



DNA Damage, Mutations and Repair Mechanisms



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* DNA repair لازم يكون قبل ال cell division (قبل DNA replication) لانه إذا صار damage بعد الانقسام بال daughter cell يكون permanent damage دائم وبسمية بماي الحالة mutation.



DNA Damage

- DNA molecules like all other biomolecules can be damaged in numerous ways
- DNA damage occurs at a rate of $10^4 - 10^6$ molecular lesions per cell per day (بالوضع الطبيعي)
- What are the sources and types of this damage?
- Can our cells recognise and repair this damage?
- What are the consequences of unrepaired damage on the cell fate?

DNA is repairable molecule unlike other molecule like Protein and lipid.

DNA → reversible damage
other molecule → irreversible damage.

DNA Damage

Classification of DNA Damage

- DNA damage can be classified according to the causative agents ^{العامل المسبب} into two main types:
 - A. Spontaneous damage (Endogenous)** ^{داخلي}: arising **naturally** and in the absence of known causative agents. Spontaneous DNA lesions are random events *so we can't predict when or in which cell the damage will take place.*
 - B. Induced damage (Exogenous)**: occurs in the presence of known causative agents (external factors) *like radiation.*

Spontaneous DNA Damage



- Although DNA is highly stable, nevertheless it is susceptible to the following spontaneous changes under normal cell conditions:

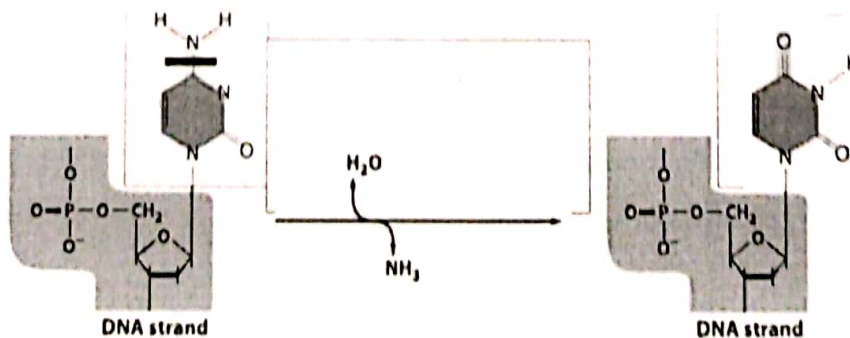
1. Deamination
2. Depurination
3. Replication errors
4. Base tautomers
5. Oxidative DNA damage

↳ the most common type of spontaneous damage.

