

Lecture #5

“DNA repair & recombination”

Jan 22, 2018

RECAP

How cells protect DNA ends (telomeres)

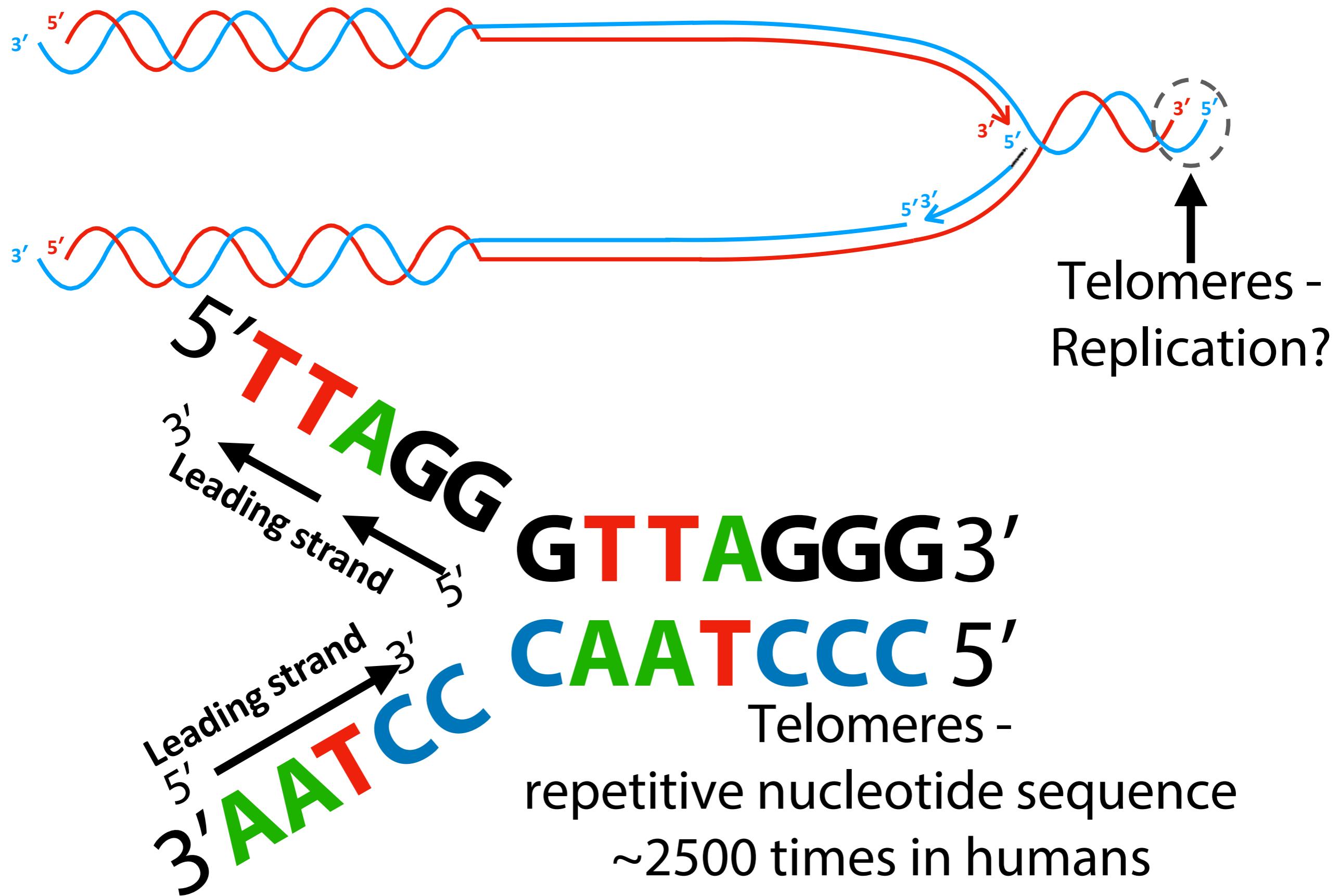


200
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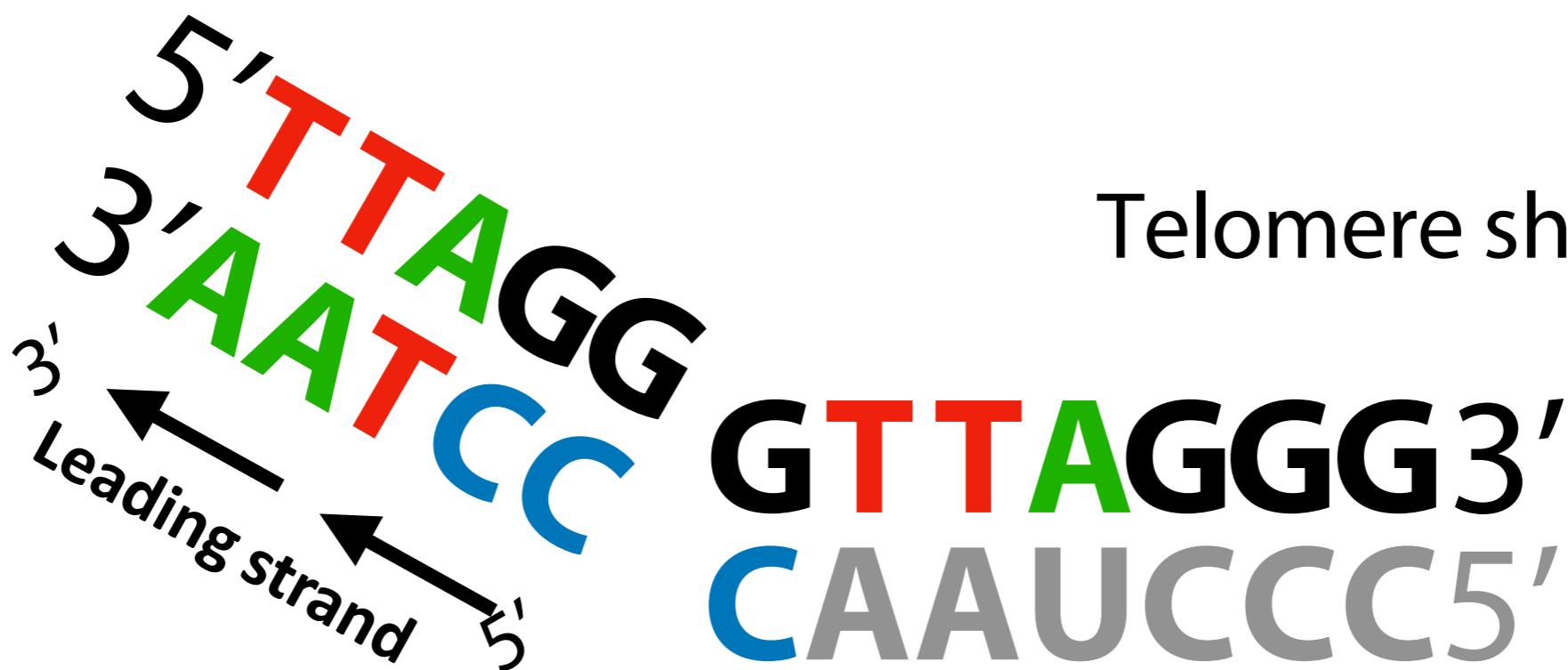
Elizabeth H. Blackburn, Carol W. Greider & Jack Szostak
(2009)

How chromosomes are protected by telomeres and the
enzyme telomerase

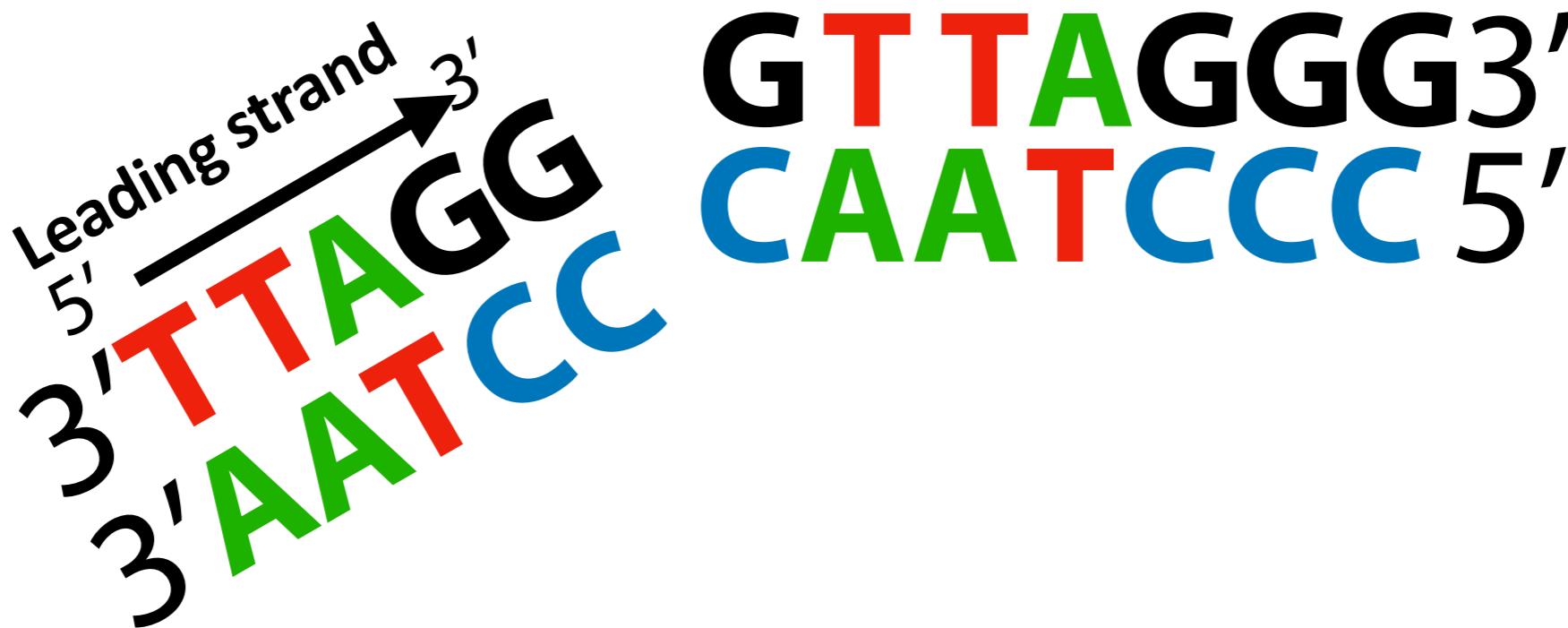
Replication at the ends of chromosome



Replication at the ends of chromosome



Telomere shortening



How cells protect telomeres

5' **TTAGGGT** A GGG 3'

3' **AATC** CCC AAUCCC 5'



Human TERT

TElomerase Reverse Transcriptase

5' **TTAGGGT** TA GGG **TTA** G 3'

3' **AATC** CCC AAUCCC AAUCCC. 5'

