

BIO-373
Genetics & Genomics

**Mutation
and DNA repair**

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Plan

1. Classification of mutations
 2. Spontaneous mutations
 3. Induced mutations
 4. DNA repair systems
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Mutations

- **Mutation = new alteration in DNA sequence**
 - Only way to introduce new genetic variation in the gene pool of a species
 - May occur anywhere in the genome, but frequency is increased in some regions called “mutational hotspots”
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1. Classification of mutations

Classification of mutations

- **Based on molecular change**

- **Point mutation or base substitution**

- **Missense mutation:** Results in a new triplet coding for a different amino acid
 - **Nonsense mutation:** Results in a new stop codon (translation terminated prematurely) – *TAG, TAA, TGA*
 - **Silent mutation:** Results in a new triplet coding for the same amino acid

- **Insertions / deletions (indels)**

- **In-frame indels:** respect the frame of triplet reading during translation
 - **Frameshift indels:** shift the frame → usually results in premature stop due to a new stop codon

- **Large structural variants**

- **Deletions, duplications, inversions, translocations...**
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Classification of mutations

- **Based on location**

- **Somatic mutations** occur in any cell except germ cells; are not heritable
 - **Germline mutations** occur in gametes; can be transmitted to the next generation
 - **Autosomal**
 - **X-linked and Y-linked**
 - **Mitochondrial**
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Classification of mutations

- **Based on detectable impact**
 - Neutral
 - Loss-of-function
 - Gain-of-function
 - Visible (morphological)
 - Nutritional (biochemical)
 - Behavioral
 - Regulatory
 - Lethal
 - Conditional/temperature-sensitive
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Classification of mutations

- **Based on mode of acquisition**
 - **Spontaneous mutations**
 - Arise naturally, from normal biological processes that alter the DNA bases
 - **Induced mutations**
 - Result from influence of extraneous factors, either natural or artificial (radiation, UV light, chemicals, etc.)
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2. Spontaneous mutations

Causes of spontaneous mutations

1. Replication errors

- Replication is imperfect
- DNA polymerase occasionally inserts incorrect nucleotides
- Repair mechanisms are not 100% effective

Two major types of replication errors:

- Replication slippage
 - Tautomeric shifts
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Causes of spontaneous mutations

- **Replication slippage**

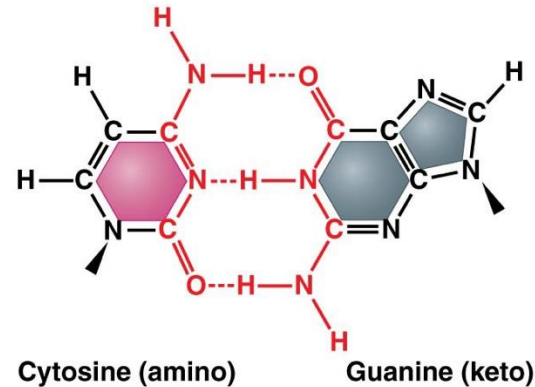
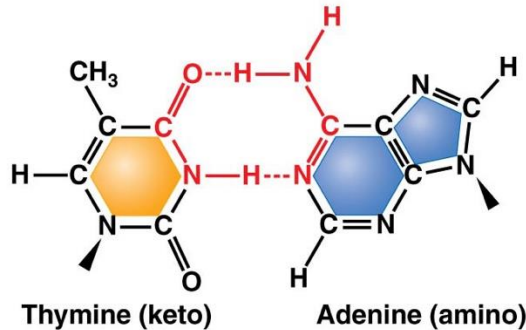
- Small insertions and deletions resulting from loops in the template strand during replication → DNA polymerase misses looped out nucleotides
 - More common in repeat sequences
 - Hot spots for DNA mutation
 - Known to cause genetic diseases
 - Fragile-X syndrome
 - Huntington disease
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Causes of spontaneous mutations

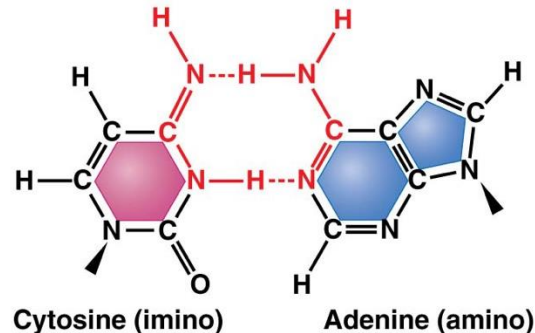
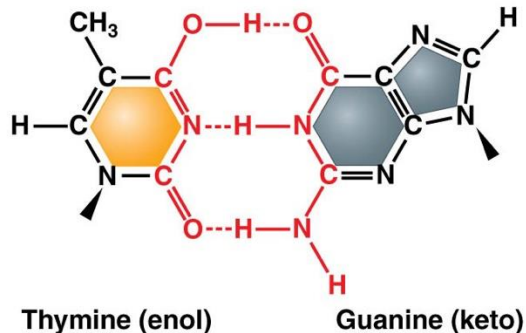
■ Tautomeric shifts