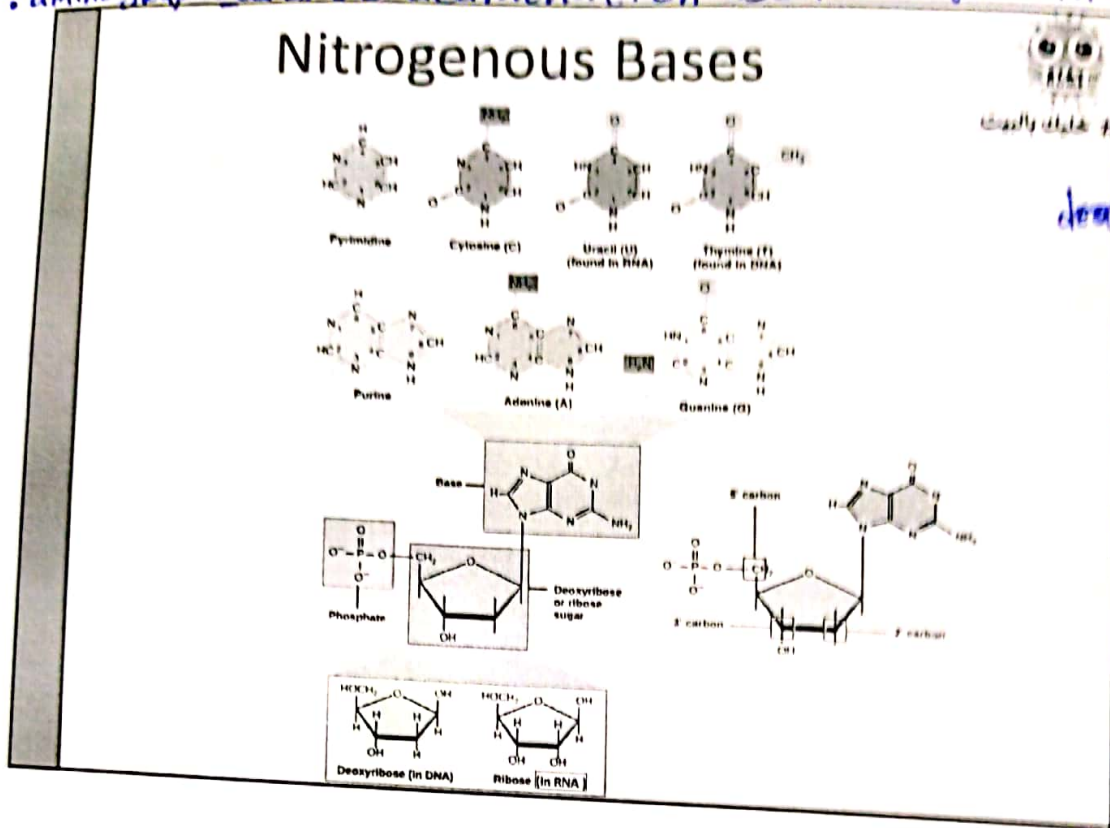
Spontaneous DNA Damage



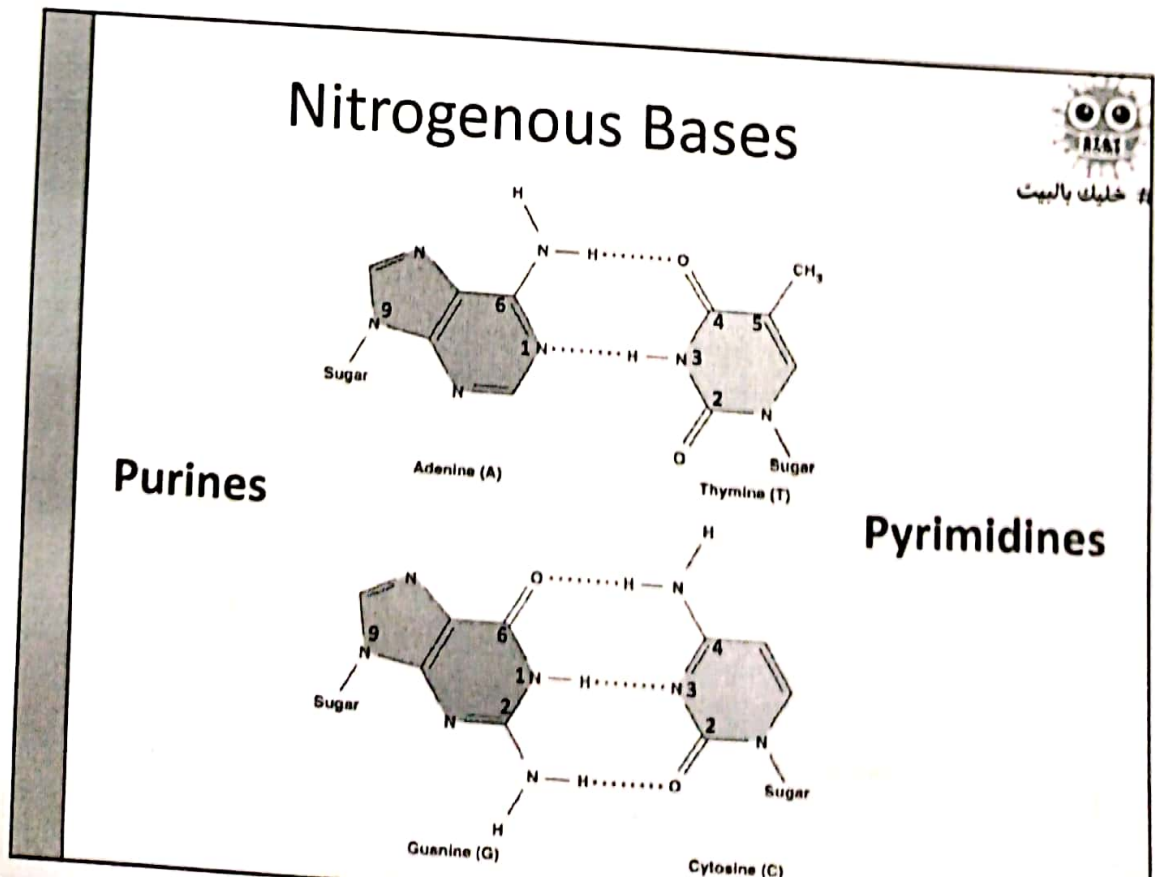
1. Deamination: the most common type is the spontaneous deamination of cytosine to uracil which occurs at a rate of about 100 bases/cell/day