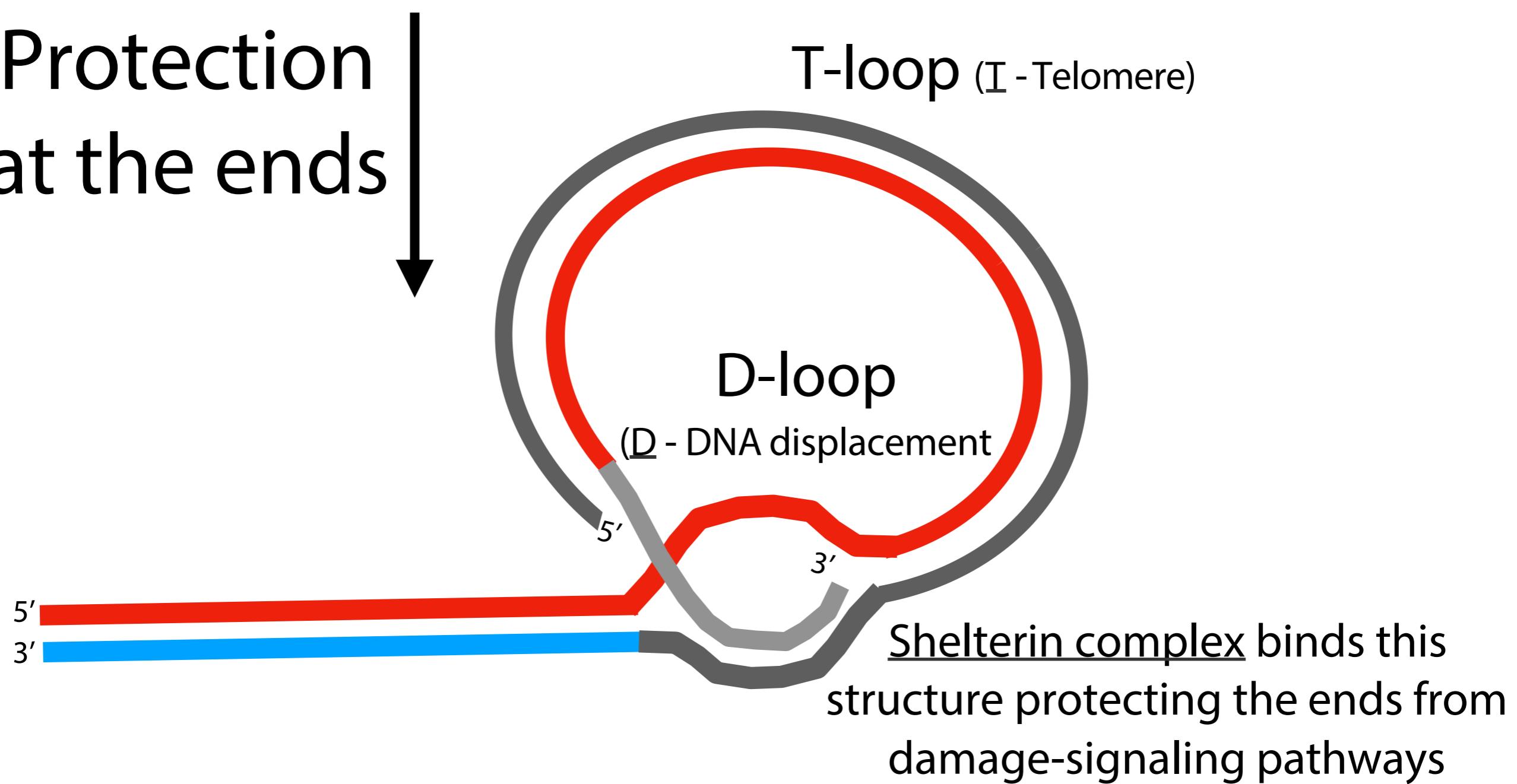
hTERT

RNA component
used as a template

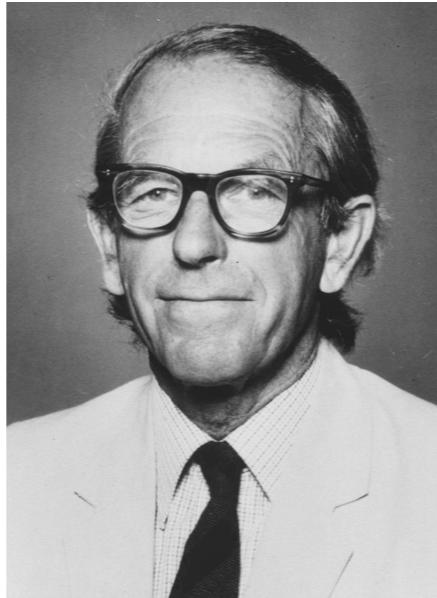
Protection at the ends of DNA



Protection
at the ends



DNA sequencing



198

0

Paul
Berg

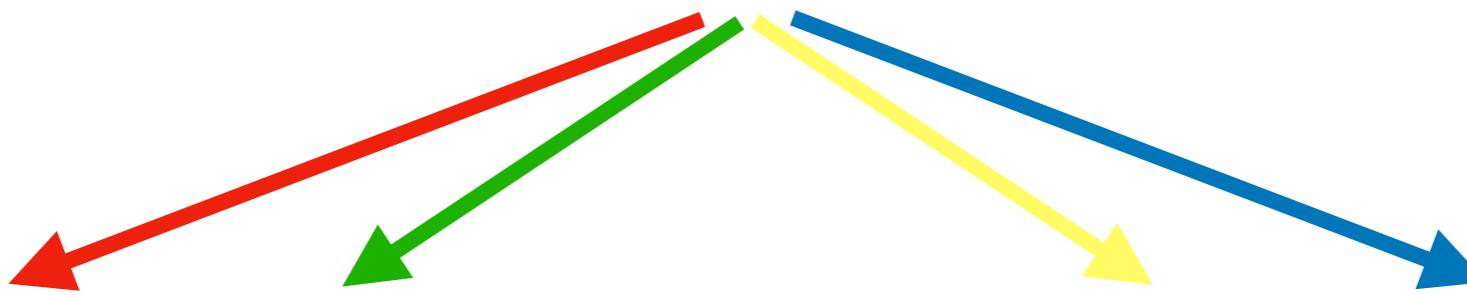
Frederick
Sanger 2
Nobel
prizes

Walter
Gilbert

One-half to Berg for his studies of nucleic acids relevant to recombinant DNA. The second half shared between Sanger and Gilbert for their work of determining base sequence of nucleic acids.

DNA sequencing (Sanger)

5't t t t **TATCGCTCTGAC**3'
3'aaaa **ATAGCGAGACTG**5'



Separate the two DNA strands

Add DNA polymerase

Add dNTPs

ddTTP*

ddATP*

ddGTP*

ddCTP*

Add primer

5't t t t
3'aaaa

ATAGCGAGACTG5'

DNA sequencing

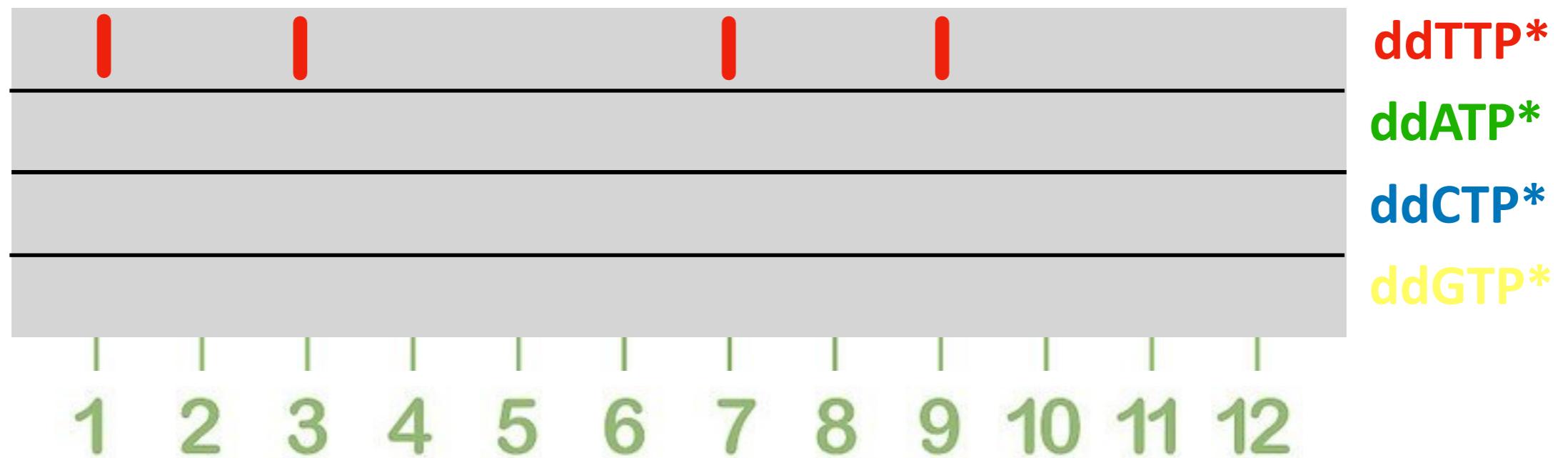
ddTTP*

5' T
3' ATAGCGAGACTG 5'

5' T A T
3' ATAGCGAGACTG 5'

5' TATCGC T
3' ATAGCGAGACTG 5'

5' T A T C G C T C T
3' ATAGCGAGACTG 5'



DNA sequencing

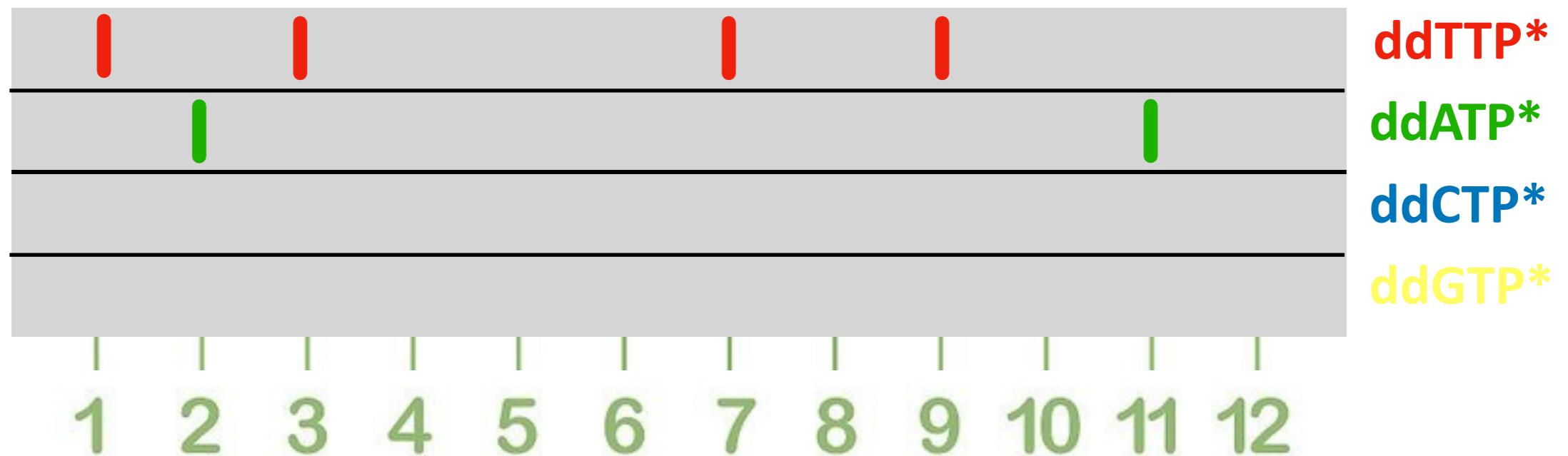
ddATP*

5' T A

3' ATAGCGAGACTG 5'

5' TATCGCTCTG A

3' ATAGCGAGACTG 5'



DNA sequencing

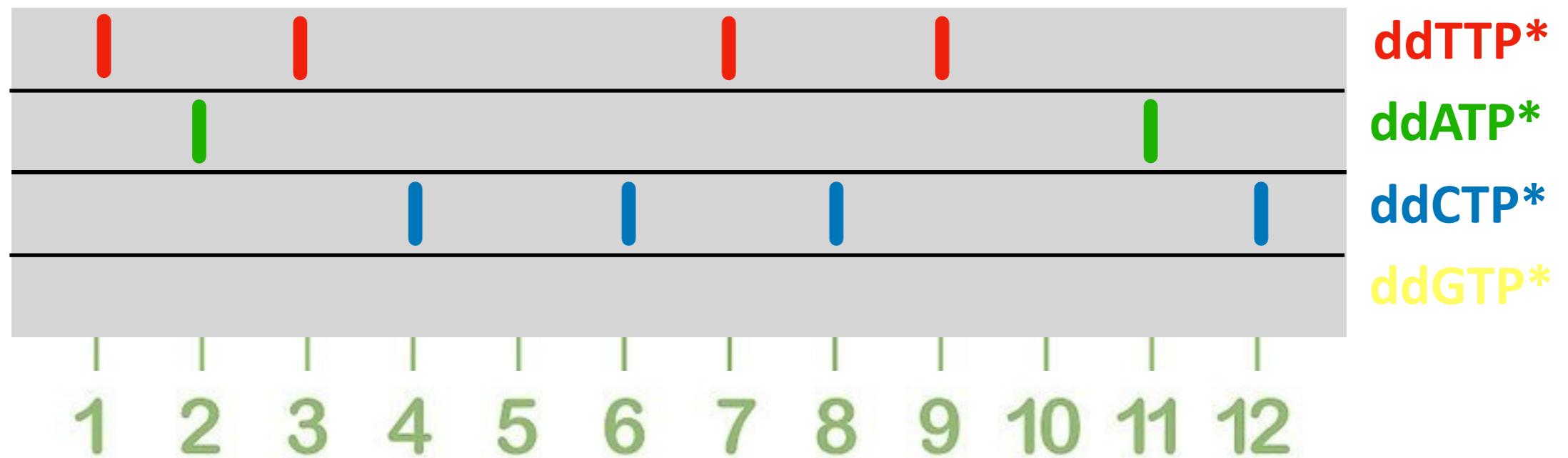
ddCTP*

5' T A T C
3' A T A G C G A G A C T G 5'

5' T A T C G C
3' A T A G C G A G A C T G 5'

5' T A T C G C T C
3' A T A G C G A G A C T G 5'

5' T A T C G C T C T G A C
3' A T A G C G A G A C T G 5'

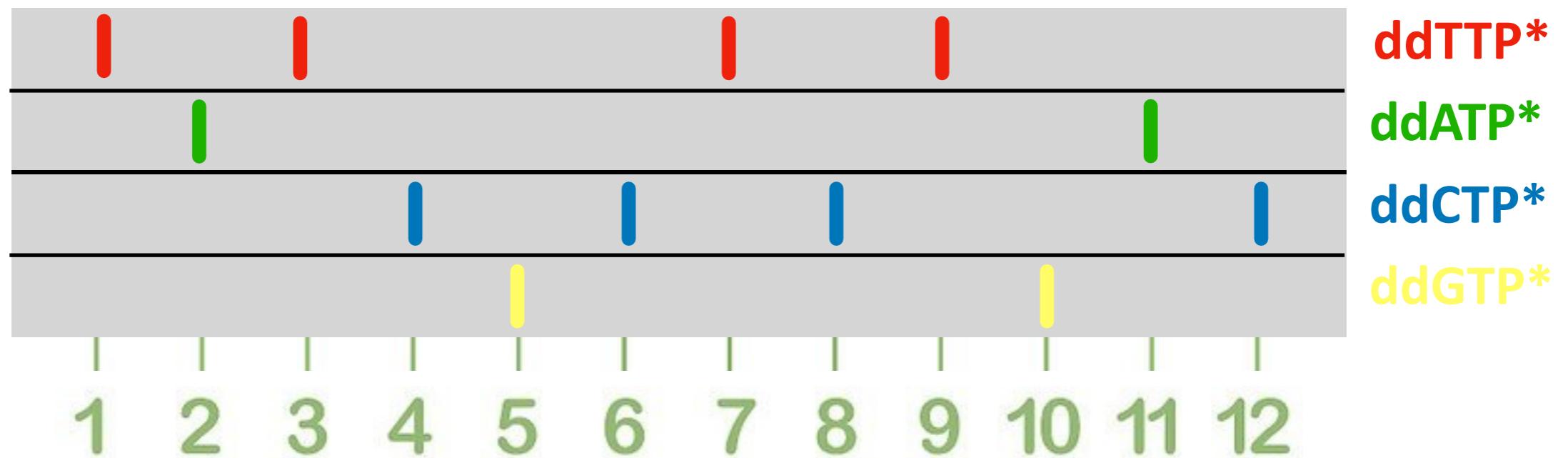


DNA sequencing

ddGTP*

5' T A T C G
3' A T A G C G A G A C T G 5'

5' T A T C G C T C T G
3' A T A G C G A G A C T G 5'

