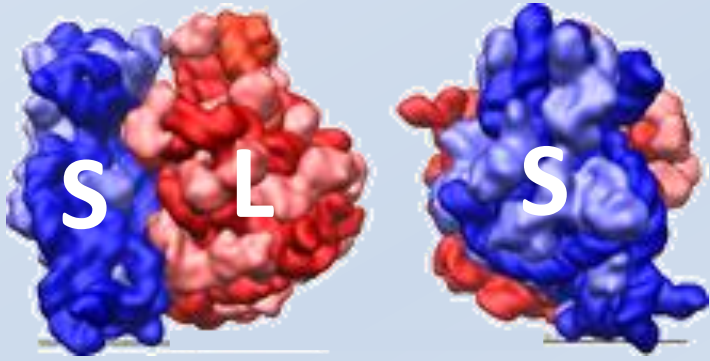


More than one
Ribosome may be
assembled on the
same mRNA strand

(PolyRibosome)

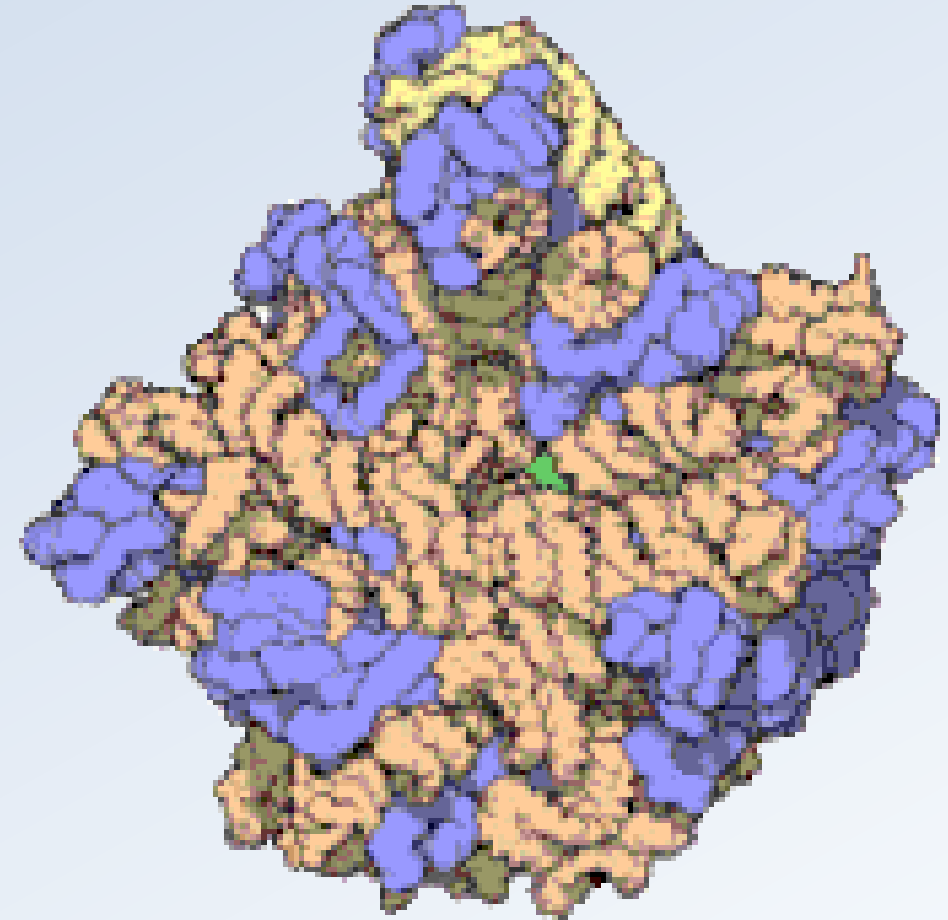
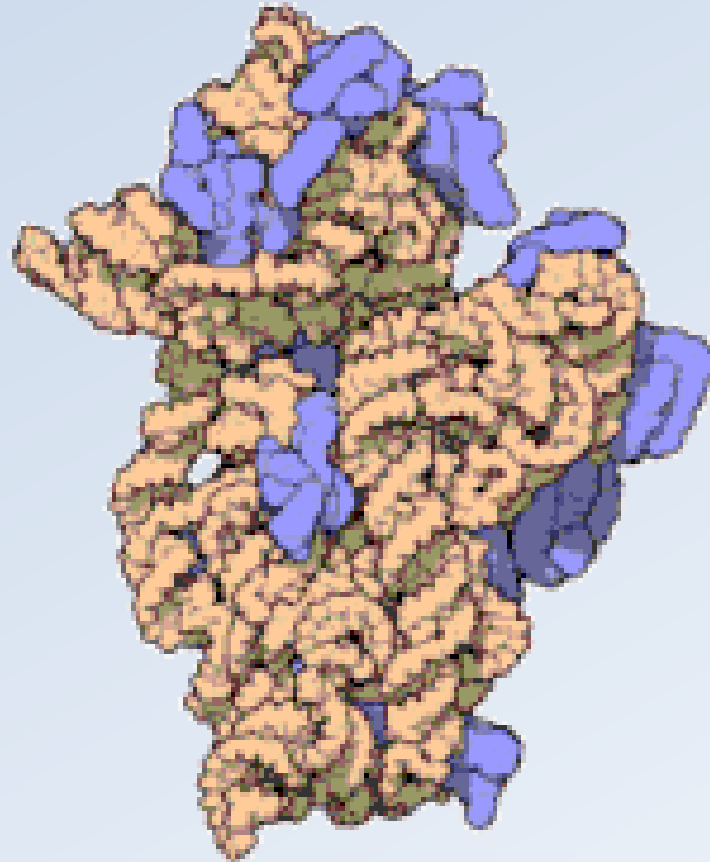