* deaminated product of thiamin is thymine because it doesn't have amino group.
 • amino group is lost in deamination. $\text{Thymine} = \text{Thiamin} - \text{NH}_2$

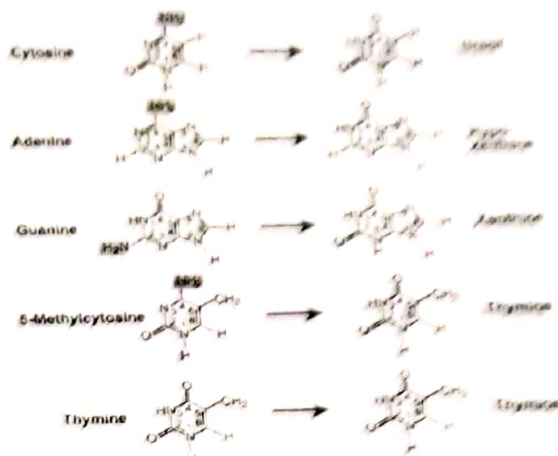


deaminated product

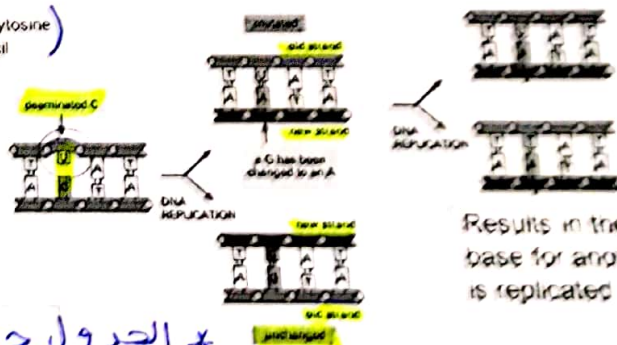


Deamination

- Other possible deamination events in DNA



Deamination



* الخلل و الحفظ

Nitrogenous base	Original base pair	Deamination product which base pairs with () 1 st round	Substituted base pair 2 nd round
Cytosine	C-G	Uracil (A)	T-A
Adenine	A-T	Hypoxanthine (C)	G-C
Guanine	G-C	Xanthine (T)	A-T
5-Me cytosine	C-G	Thymine (A)	T-A
Thymine	T-A	Thymine	T-A

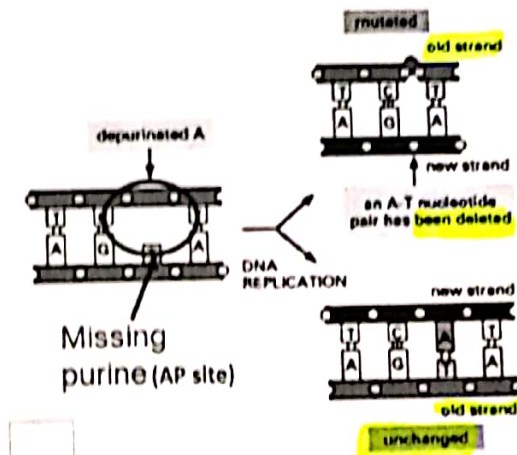
the consequences of this deamination event is the occurrence of mismatch base pairs.