– **Tautomers:** alternate chemical forms of nucleotides

(a) Standard base-pairing arrangements



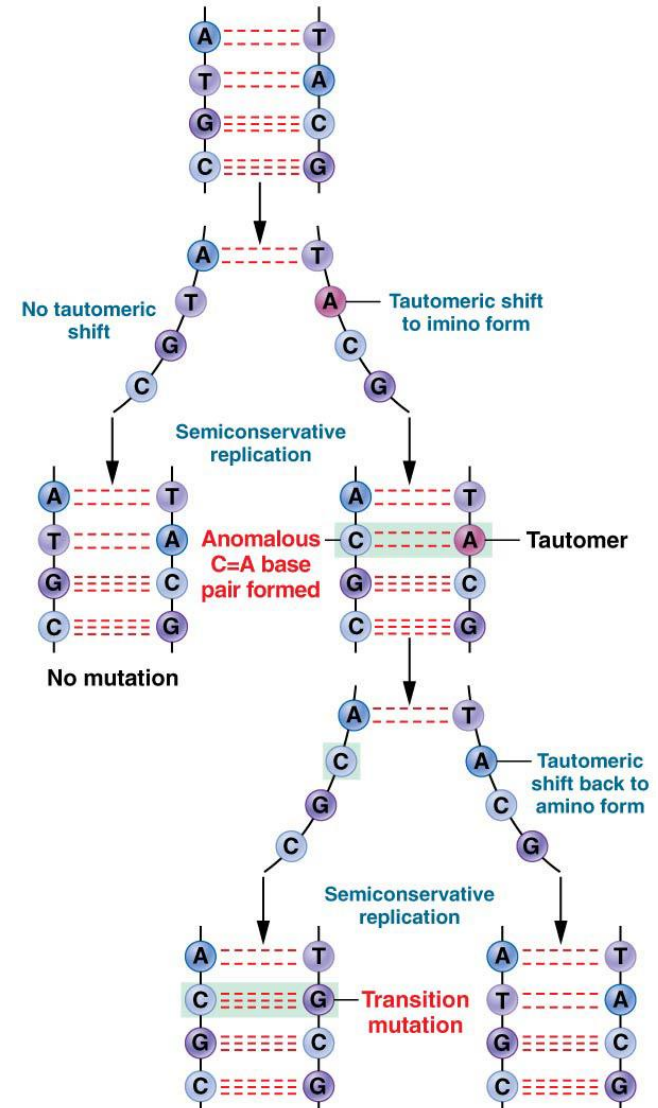
(b) Anomalous base-pairing arrangements



Causes of spontaneous mutations

■ Tautomeric shifts

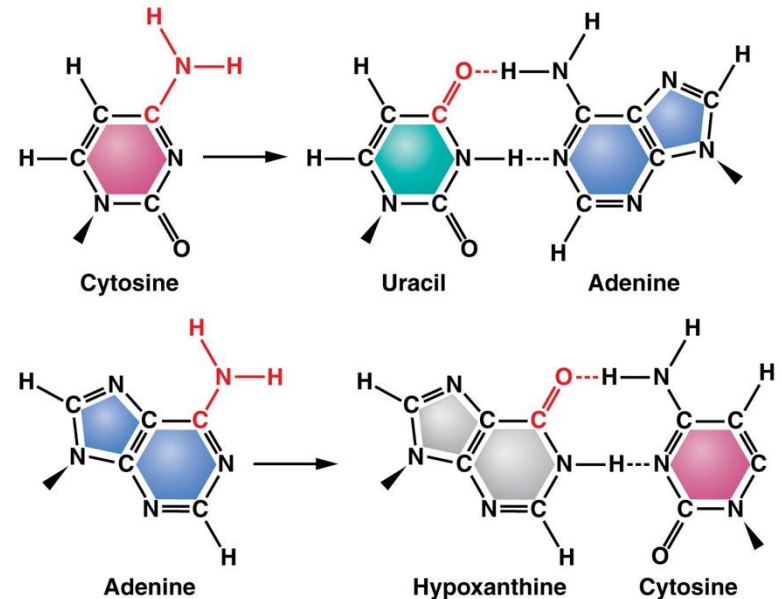
- A tautomeric shift changes the bonding structure, resulting in noncomplementary base pairing
- May lead to permanent base-pair changes and mutations