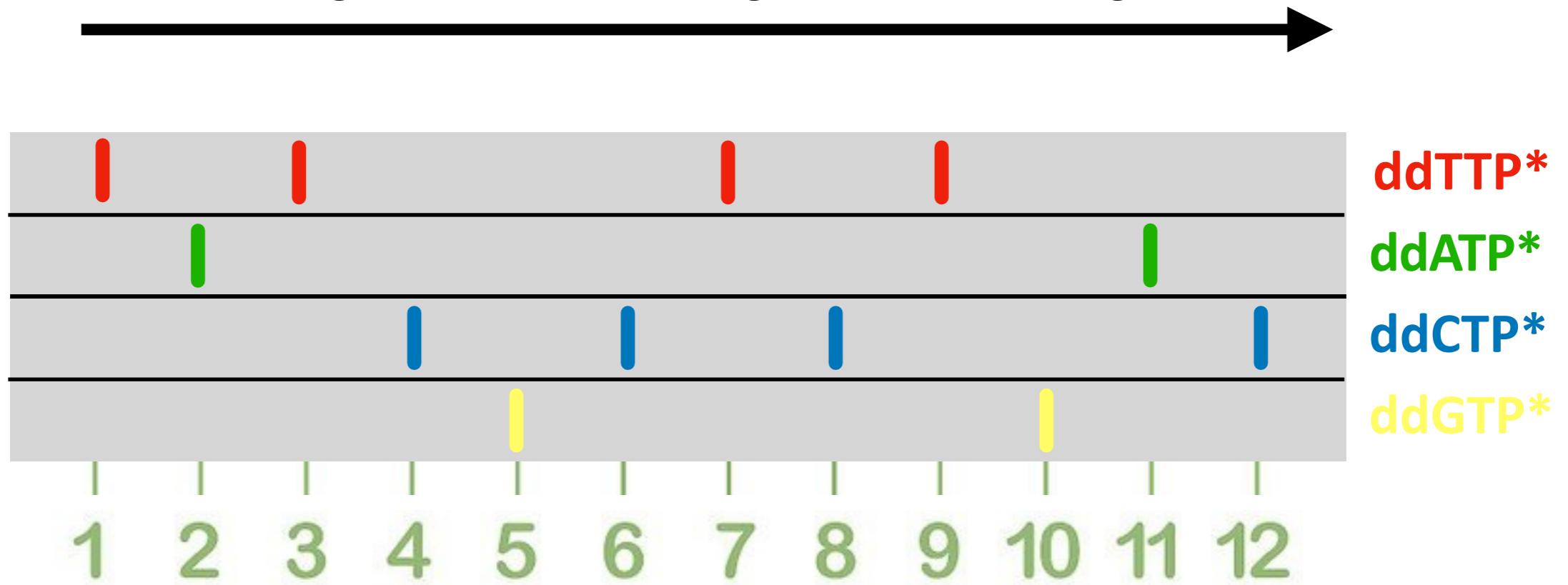


DNA sequencing

5' **TATCGCTCTGAC** 3'

3' ATAGCGAGACTG 5'

Reading direction lower to higher molecular weight



DNA sequencing

Sequencing method	Read length	Accuracy	Reads per run	Time per run
Sanger	~1000bp	99.9%	500-1000bp	~1-2hrs
Single molecule pacific biosciences	~14,000bp - 40,000bp	87%	50,000bp-~1000 Mbp	~2-4hrs
Sequencing by synthesis (Illumina)	~75-1000bp	99.9%	1-3billion bp	1-10 days
Nanopore sequencing	~500 kbp	~90%	variable	In real time as sequencing takes place

DNA sequencing

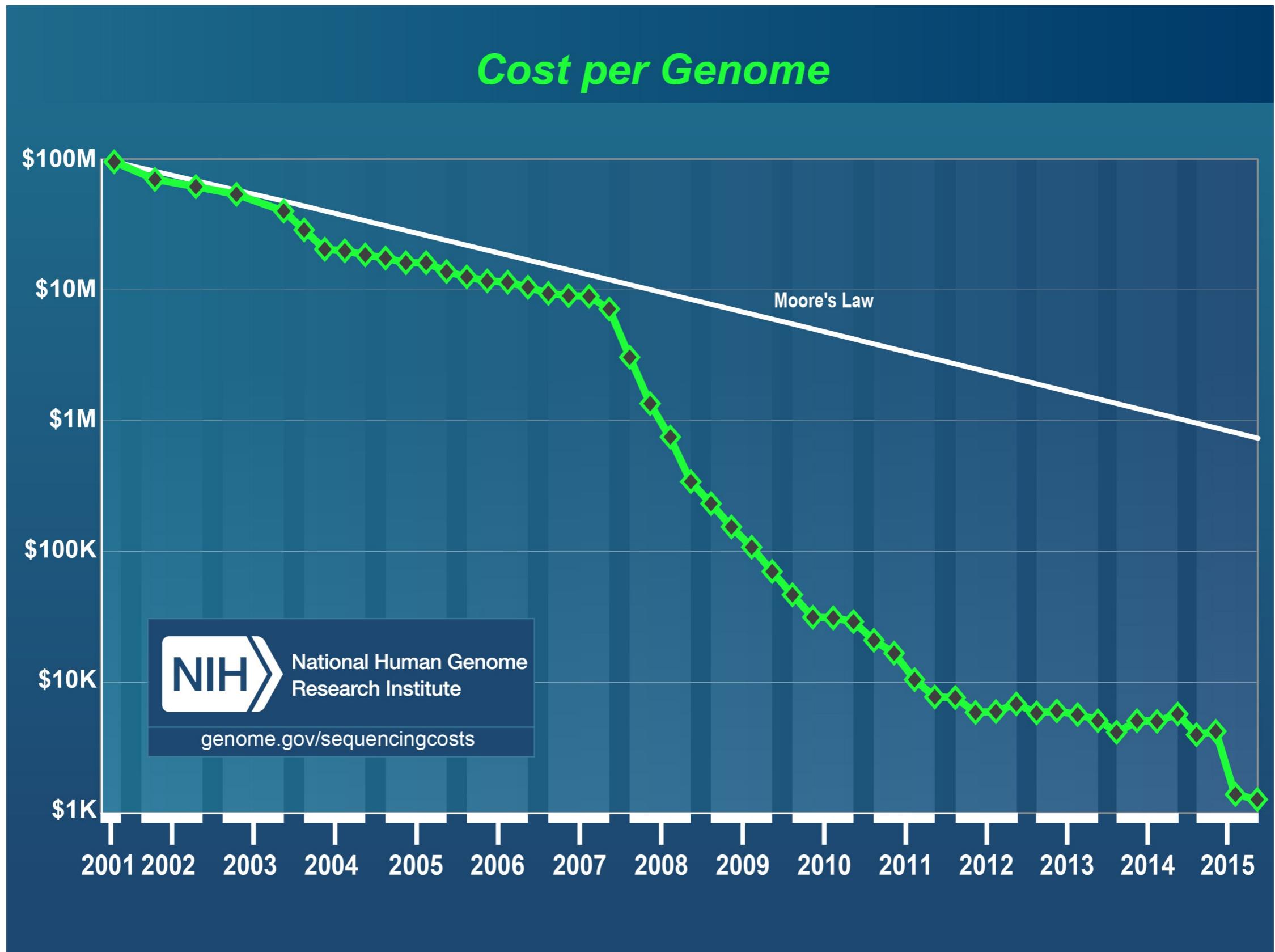
Pop quiz

Sequencing the first human genome cost \$1 billion and took >10 years. Now the cost has dropped by a factor of:

- (A) ten
- (B) a hundred
- (C) a thousand
- (D) a hundred-thousand
- (E) a million

& only takes few days. Now, millions of genomes are being sequenced

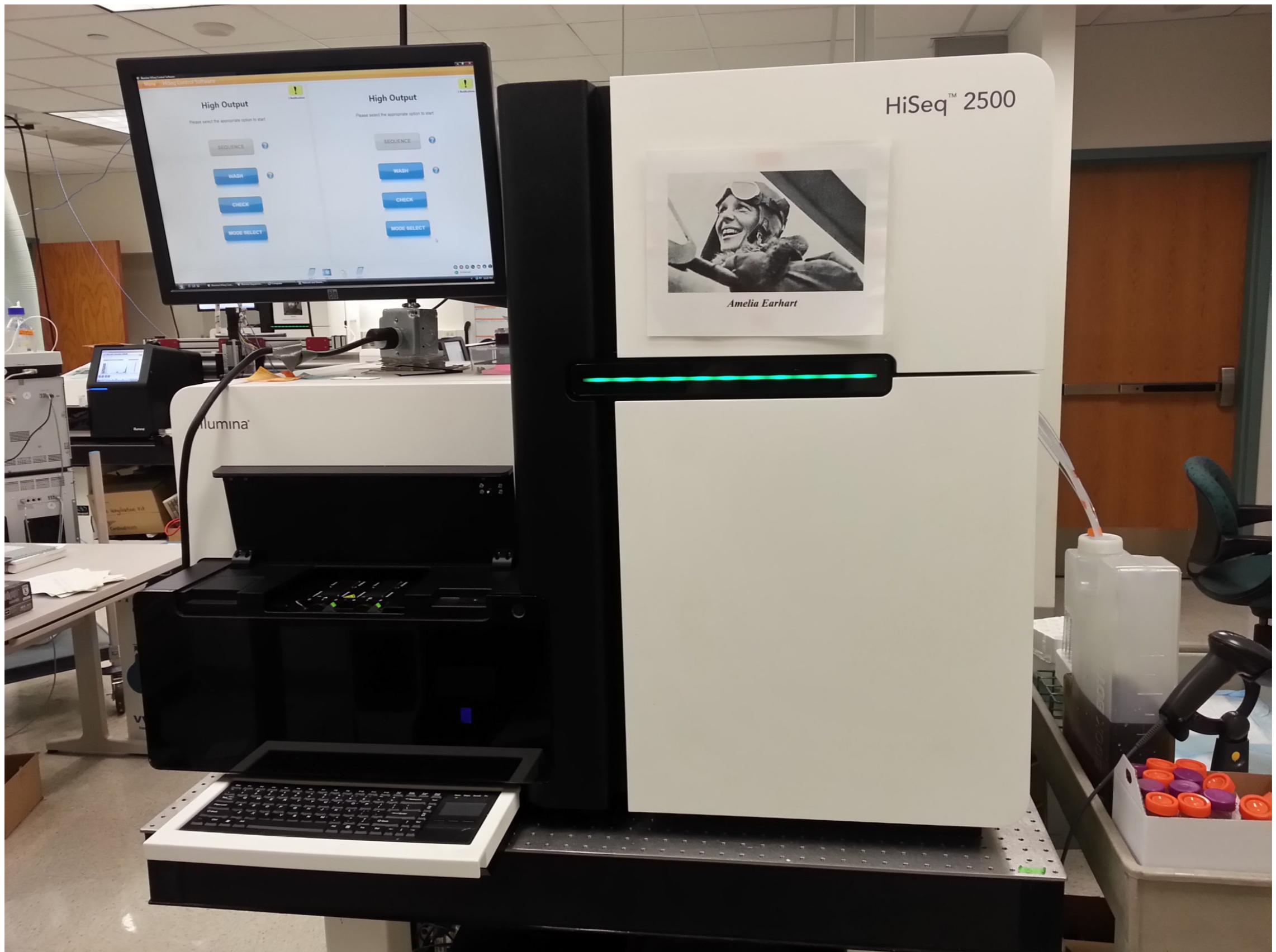
DNA sequencing cost



Human genome project



Now



Future of DNA sequencing



Personalized DNA sequencing



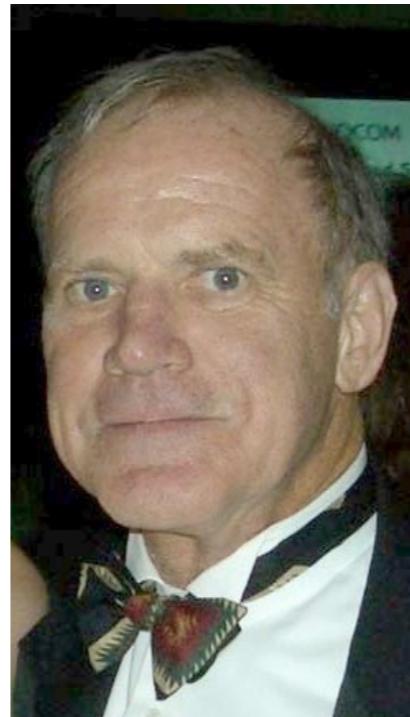
**Sequencing fetus
using DNA from the
mother**



**Sequence your
genome for \$99**

Polymerase Chain Reaction

How to make many copies of DNA in a test tube?