Depurination



- Under physiological conditions, depurination occurs at a rate of about 5000 bases/cell/day
- Depurination results in apurinic site (AP site) which can be recognized and repaired by specific repair mechanisms
- If left uncorrected, during DNA replication these changes would lead to mutations in the daughter DNA chain (particularly base pair deletion) *تؤدي الى طفرات*
- This error will propagate throughout subsequent generations (inherited)

Depurination



DNA is repairable molecule because there are 2 types of backup system in DNA.

1- second complementary strand: guide the DNA repair enzyme to add original base pair

2- homologous chromosome: (بإذا كان الخلل بإل 2 strands of DNA)

Spontaneous

3. Replication errors: spontaneous lesions may occur during DNA replication in which the wrong base is added to the newly synthesized strand (base substitution), a DNA base is skipped (base deletion) or extra base is added (base insertion)
- Such errors are normally detected and repaired immediately by the proofreading/editing activity of DNA polymerase enzyme (3'-5' exonuclease activity)

DNA proofreading



- Otherwise, DNA repair enzymes will recognize the mismatched base pairs and repair them

Replication errors: spontaneous errors which escaped from proofreading activity if they aren't recognised by DNA polymerase enzyme.

Spontaneous DNA Damage

4. Base tautomers: DNA bases exist in one of several forms called tautomers (structural isomers)

1. Keto/Enol pair

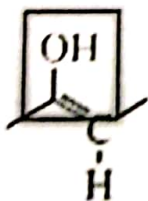
2. Amine/Imine pair

Enol form

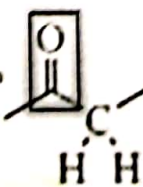
Keto form

Amine form

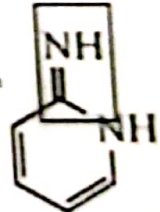
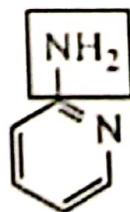
Imine form (rare)



tautomerization



tautomerization



enol form is rare

الكين من OH

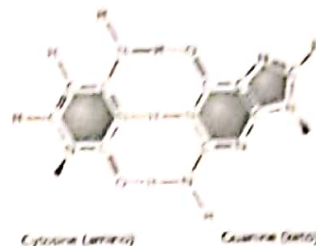
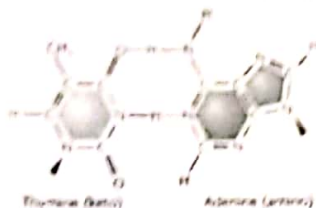
cytosine و adenine موجودين بال amino form
 thymine و guanine موجودين بال keto form

Base Tautomers

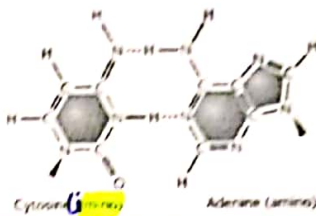


- Actually enol/imine forms are rare and tend to cause mispairing

(الوضع الطبيعي) Standard base pairing arrangements



(الوضع غير الطبيعي) Anomalous base pairing arrangements



3 H bond

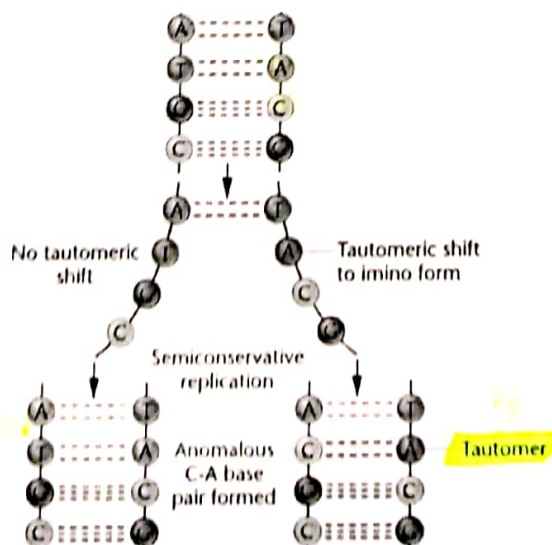
بينهم

(بينهم 2 H bond)