Causes of spontaneous mutations

2. DNA base damage

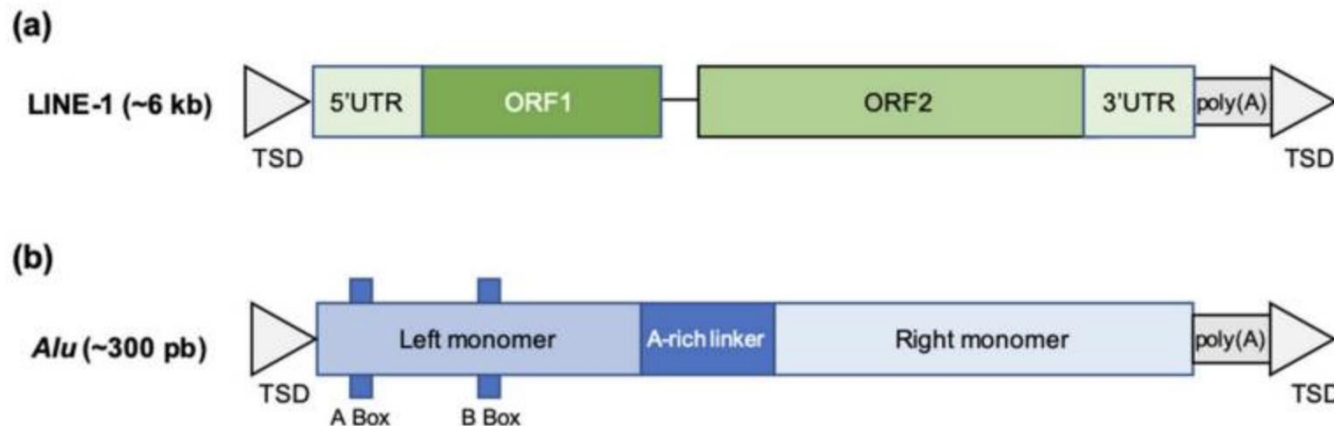
- **Depurination:** loss of a base, most often a purine (guanine or adenine)
- **Deamination:** the amino group in cytosine or adenine is converted to a keto group: cytosine is converted to uracil, and adenine to hypoxanthine



Causes of spontaneous mutations

3. Transposable elements

- Also called transposons or “jumping genes”
- DNA elements that move within or between genomes
- Can act as naturally occurring **mutagens**
- Examples in humans: *Alu* and LINE-1 repeats, which together make close to 30% of the genome (only a very small portion still active, i.e., capable of transposition)