Ribosome constructed of 2 Subunits: Large and Small



Small subunit

Large subunit



- Proteins ■ 53 79
- rRNA's ■■■ 5 6
- Nucleotides 4343 5107

~65% RNA

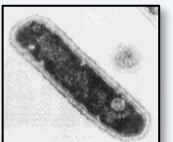
Eukaryotes

100% of key components are RNA's!

Thermus thermophilus

nanoMachines 4

Wikipedia



E Coli Ribosome

(Large & Small Subunits assembled, but without tRNA's or mRNA)

Note RNA elements



Settings / Controls Info

Animate	Spin	
Camera	Perspective	...
Background		...
Occlusion	✓ On	...
Shadow	✗ Off	...
Outline	✗ Off	...
Fog	✓ On	...
Clipping	<input type="range" value="74"/>	74 ...
Layout	1 of 1	

▼ Moving in 3D

Rotate
Drag using left mouse button

Rotate around z-axis

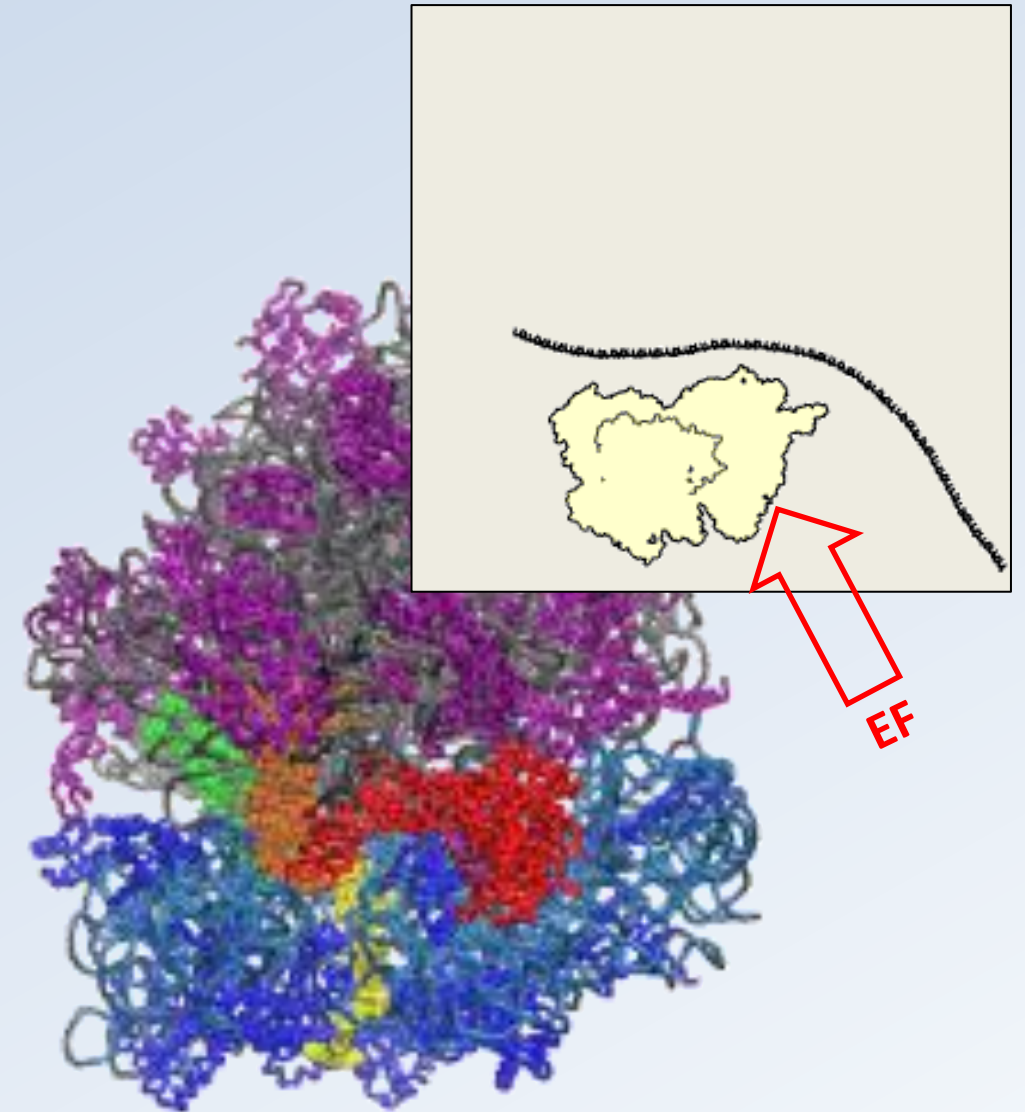
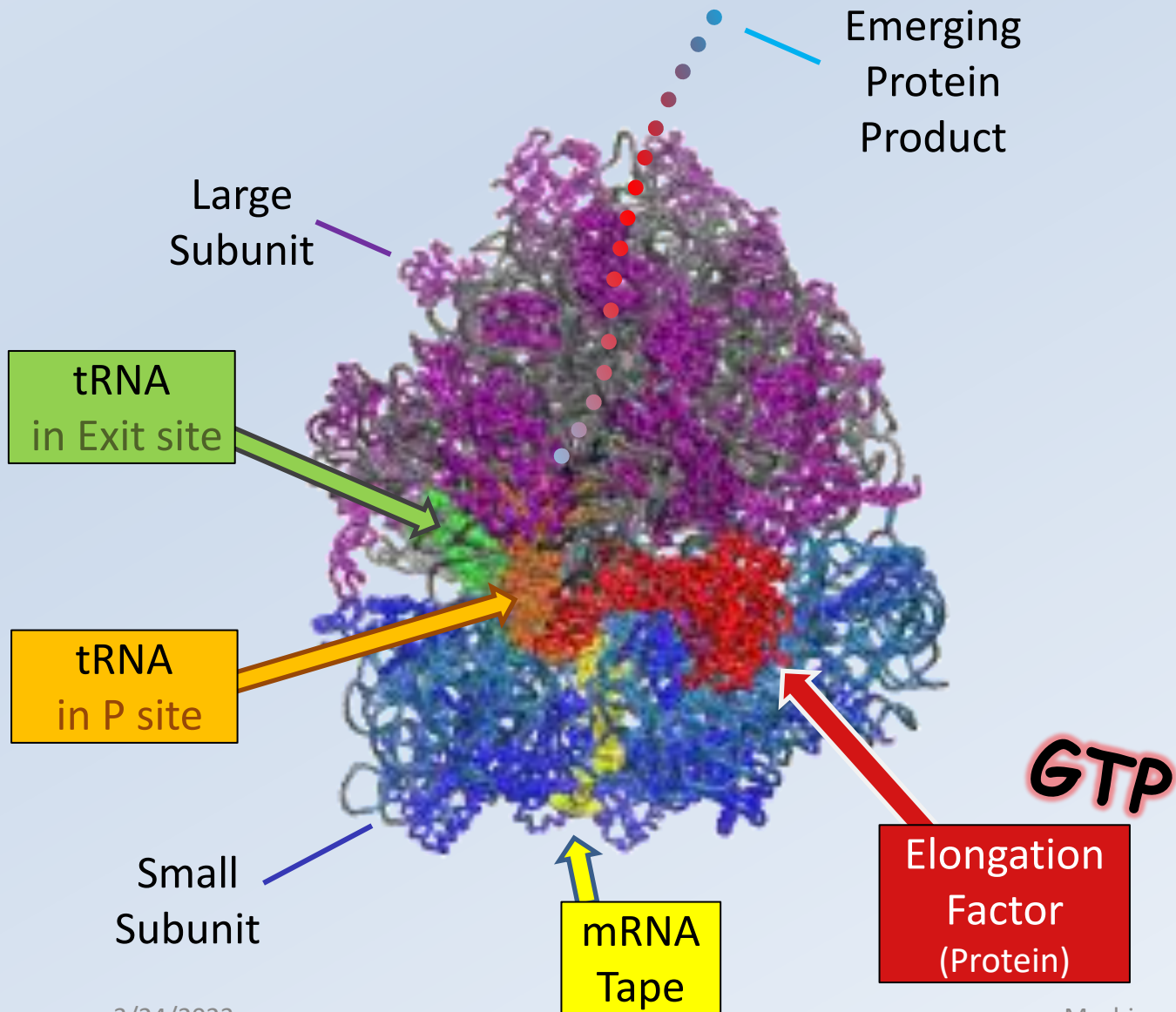
PDB: code 4v4a