Base Tautomers



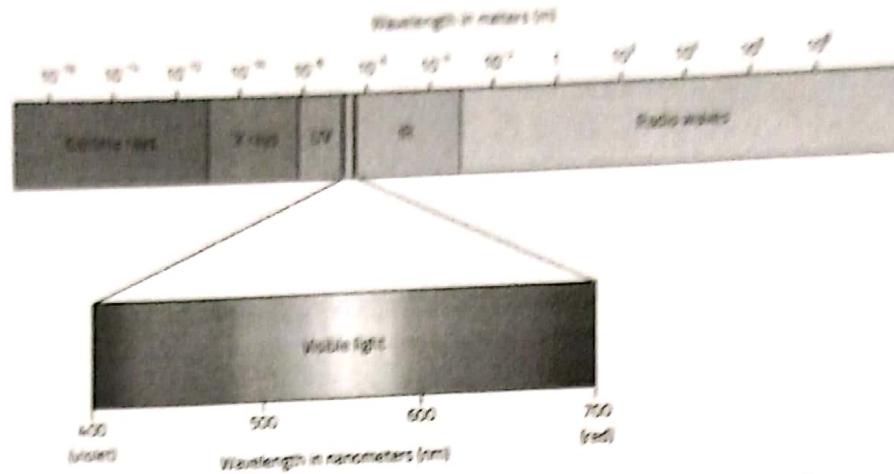
- If not repaired, it can lead to tautomeric shift mutation



Induced DNA damage



1. Radiation damage: which includes both UV light and ionizing radiation like x-rays

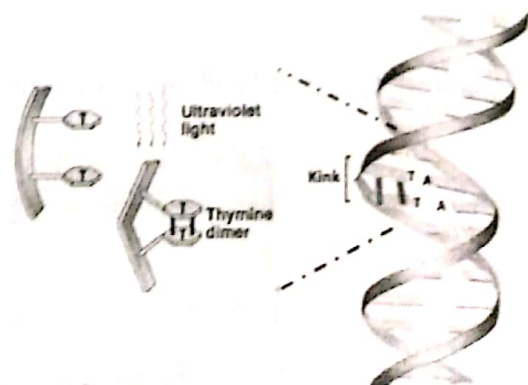


Induced DNA damage



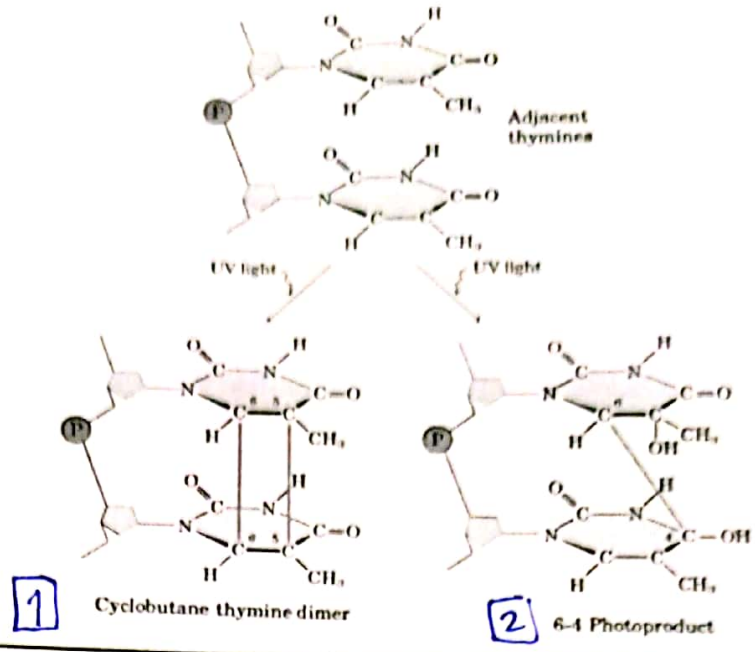
A. Ultraviolet Radiation

- Pyrimidines are highly sensitive to UV light. They form pyrimidine dimer (intra-strand crosslinking) particularly thymine dimer (T-dimer)
- Dimers alter DNA structure (kink or knot in DNA strand)
- Thymine dimers prevent proper replication. The cell either dies (apoptosis) or forms a malignant tumour (cancer)



this type of damage is called intra strand cross linking.