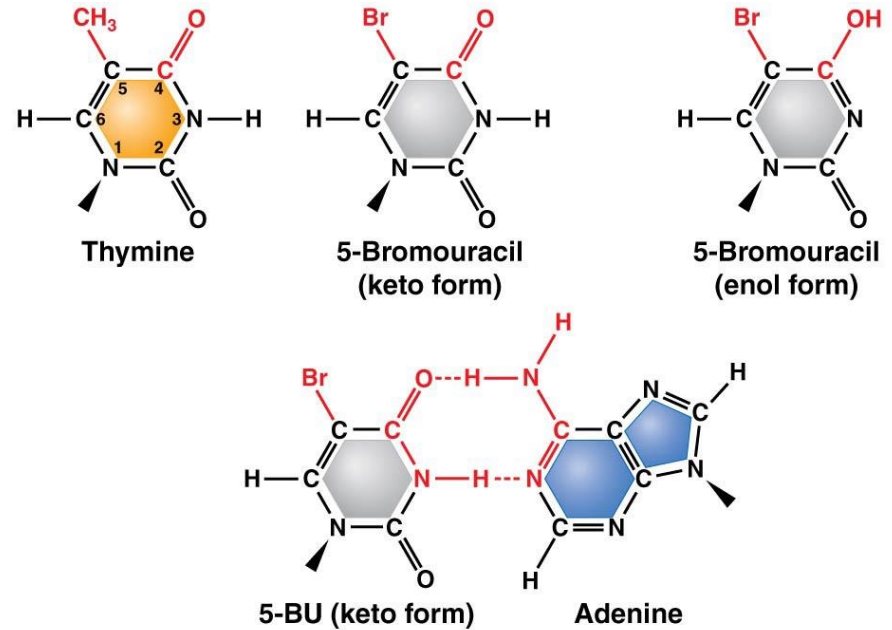
3. Induced mutations

Induced mutations

- Caused by **mutagens** = agents that induce mutations
 - Base analogs
 - Alkylating agents
 - Intercalating agents
 - Adduct-forming agents
 - UV light
 - Ionizing radiation
 - Free radicals
 - Viruses
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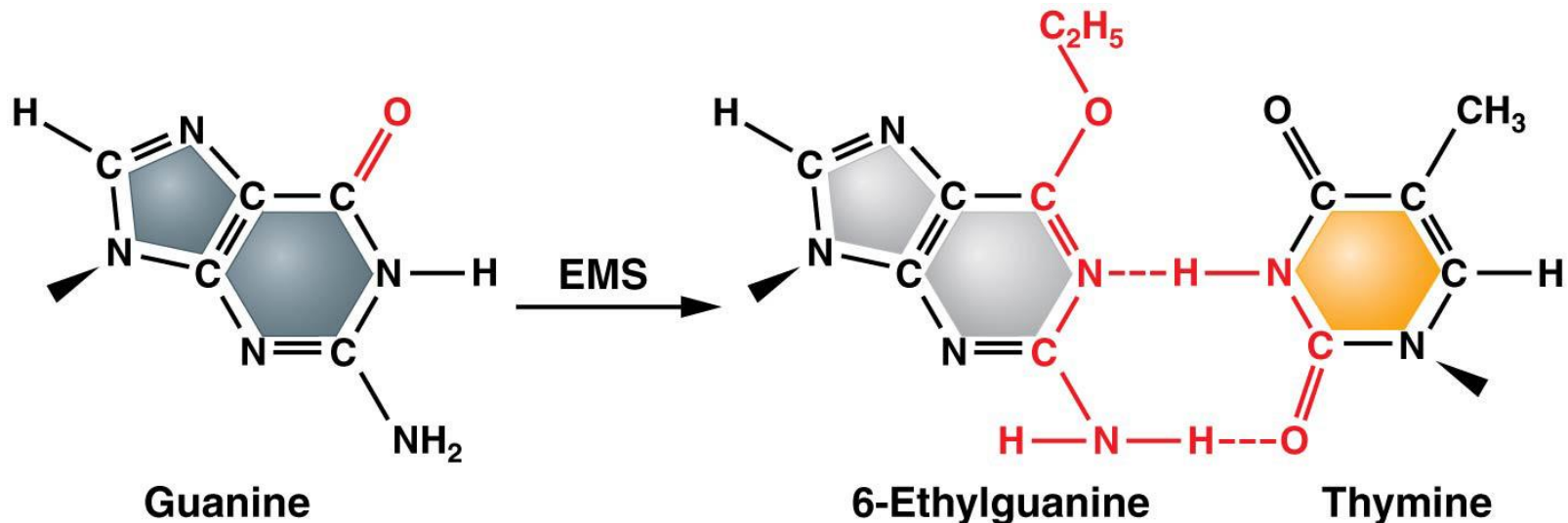
Base analogs

- Can substitute for purines or pyrimidines during nucleic acid biosynthesis
- Example: 5-Bromouracil behaves as thymine analog



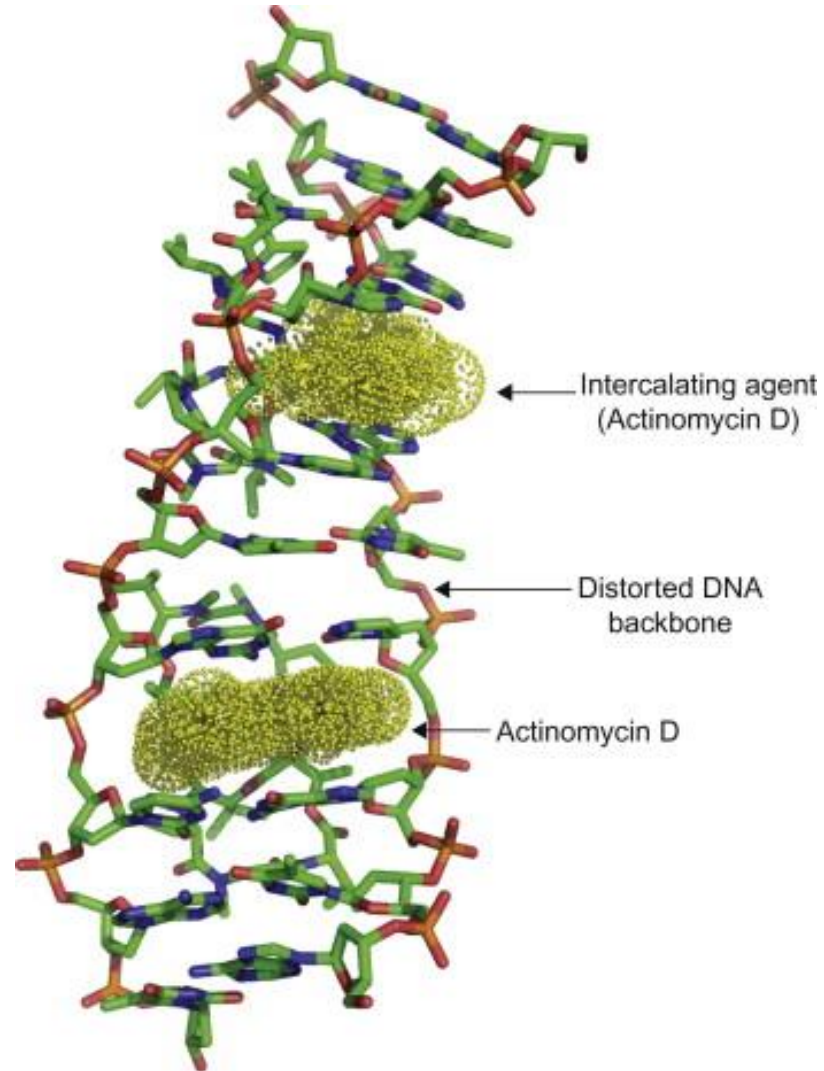
Alkylating agents

- Donate alkyl group (CH_3 or CH_3CH_3) to amino or keto groups in nucleotides
- Alter base-pairing affinity
- Examples: Ethyl Methyl Sulfonate, mustard gas