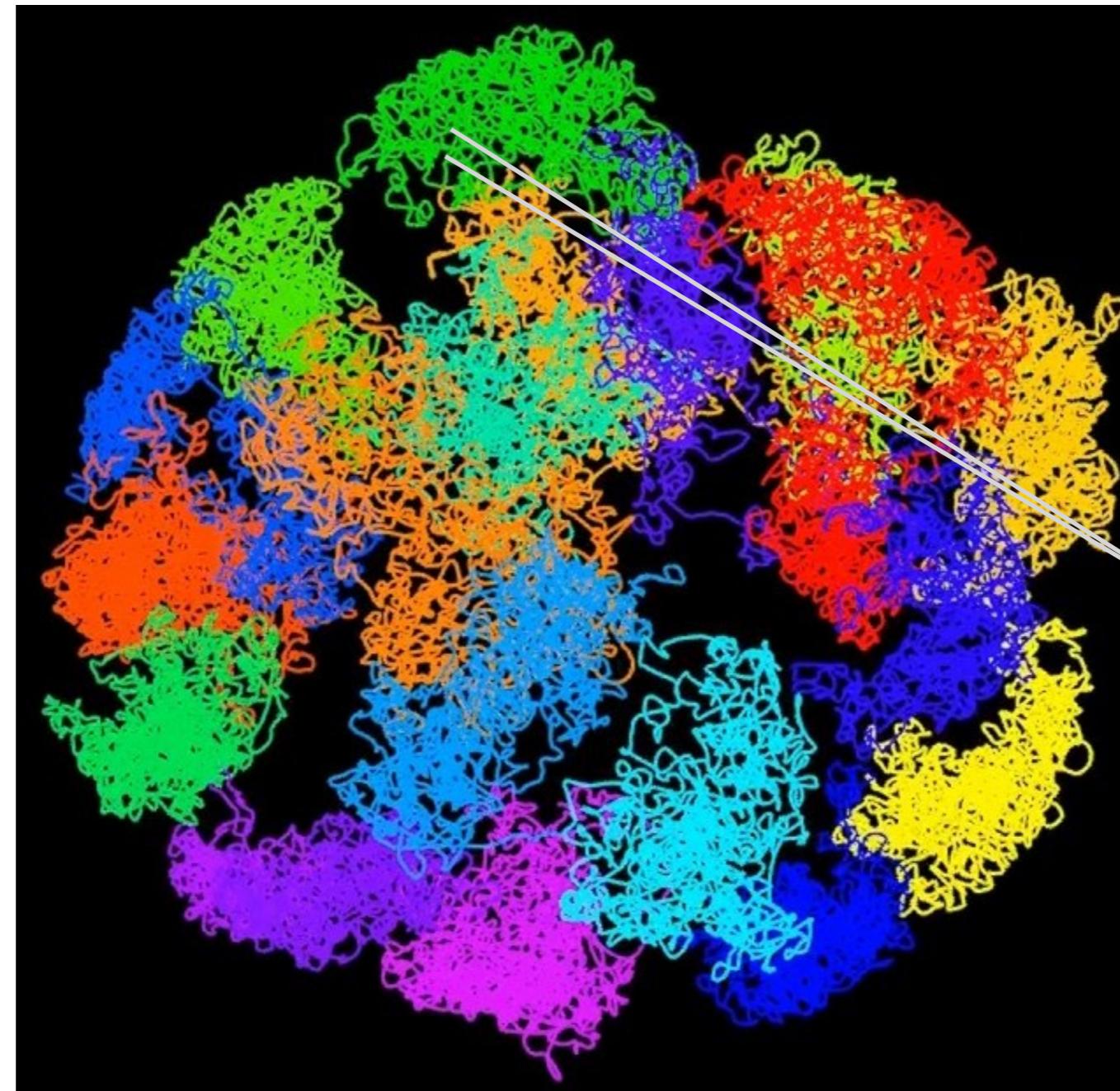


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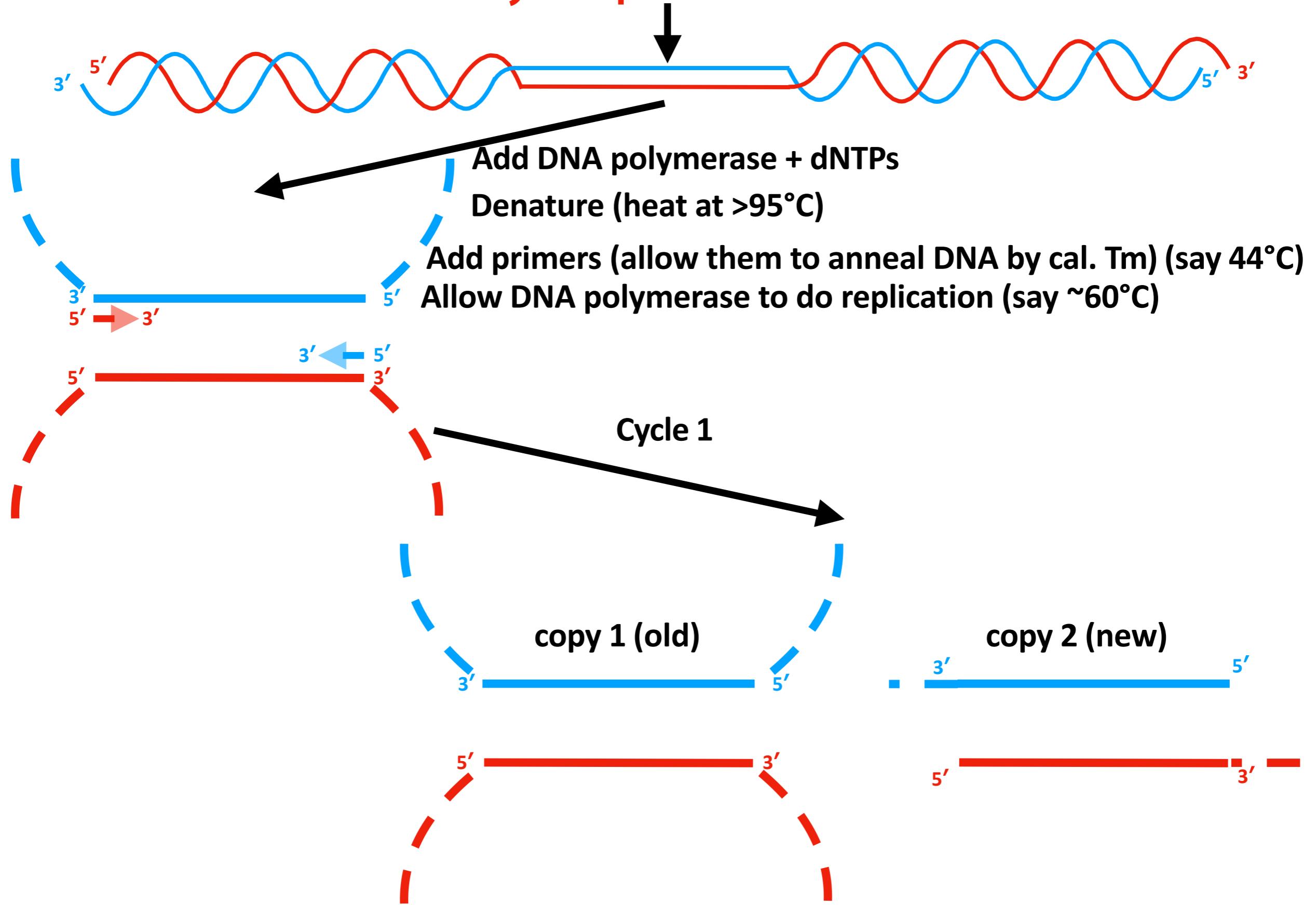
Micheal Smith & Kary
Mullis
Site-directed
mutagenesis & PCR

The human genome in 3D

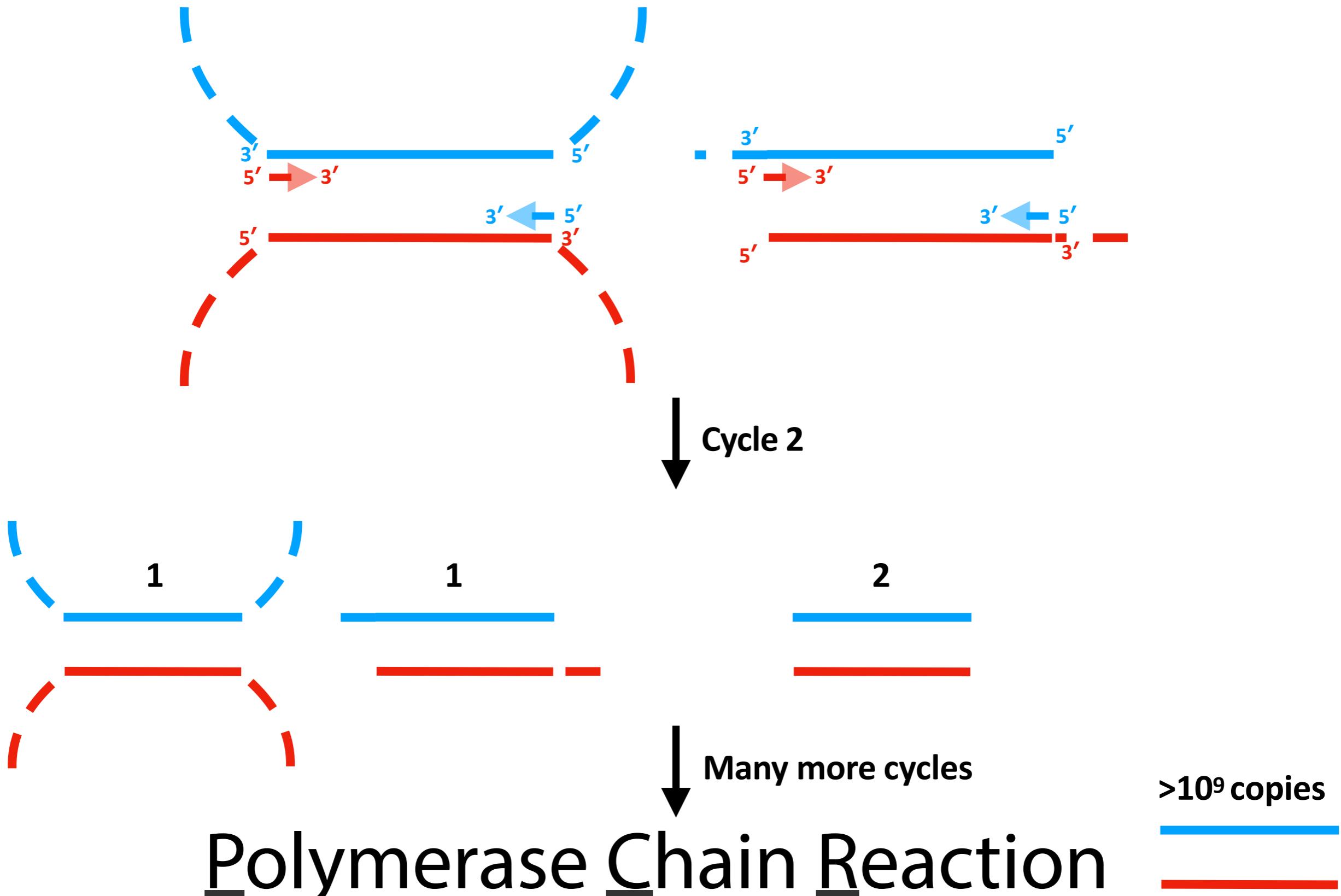


How to make a copy of this region in our genome?

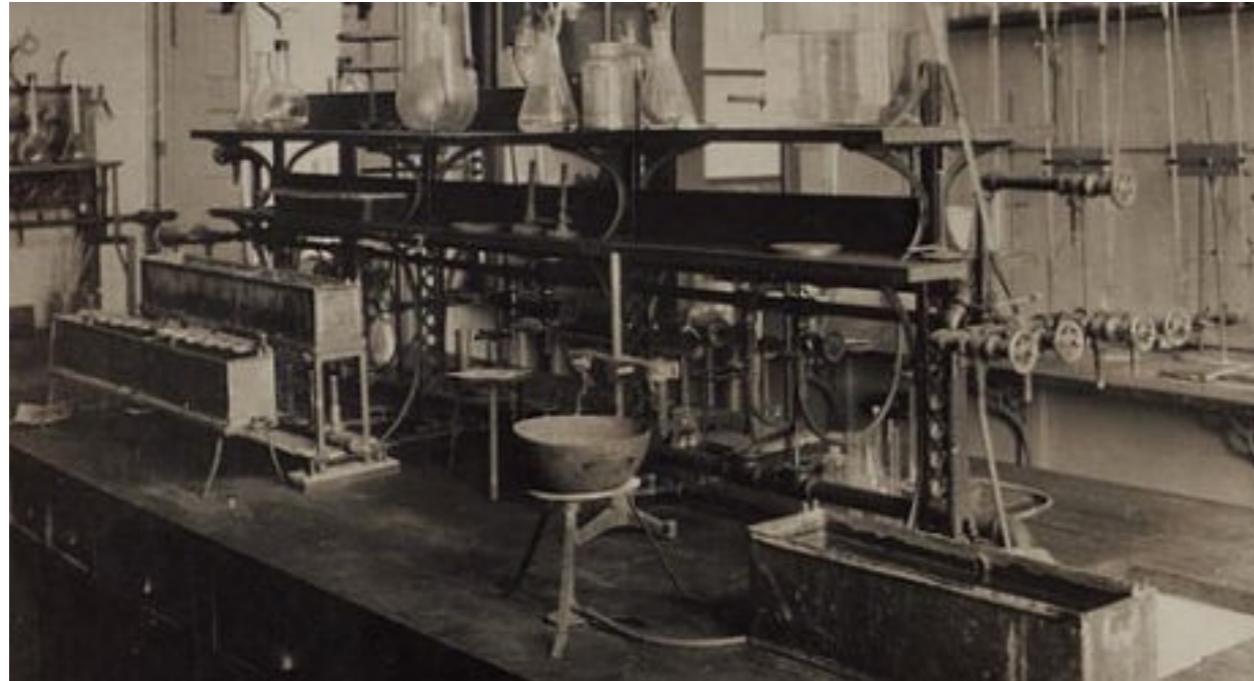
How to make many copies of DNA in a test tube?