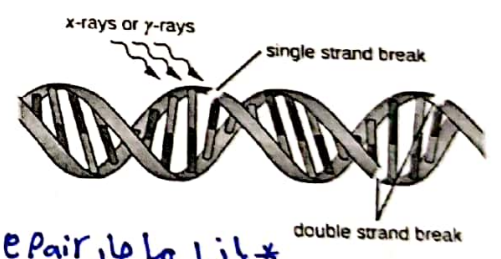
T-Dimer Types



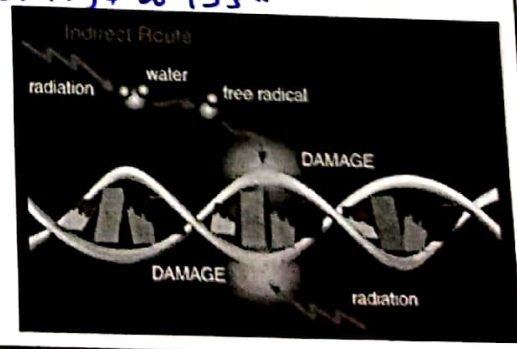
Induced DNA damage

B. Ionizing Radiation: like cosmic rays, X-rays and gamma rays can damage DNA molecules in 2 ways:

- Direct DNA damage by producing single strand break (SSB) and the more severe double strand break (DSB)



- Indirect DNA damage by production of free radicals which alter the structure of bases by generation of exogenous ROS.



(loss of genes) chromosomal aberration damage repair لهذا

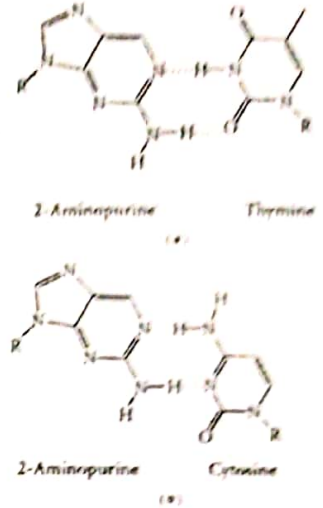
* اذا تعرضنا لا x-ray بشكل متكرر وعلى فترات متقاربة حيسير من الهعب
 باصلاح هذا damage بالتالي حيسهل cancer.



خليك بالبيت

Base analogs

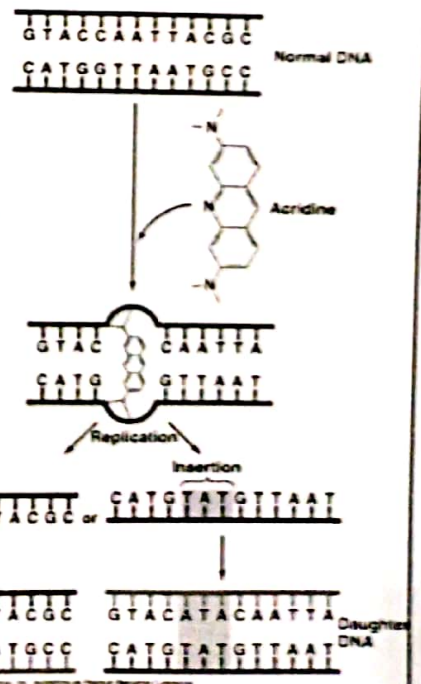
- Chemicals with structures similar to that of any of the four standard bases of DNA like 2-amino purine the base analog of adenine (6-amino purine). They replace them in DNA strand but do not **always** pair with normal bases leading to **base pair substitution** (e.g. AT bp is replaced with GC bp)



cytosine بال cytosine بال
* ال 2 amino Purin عند قابلية للارتباط بال cytosine بال
بالتالي جيني DNA damaged