Intercalating agents

- Chemicals with dimensions and shapes that wedge between DNA base pairs
- This causes base-pair distortions and DNA unwinding
- Example: Actinomycin D, Ethidium bromide

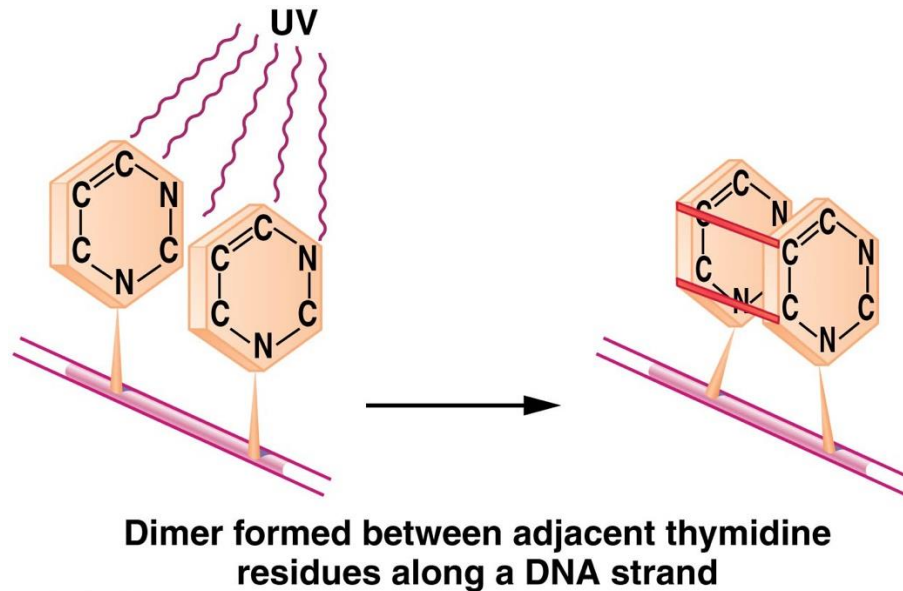


Adduct-forming agents

- Chemicals that covalently binds to DNA, altering conformation and interfering with replication and repair
 - A **DNA adduct** is a segment of DNA bound to a carcinogen
 - Examples:
 - Acetaldehyde (component of cigarette smoke)
 - Nitrosamines (used as meat preservatives)
 - Aflatoxin (produced by some *Aspergillus*)
 - Cisplatin (anti-cancer drug)
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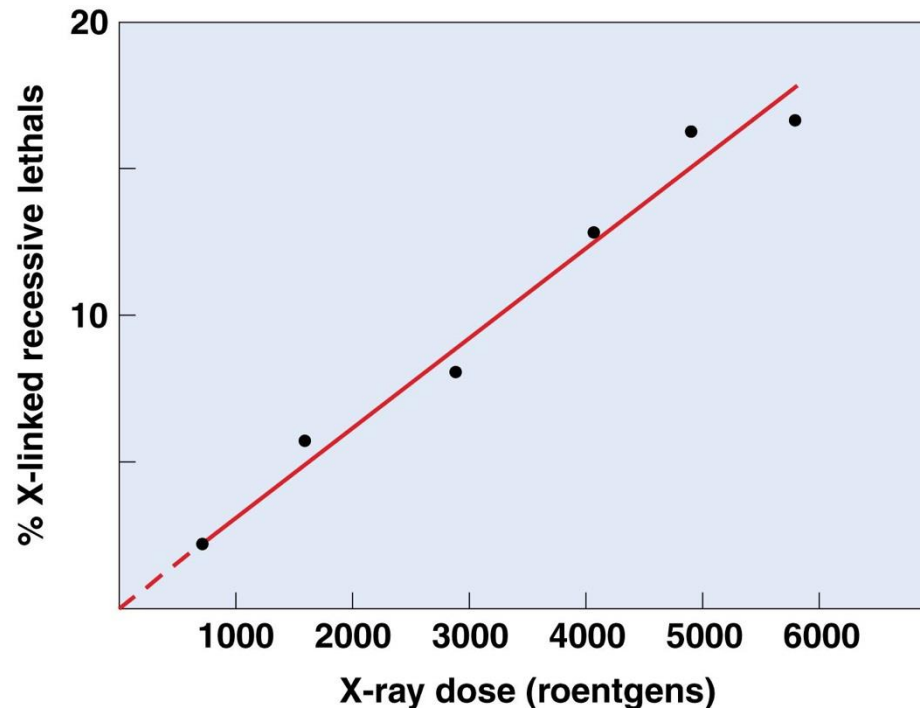
UV light

- UV light creates **pyrimidine dimers**
 - Two identical, adjacent pyrimidines create a dimer
 - Lead to distortion of DNA conformation
 - Errors are then introduced during DNA replication