Polymerase Chain Reaction



PCR machines

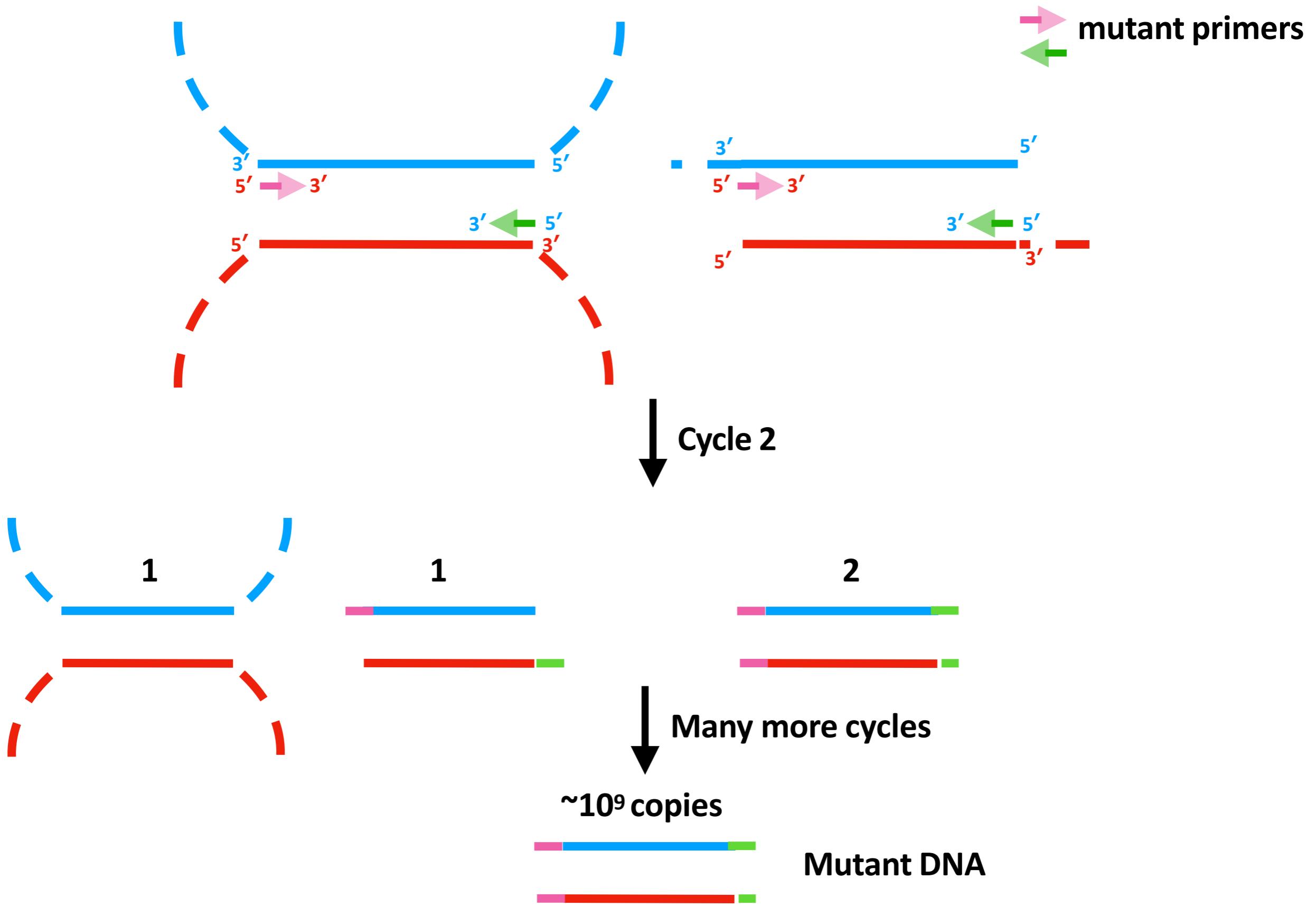


Running around the water baths (1980s)



And
now!!
!

Site-directed mutagenesis



Lecture #5

“DNA repair & recombination”

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DNA Damage

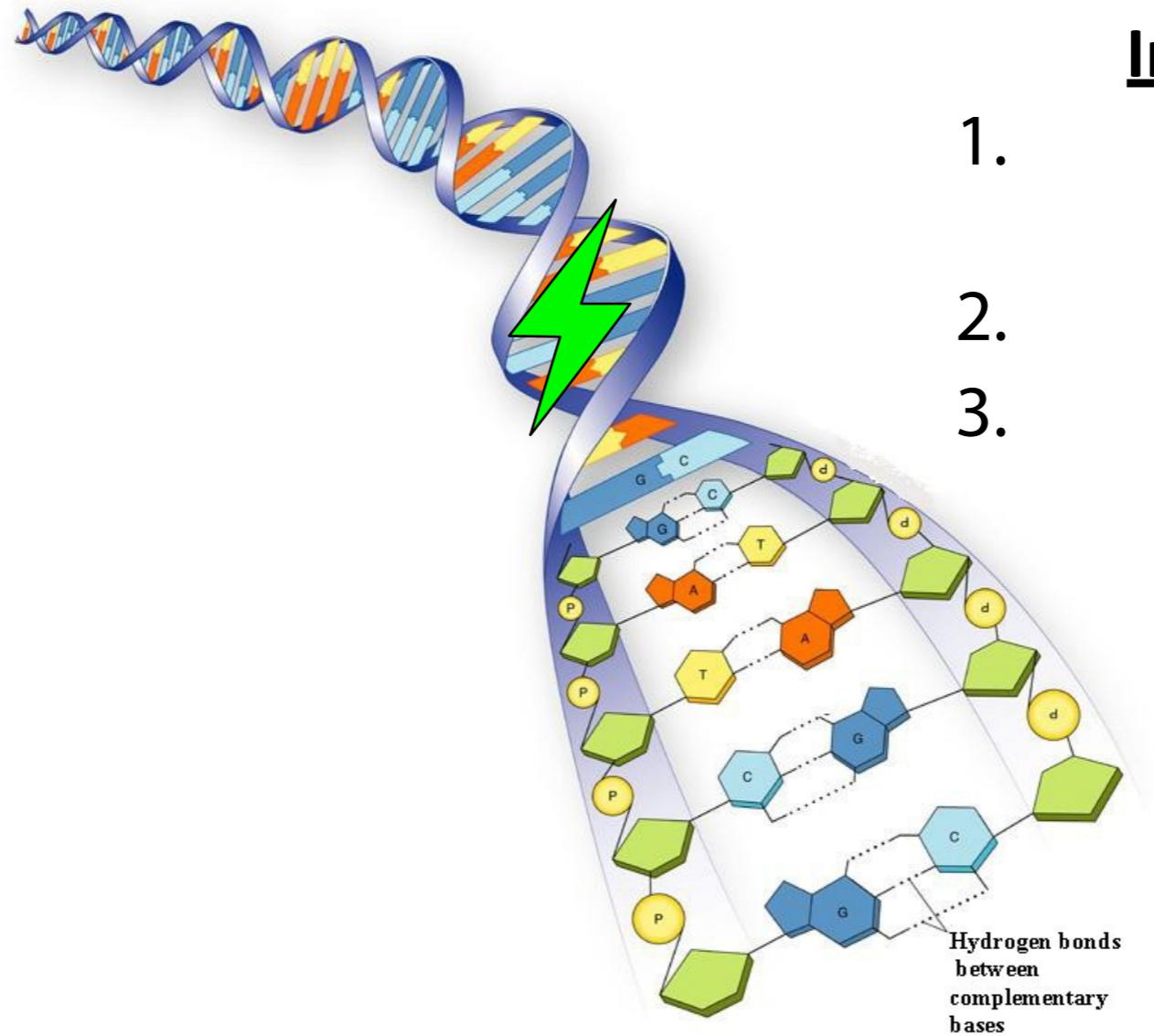
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External agents:

1. Sunlight
2. Tobacco smoke
3. Pollution
4. Toxins in food

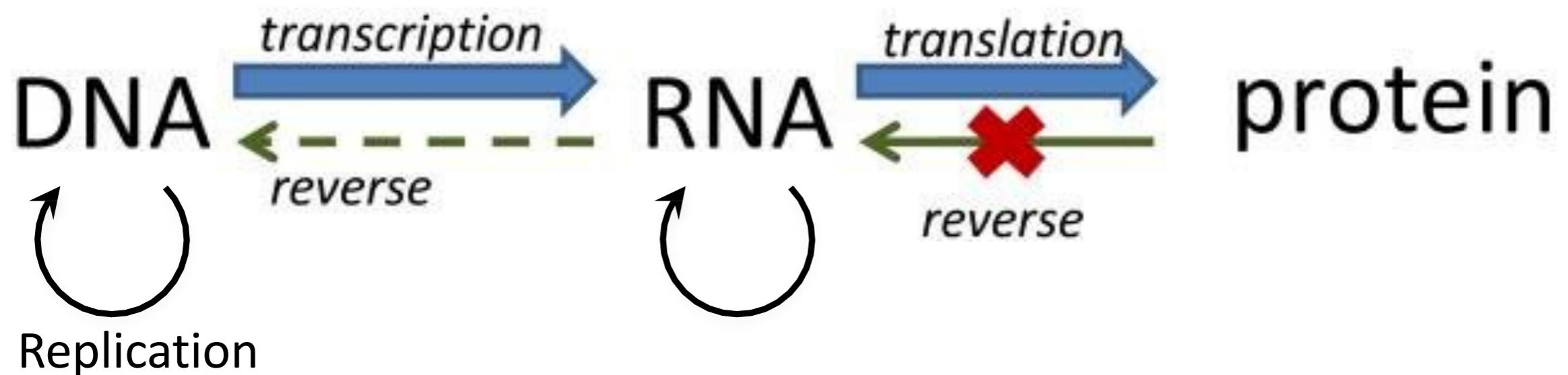
Internal agents:

1. Oxidation (ROS)
2. Reactive Nitrogen species - NO
3. Aging



DNA is constantly damaged by endogenous or exogenous agents

The central dogma



Damaging DNA results in a permanent change in the genetic material

DNA Damage is Bad

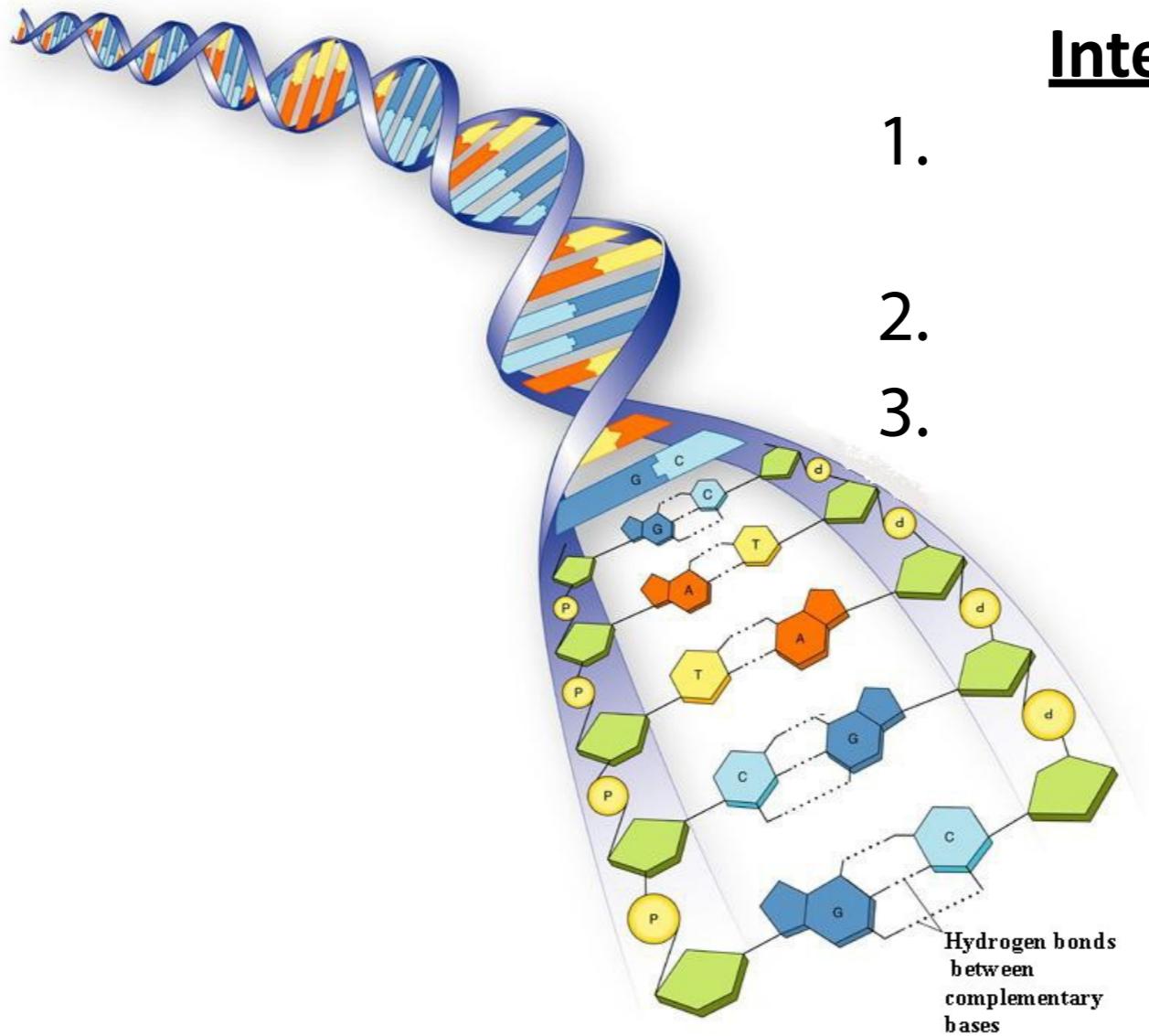
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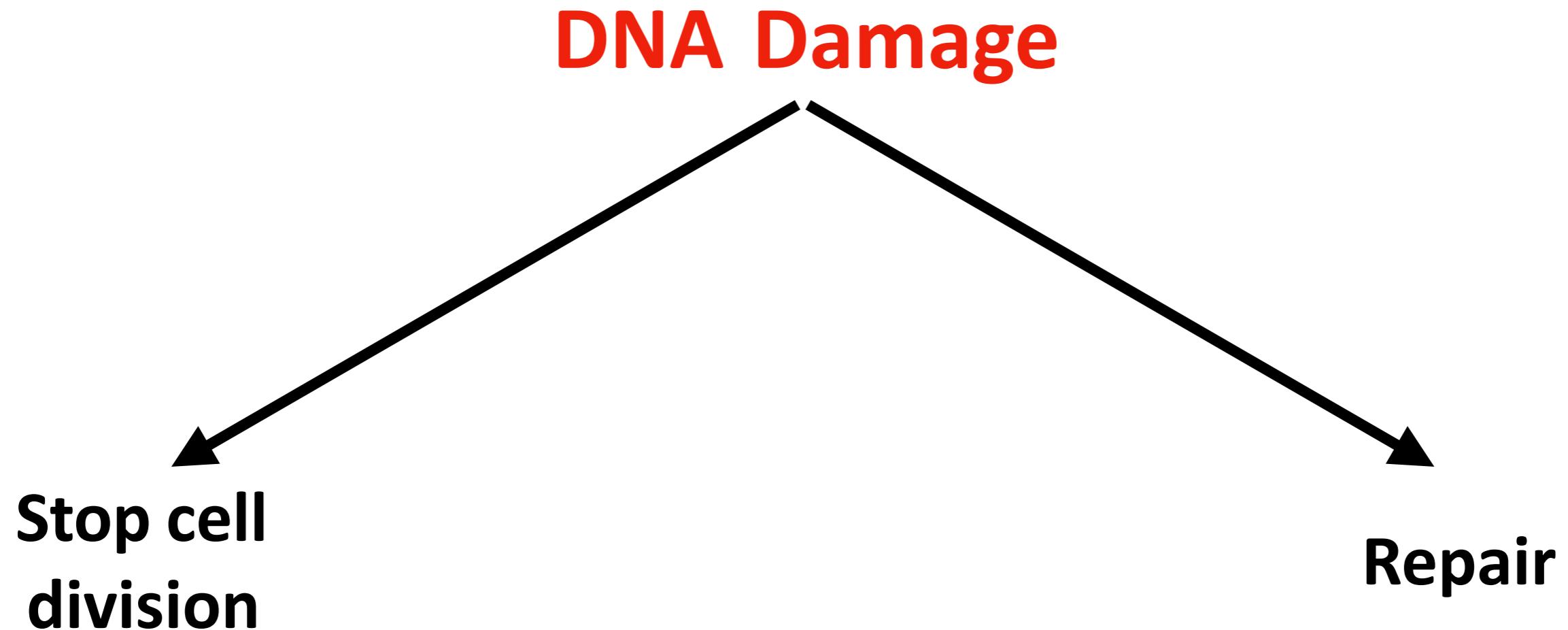


DNA damage causes mutations that may lead to Disease!!!