Intercalating Agents

- Sandwich themselves between adjacent DNA bases like acridine orange, benzopyrene (cigarette smoke), aflatoxin B1 (mycotoxins produced by some fungi)
- They affect DNA structure causing insertion or deletion of an entire base pair leading to **frameshift mutation**



خليك بالبيت



DNA Repair Pathways

DNA Repair Mechanisms



- DNA repair system : is a collection of processes by which a cell identifies and corrects various DNA lesions
- Several repair strategies are available:
 - A. Direct/reversal repair
 - B. Base excision repair (BER)
 - C. Nucleotide excision repair (NER)
 - D. Strand-directed Mismatch repair (MMR)
 - E. Double strand breaks repair (DSB)

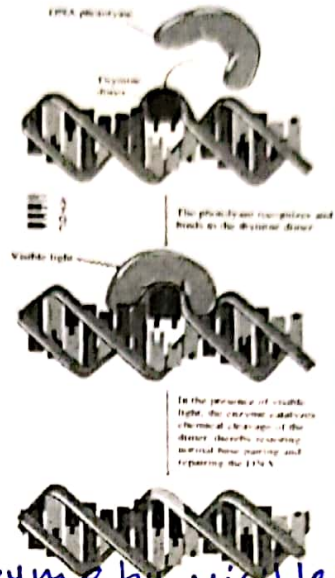
* حفظ الاختبارات ٢٥٥

Direct Repair system



- Direct repair also called direct reversal because the damage can be directly recognized and reversed
- Two specific enzymes are involved in direct repair:

1. Photolyases which repair UV induced damage in plants, bacteria and some animals (excluding humans) by splitting the dimers



* mechanism of repair
 1- activation of photolyase enzyme by visible light (Photoactivation)
 2- cleavage the T-dimer

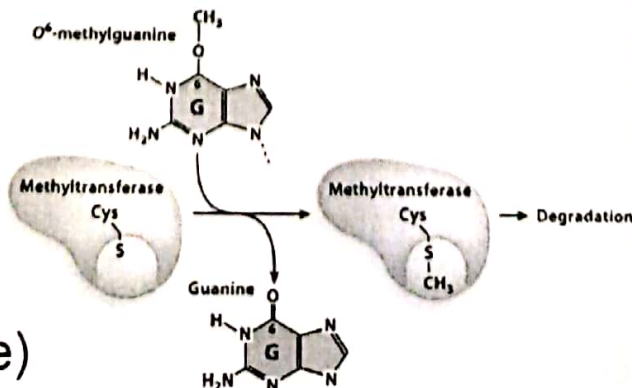
Direct Repair system



2. O6-methylguanine methyltransferase (MGMT) *حفظ الاختصار* which transfer methyl group from G to a cysteine residue within the enzyme itself. Ada and Ogt are the two bacterial isoforms of MGMT

* repair the damage result from base modifying agent.

- This reaction is stoichiometric rather than catalytic because each enzyme can be used only once (suicide)



This repair is specific to remove the modified pairs such as deamination or deamination or oxidative damage.

Base Excision Repair



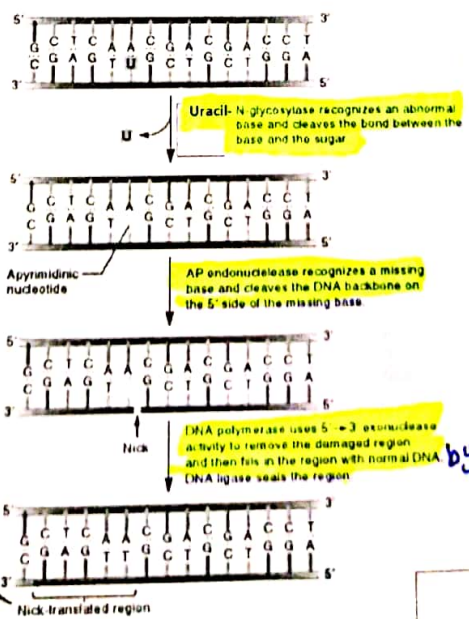
- Base excision repair (BER) involves a category of enzymes known as DNA-N-glycosylases like uracil DNA glycosylase
- Glycosylases recognize damaged bases and remove them resulting in apurinic or apyrimidinic (AP) site
- AP endonucleases enzymes nick the damaged backbone at 5' end of AP site
- DNA polymerase removes the damaged region using its 5' to 3' exonuclease activity and correctly synthesizes the new strand. Finally, DNA ligase seals the strand.