Credit: RCSB.org

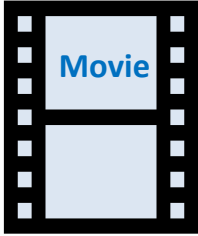


Ribosome "Innards" in Action

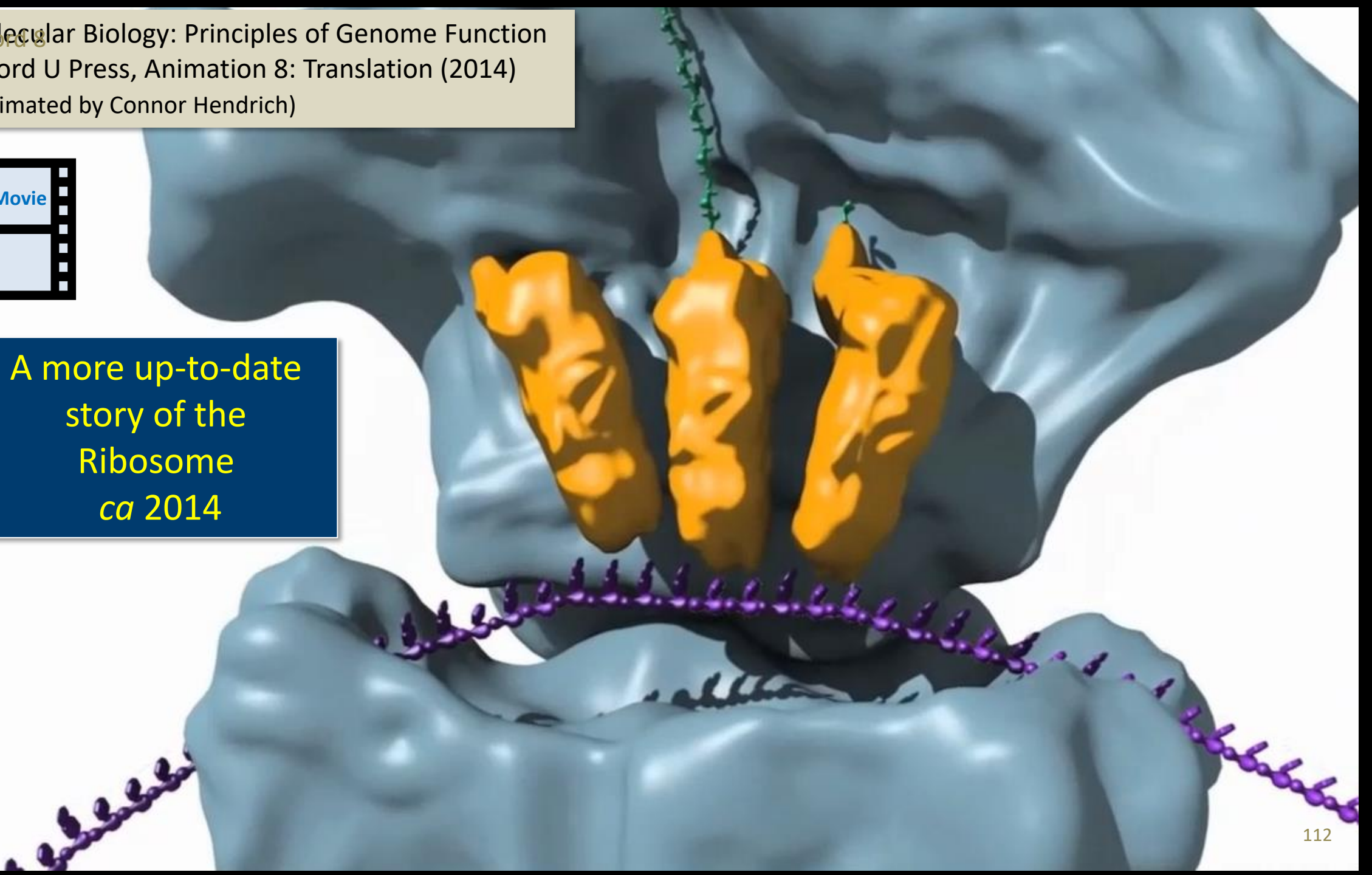
Thermus thermophilus Bacterium



Oxford
Molecular Biology: Principles of Genome Function
Oxford U Press, Animation 8: Translation (2014)
(Animated by Connor Hendrich)



A more up-to-date
story of the
Ribosome
ca 2014



Questions?