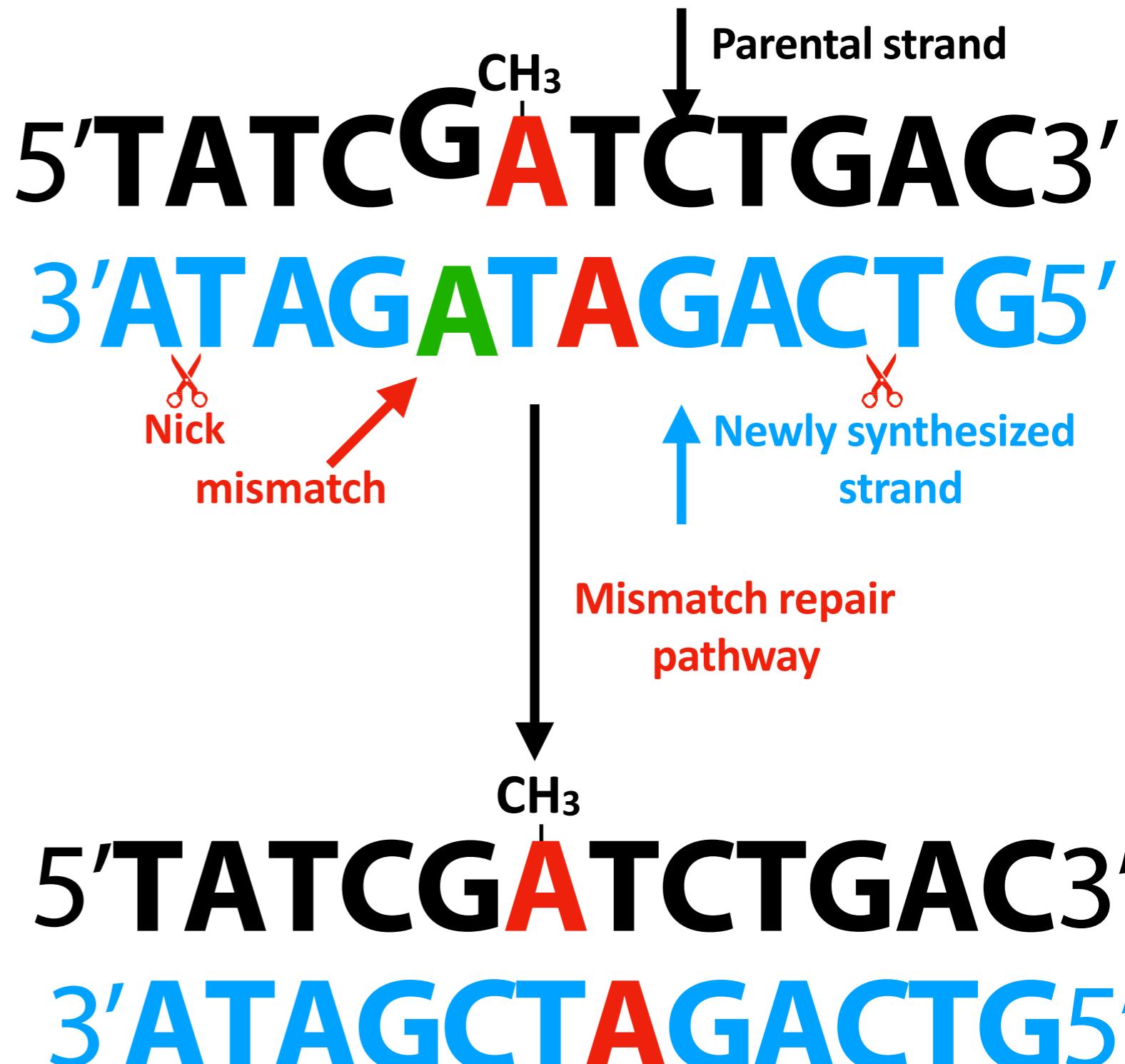
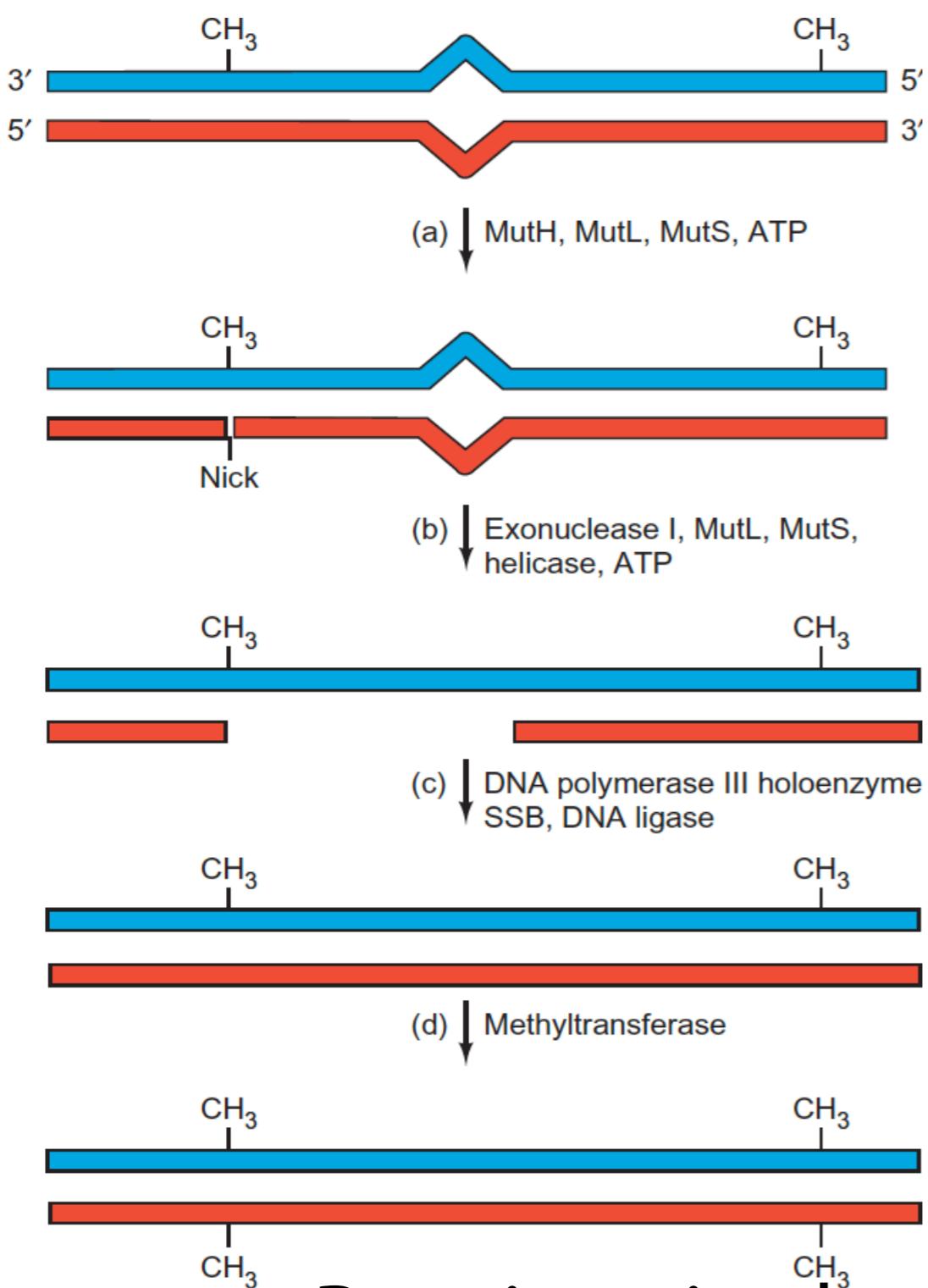
Mutation



What do cells do to repair DNA damage



Mismatch Repair

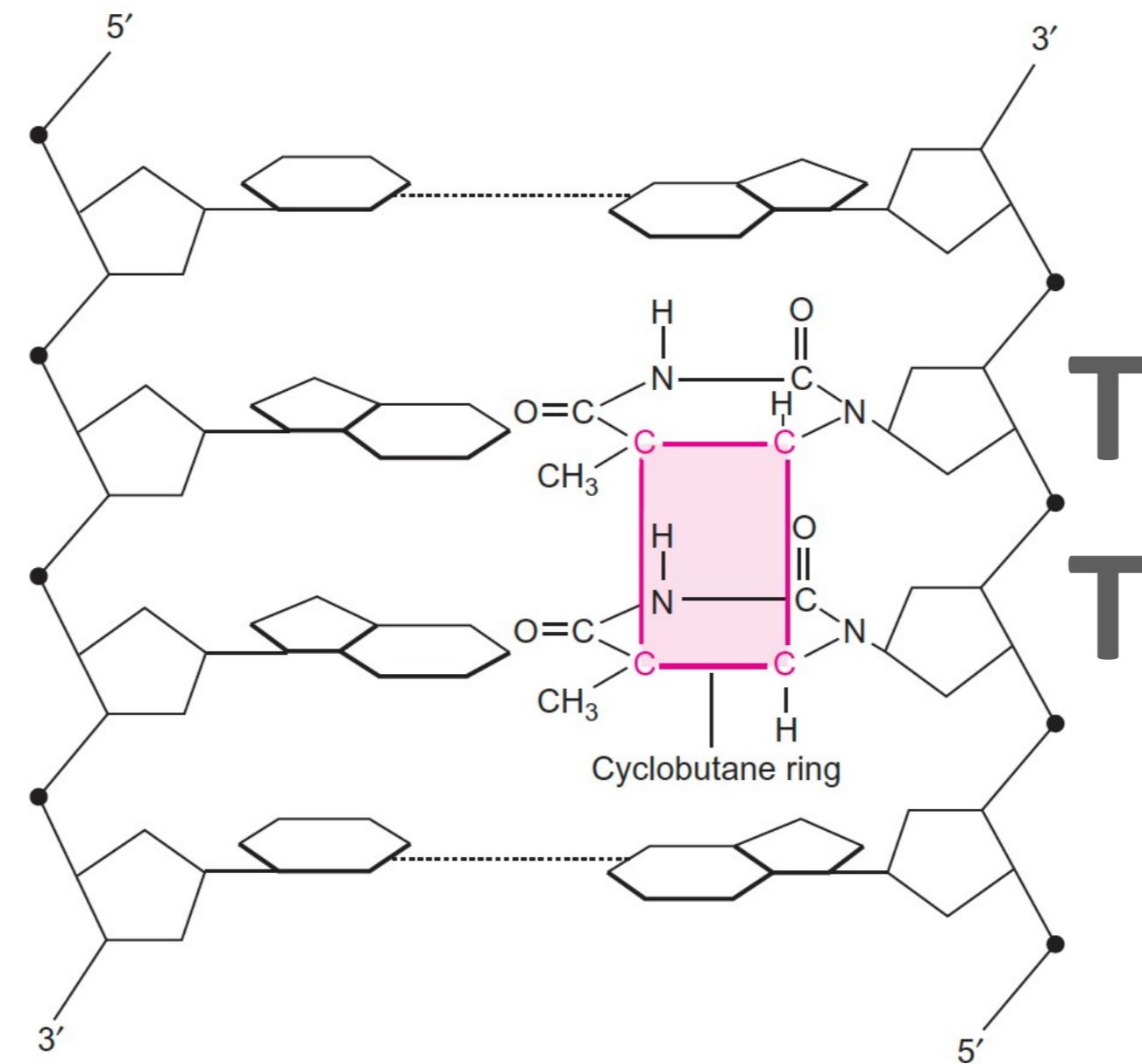
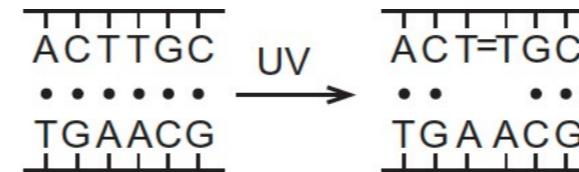


Repairs mistakes made by DNA polymerase

Mutation in mismatch repair causes Hereditary nonpolyposis colorectal cancer (HNPCC) or Lynch syndrome