Ionizing radiation

- X rays, gamma rays, cosmic rays
- Penetrates deeply into tissues
- Causes ionization of molecules



Free radicals

- By-products of biochemical reactions in the body, including some metabolic processes and immune system responses
 - Highly reactive because they contain one or more unpaired electrons
 - Free radicals can affect DNA directly or indirectly:
 - Alter purines and pyrimidines
 - Break phosphodiester bonds
 - Produce deletions, translocations, and DNA fragmentation
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4. DNA repair systems

DNA Repair

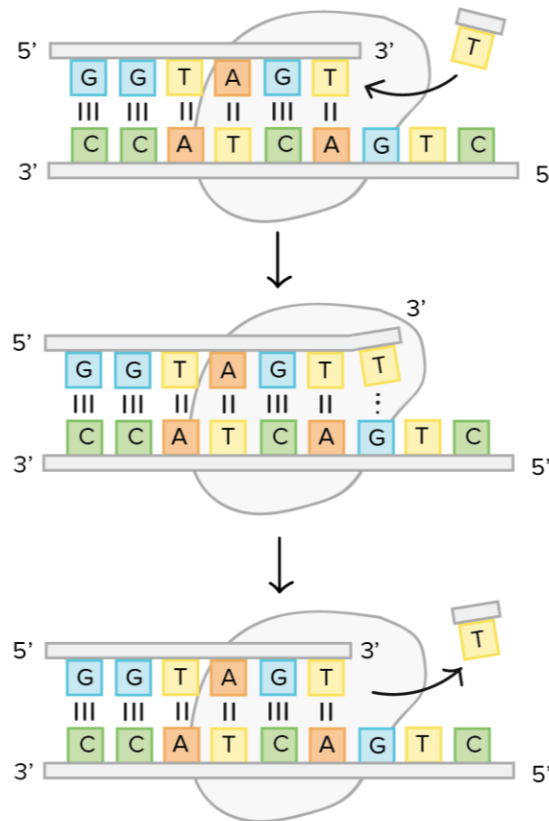
- **Repair systems counteract spontaneous and induced DNA damage**
 - Maintains integrity of genetic material
 - Counteract genetic damage that would result in genetic diseases and cancer
 - The balance between mutation and repair results in the **observed mutation rates** of individual genes and organisms.
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Systems of DNA repair

1. During DNA synthesis, most DNA polymerases "check their work," fixing the majority of mispaired bases in a process called **proofreading**
 2. Immediately after DNA synthesis, any remaining mispaired bases can be detected and replaced in a process called **mismatch repair**
 3. Outside of synthesis, if DNA gets damaged, it can be repaired by various mechanisms, including **chemical reversal**, **excision repair**, and **double-stranded break repair**
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Proofreading

- Happens during DNA synthesis
- When an incorrect base pair is recognized, the **DNA polymerase** reverses its direction and excises the mismatched base
- This implies that the polymerase has a **3' → 5' exonuclease activity**



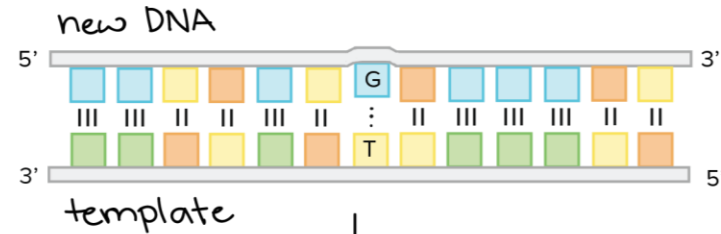
Polymerase adds an incorrect nucleotide to the new strand of DNA.

Polymerase detects that bases are mispaired.

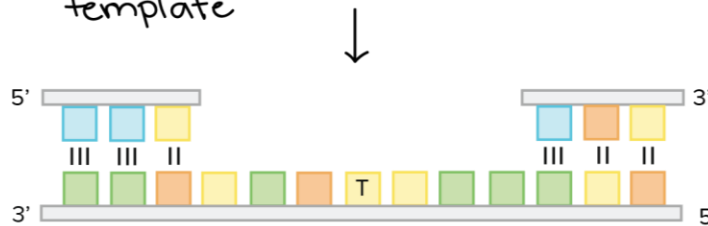
Polymerase uses 3' → 5' exonuclease activity to remove incorrect nucleotide.