Types of DNA damage

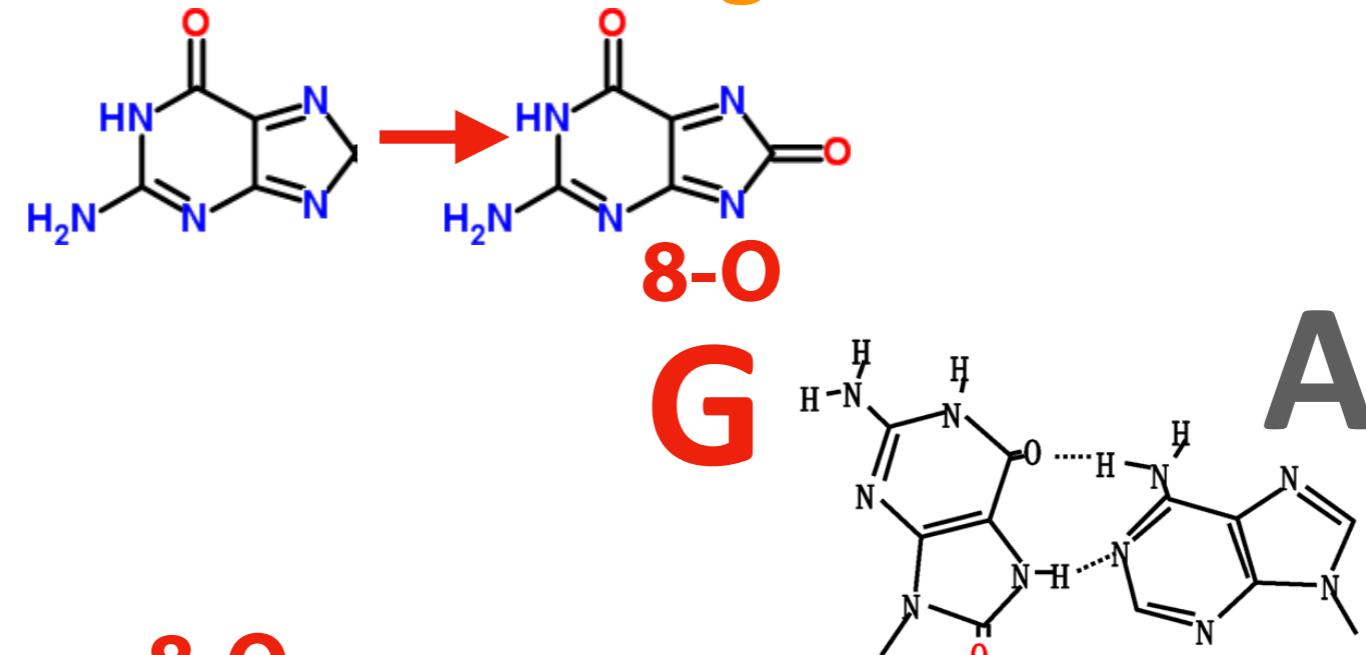
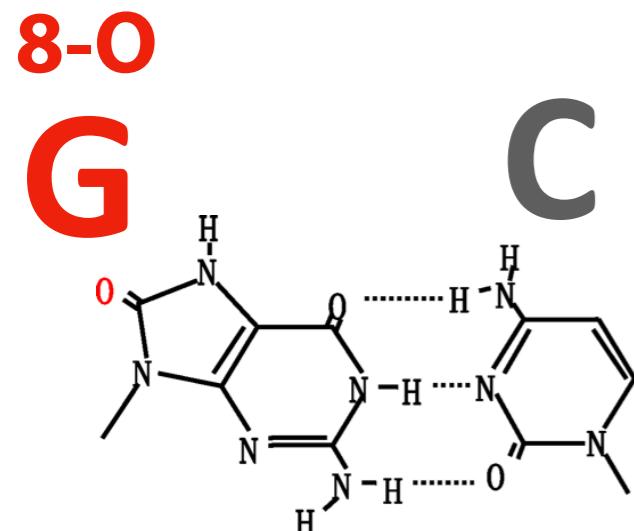
Ultra violet light



Cyclobutane Pyrimidine Dimer (CPD) formation

Types of DNA damage

Oxidative damage



TCG^{8-O}ATCT 3'

AGATAGA5'

TCGATCT 3'
AGCTAGA5'

GC to AT
Transition

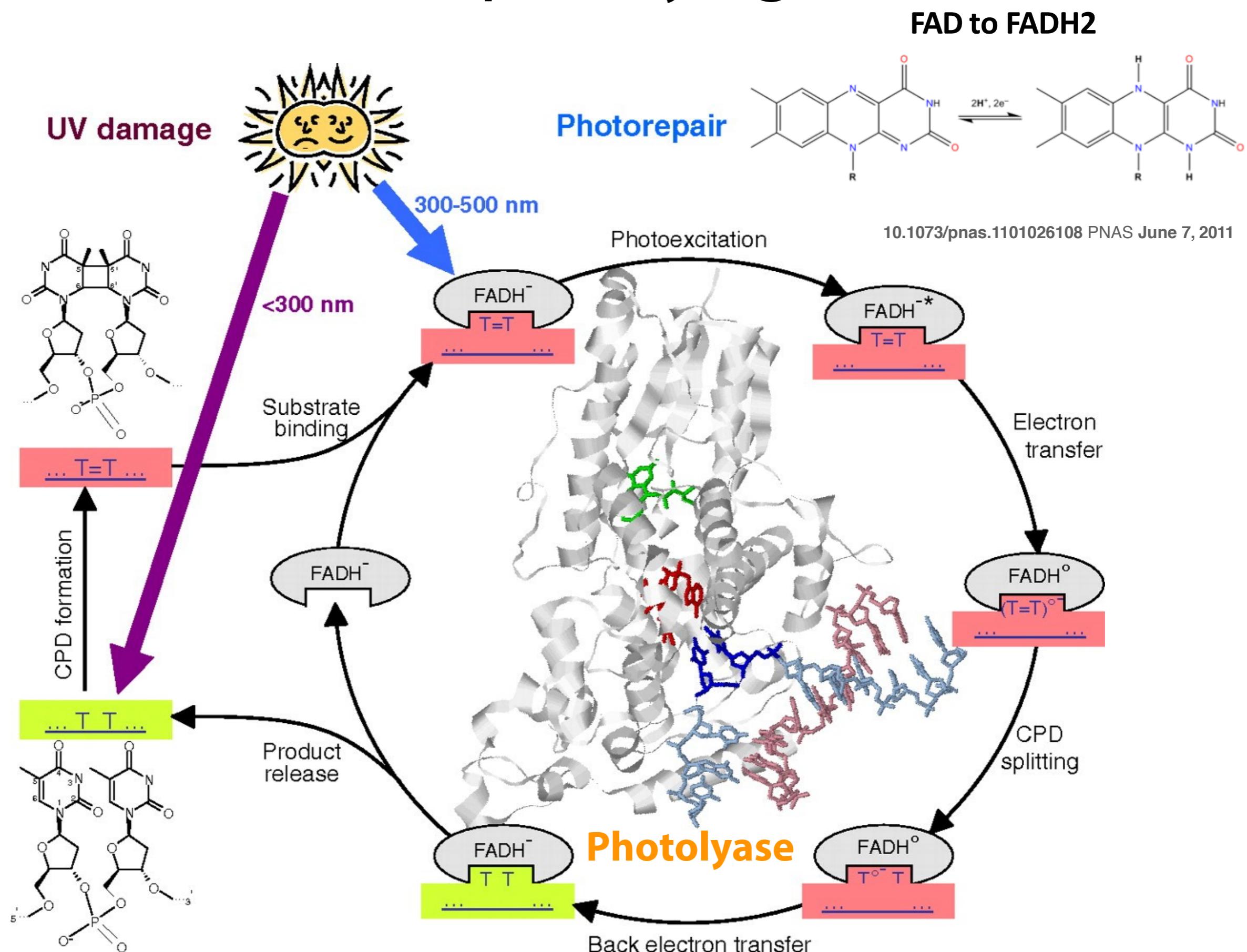
TC TATCT 3'
AGATAGA5'

Types of DNA damage

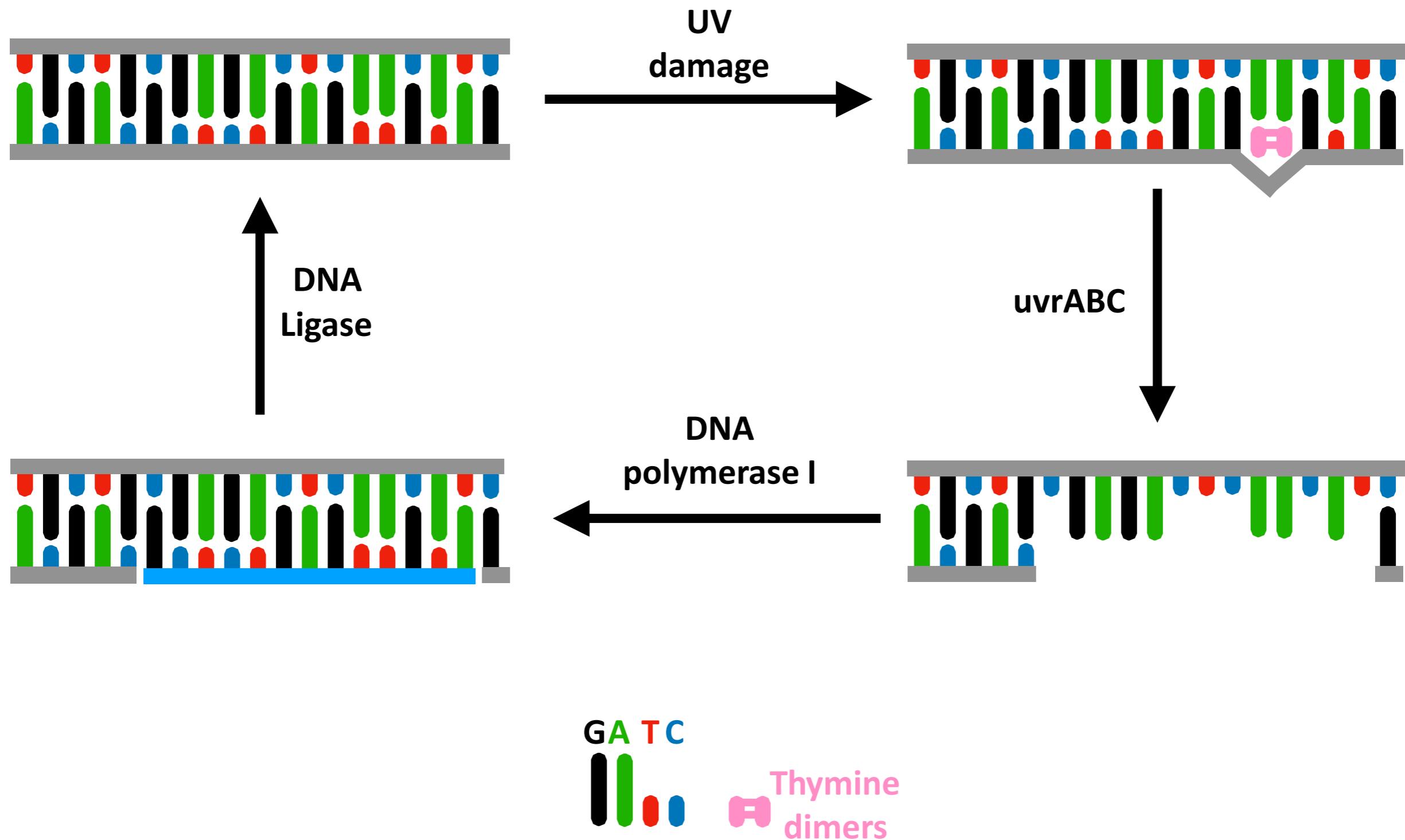
Table 1. DNA Lesions Generated by Endogenous and Exogenous DNA Damage

Endogenous DNA Damage	DNA Lesions Generated	Number Lesions/Cell/Day
Depurination	AP site	10000 ^a
Cytosine deamination	Base transition	100–500 ^a
SAM-induced methylation	3meA	600 ^a
	7meG	4000 ^a
	O ⁶ meG	10–30 ^b
Oxidation	8oxoG	400–1500 ^c
Exogenous DNA Damage	Dose Exposure (mSv)	DNA Lesions Generated
Peak hr sunlight	—	Pyrimidine dimers, (6–4) photoproducts
Cigarette smoke	—	aromatic DNA adducts
Chest X-rays	0.02 ^{f,g,h}	DSBs
Dental X-rays	0.005 ^{f,g,h}	DSBs
Mammography	0.4 ^{f,g,h}	DSBs
Body CT	7 ^f	DSBs
Head CT	2 ^{f,g}	DSBs
Coronary angioplasty	22 ^h	DSBs
Tumor PET scan (¹⁸ F)	10 ^h	DSBs
¹³¹ I treatment	70–150 ^h	DSBs
External beam therapy	1800–2000 ^j	DSBs
Airline travel	0.005/hr ^f	DSBs
Space mission (60 days)	50 ^k	DSBs
Chernobyl accident	300 ^l	DSBs
Hiroshima and Nagasaki atomic bombs	5–4000 ^k	DSBs

DNA repair by light



Nucleotide excision repair



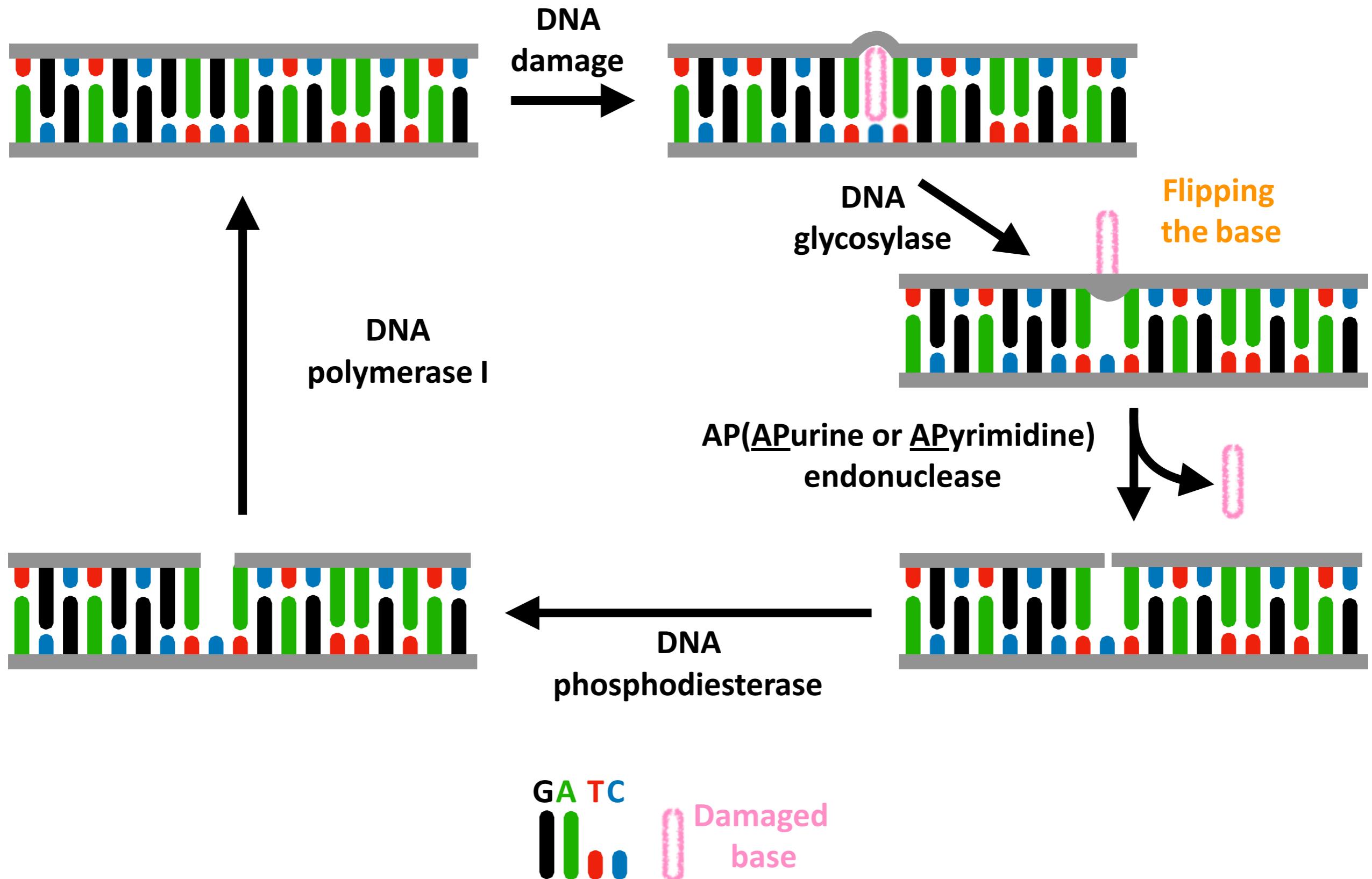
Inability to repair DNA damage causes disease



Xeroderma pigmentosum (XP)
mutation in DNA repair prevents skin cells
to repair damage by ultra violet light.

Inability to repair the DNA damage is
bad news

Base excision repair



Cytosine deamination

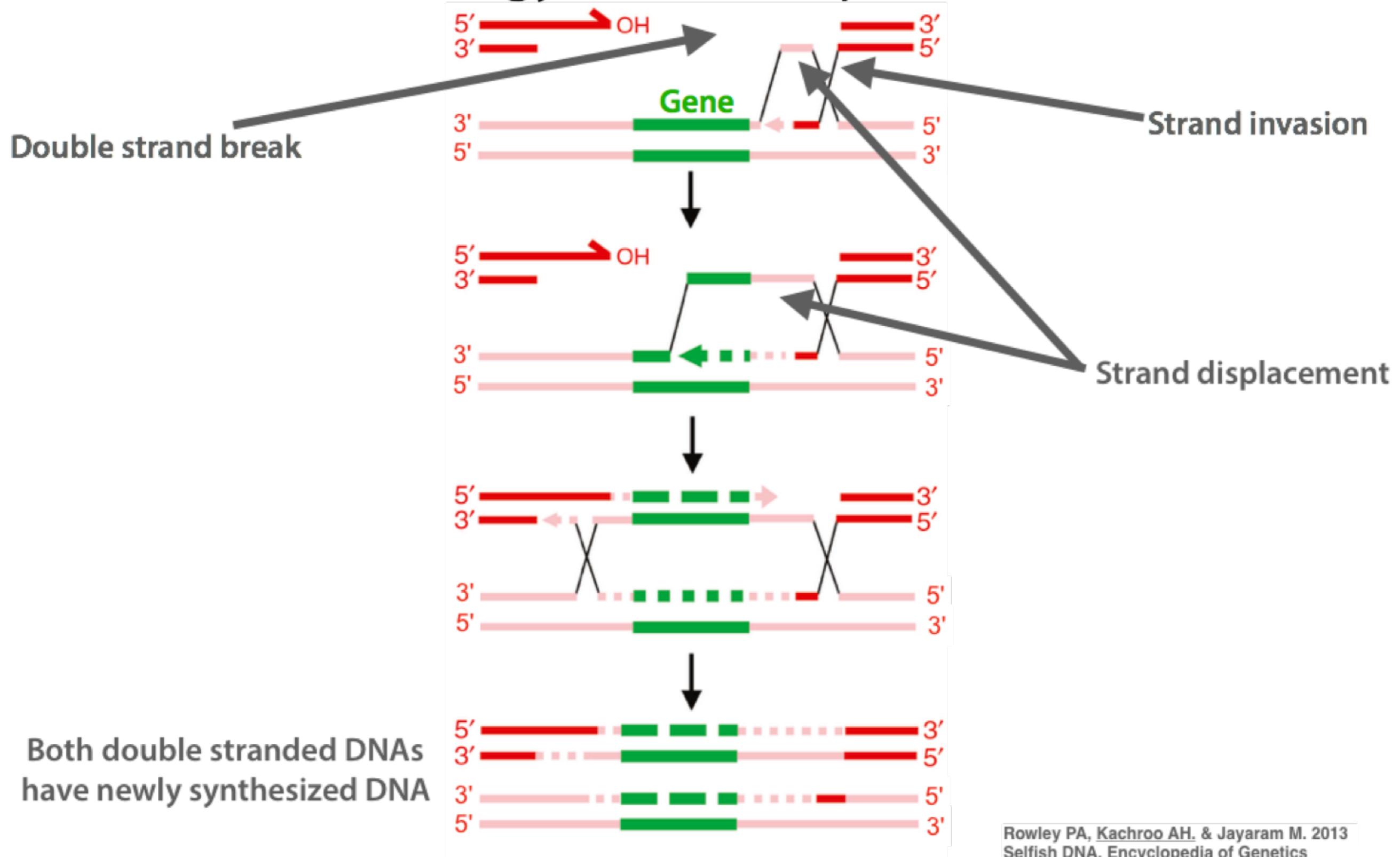


The diagram illustrates a sequence of DNA strands. The top row shows the strand: **T**
C **U** **A** **T** **C** **T** and the bottom row shows the strand: **G**
A → **A** **G** **T** **T** **A** **G** **A**. An orange arrow labeled "Cytosine deamination" points to the U in the first strand. A red arrow labeled "Mutation" points to the A in the second strand. An orange label "Replication without repair" is positioned between the two strands.

TCUATCT^{BER}TC ATCT → TCCATCT
AGGTAGA → AGGTAGA → AGGTAGA

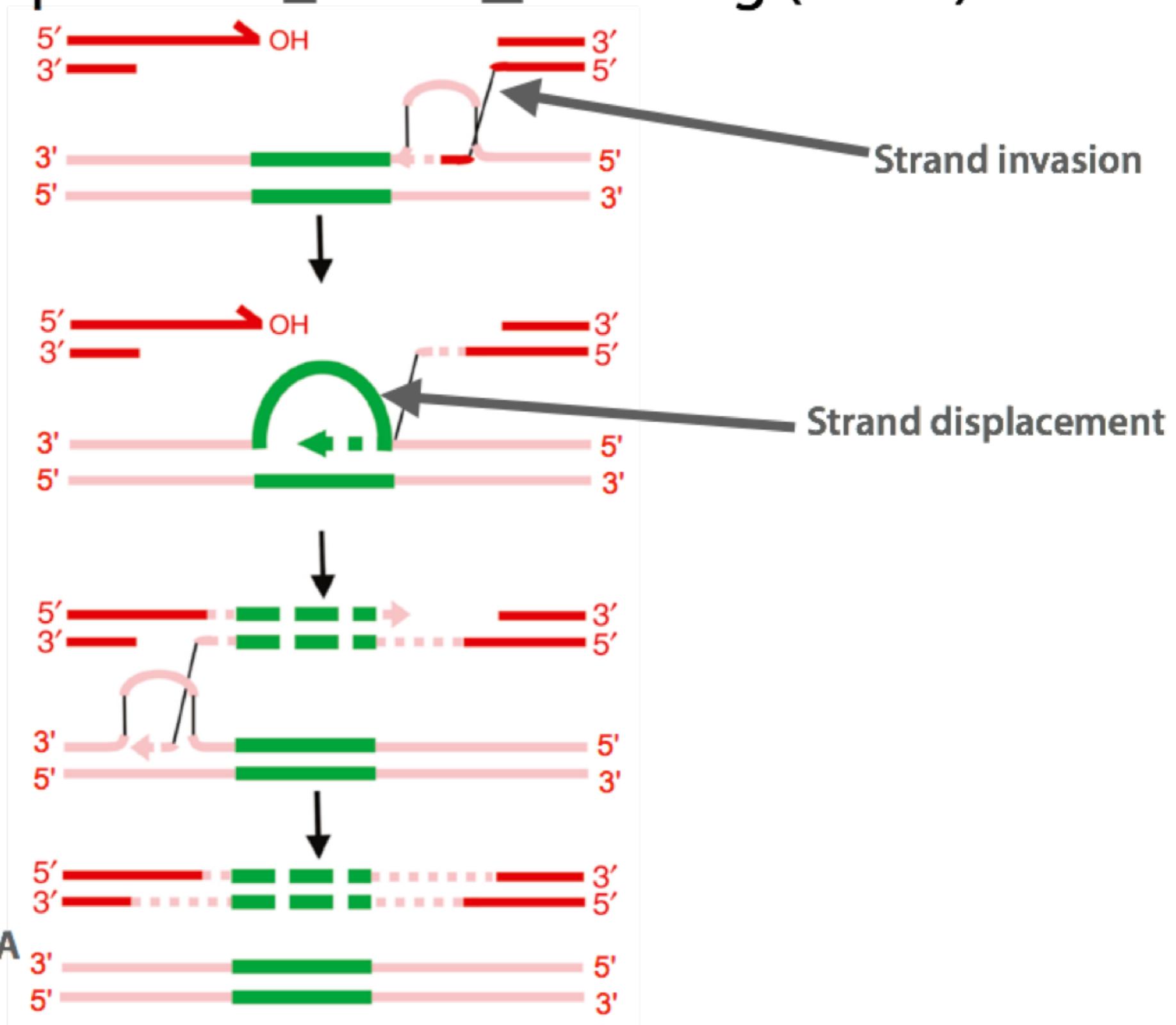
Double Strand Break Repair (DSBR)

Homology Directed Repair (HDR)



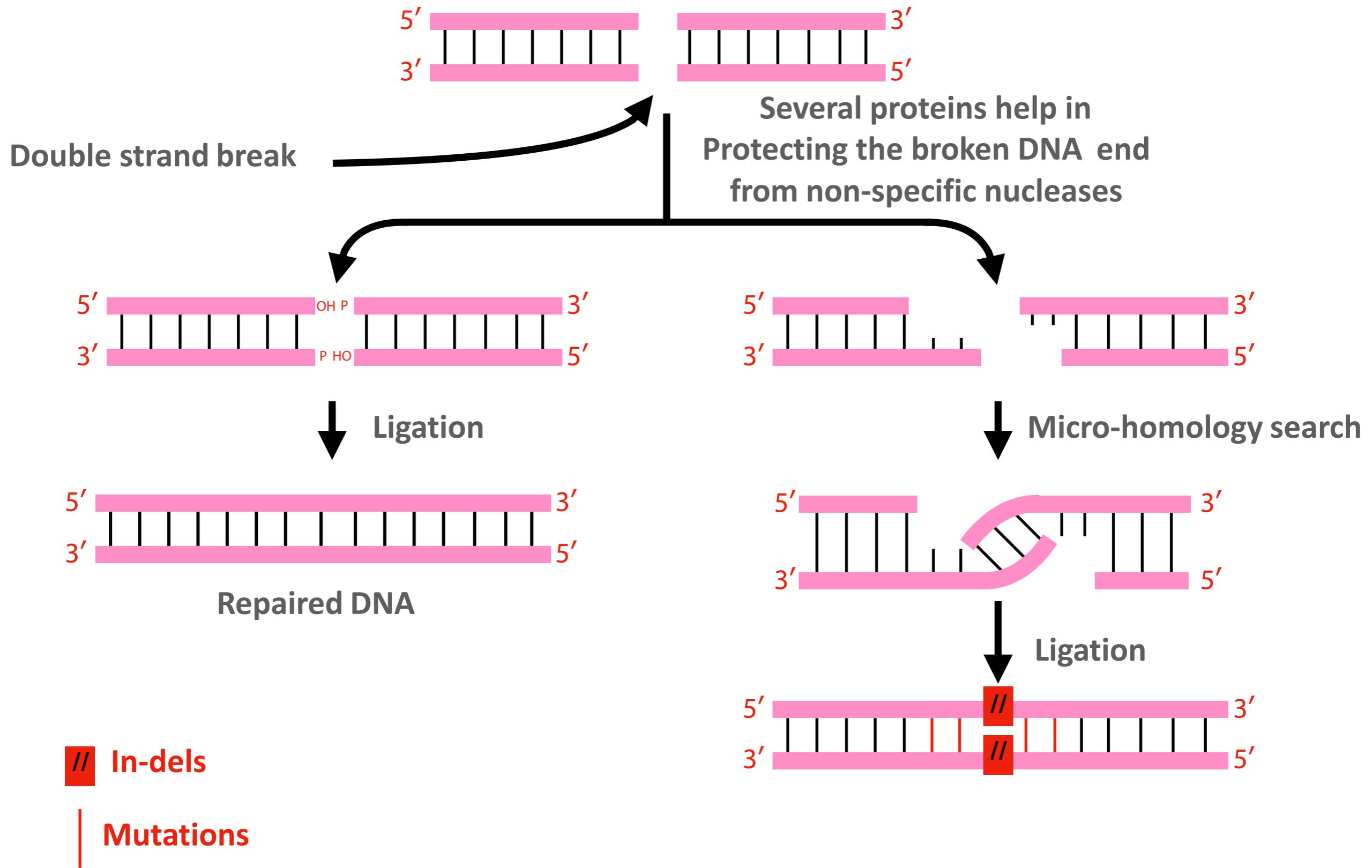
Double Strand Break Repair (DSBR)

Synthesis Dependent Strand Annealing (SDSA)



Double Strand Break Repair (DSBR)

Non Homologous End Joining (NHEJ)



“DNA repair movie”