Mismatch repair

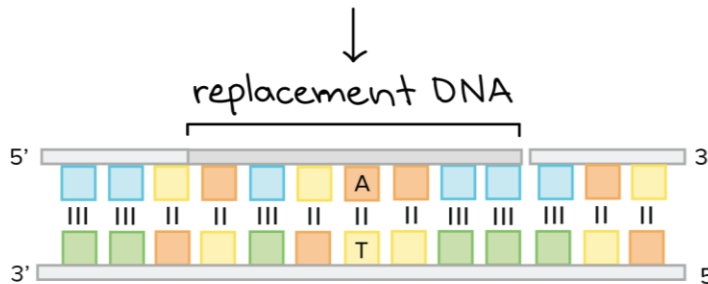
- Happens right after DNA synthesis
- Removes mismatches that were missed by proofreading
- Needs **nuclease**, **DNA polymerase** and **DNA ligase**



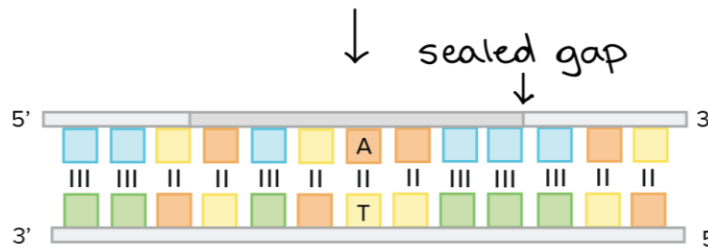
A mismatch is detected in newly synthesized DNA.



The new DNA strand is cut, and the mispaired nucleotide and its neighbors are removed.



The missing patch is replaced with correct nucleotides by a DNA polymerase.



A DNA ligase seals the gap in the DNA backbone.

DNA damage repair

1. Chemical reversal

- DNA damage is fixed by reversing the chemical reaction that caused it
 - Example: **photoreactivation** in bacteria: DNA photolyase (a light-driven enzyme) can destroy the abnormal covalent bond present in pyrimidine dimers induced by UV light
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DNA damage repair

2. Excision repair

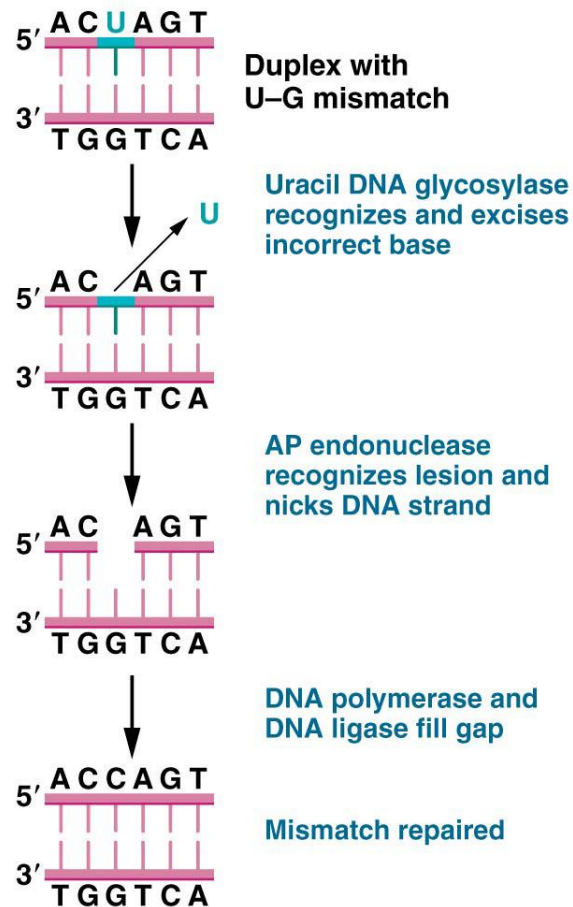
■ Base excision repair (BER)

- Corrects DNA containing a damaged DNA base
- A DNA glycosylase recognizes an altered base

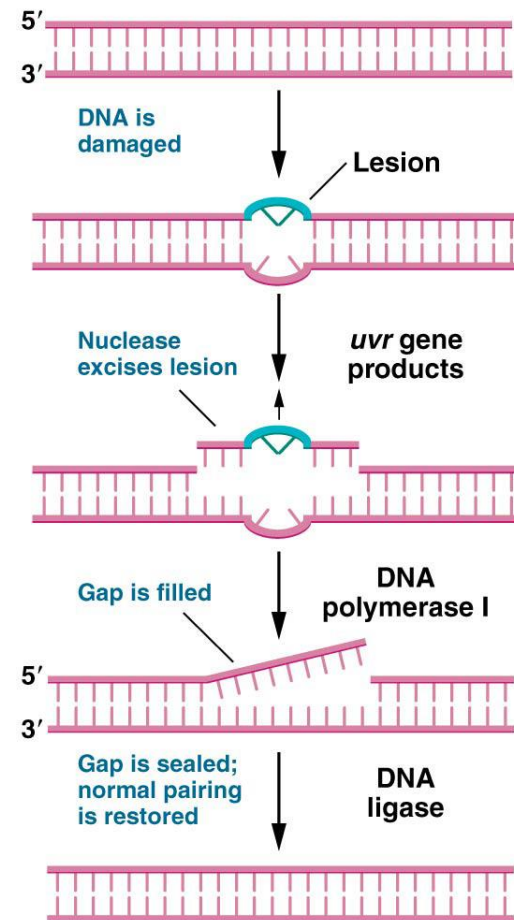
■ Nucleotide excision repair (NER)

- Repairs bulky lesions that alter/distort the DNA double helix

Base excision repair



Nucleotide excision repair



DNA damage repair

3. Double-stranded break repair

- Double strand breaks are extremely dangerous
 - Results in chromosomal rearrangements, cancer, cell death
 - Two pathways involved in DSB repair
 - **Nonhomologous end joining**
 - **Homologous recombination**
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Double-stranded break repair

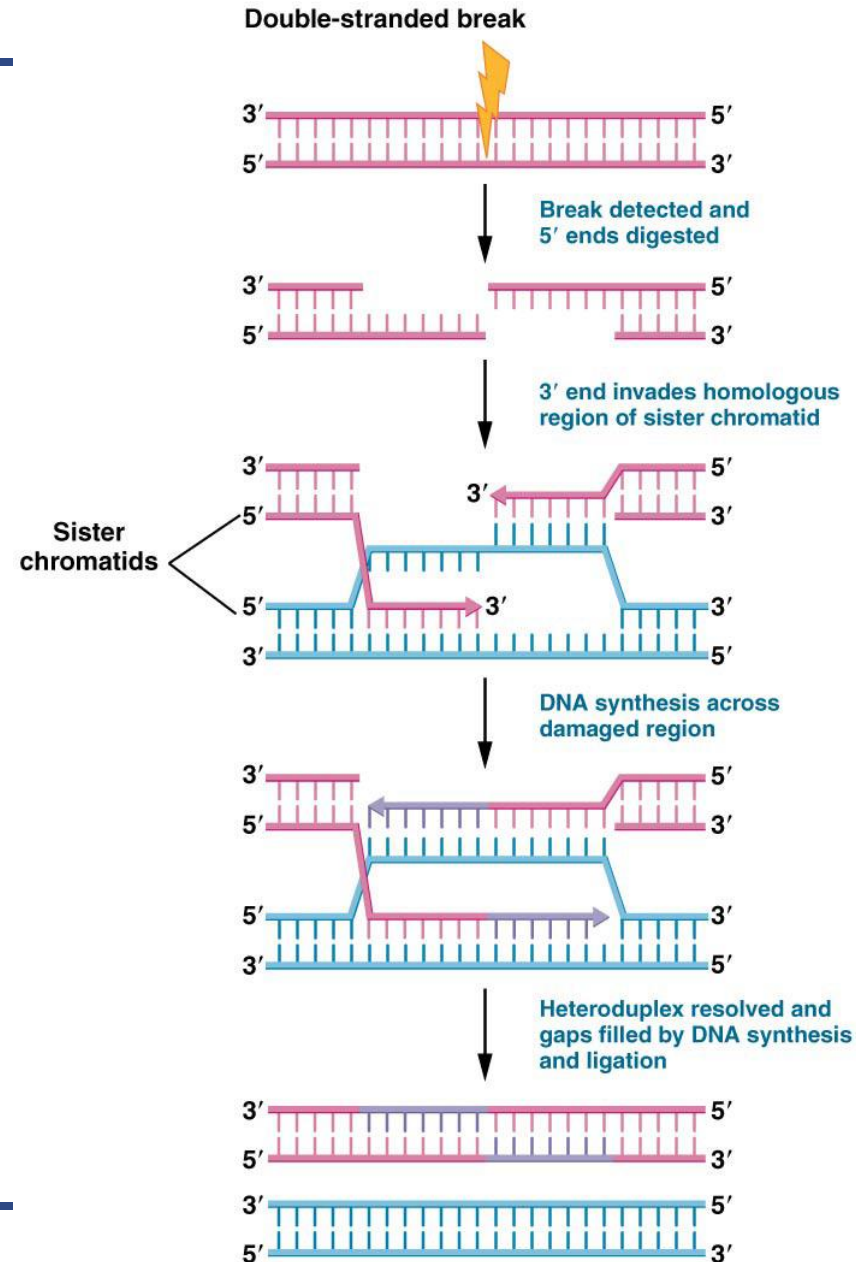
Nonhomologous end joining

- A complex of proteins binds the 2 free ends and ligate them back together
 - Error-prone: typically involves the loss of a few nucleotides at the cut site → non-homologous end joining tends to produce mutations, but this is better than the alternative (loss of an entire chromosome arm)
 - Occurs during G1 phase of the cell cycle
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Double-stranded break repair

Homologous Recombination

- Recognizes break, digests 5' end, and leaves 3' overhang
- 3' end aligns with sequence complementary on sister chromatid (homologous chromosome)
- Occurs during late S or early G2 phase of cell cycle



Defects in DNA repair

Consequences of defective DNA repair mechanism:

- Accumulation of mutations
- Premature aging and/or cancer

Example of disease: **Xeroderma pigmentosum**

- Autosomal recessive, can be caused by mutations in 9 different genes (*XPA*, *XPB*, *XPC*, etc.)
 - The nucleotide excision repair mechanism doesn't work
 - Accumulation of DNA damage due to UV light in skin cells → severe sunburn, keratosis, skin cancers...
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