Base Excision Repair



Base Excision Repair System

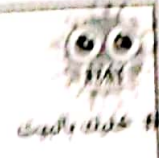
Depending on whether a purine or pyrimidine is removed, this creates an apurinic and an apyrimidinic site, respectively



Nick replication would be a more accurate term

by DNA polymerase elongation activity.

Nucleotide Excision Repair



- Nucleotide excision repair system (NER) corrects lesions which commonly cause bulk distortions in DNA helix like UV-induced pyrimidine dimers. NER is highly conserved used in both eukaryotes and prokaryotes

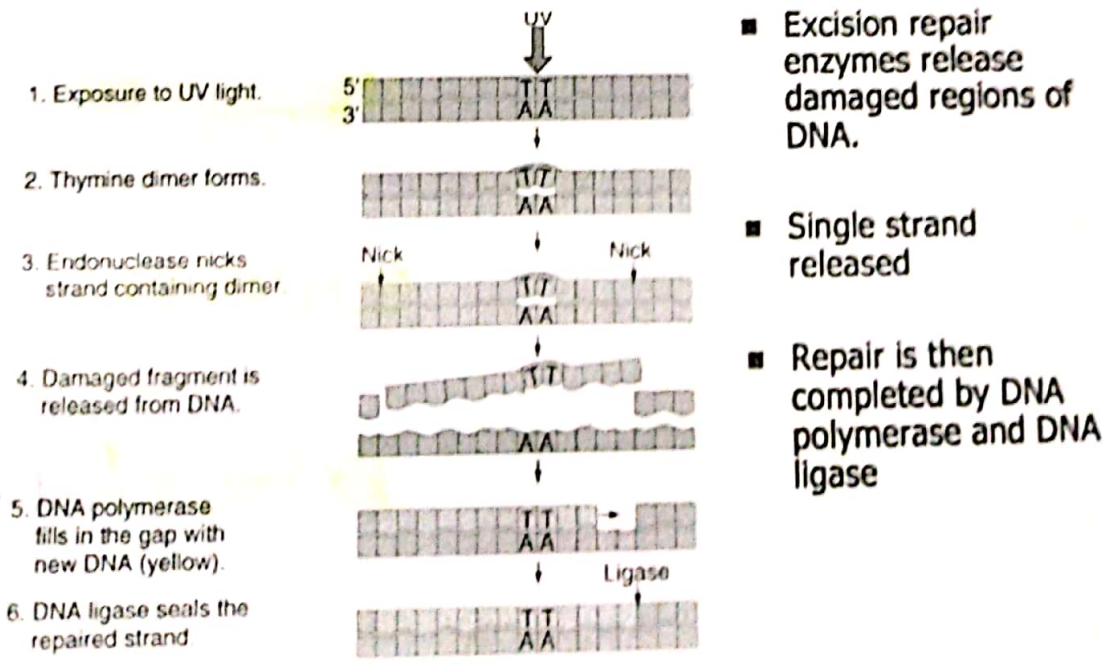
- The damaged region is removed in 3 steps process:
 1. Recognition of the damage by NER enzymes
 2. Excision of damaged DNA (12-24 nucleotides long) by endonucleases
 3. Resynthesis of removed DNA region by DNA polymerase followed by ligase to seal the region

السي با لصندوق الازرق
للإطلاق فقط

Nucleotide Excision Repair



* الـ سي كامل للإطلاق فقط



Nucleotide Excision Repair



- Xeroderma pigmentosum (XP) is a recessive disorder in which victims lack the normal UV repair enzymes (NER genes). This creates hypersensitivity to sunlight and a tendency to develop skin cancer

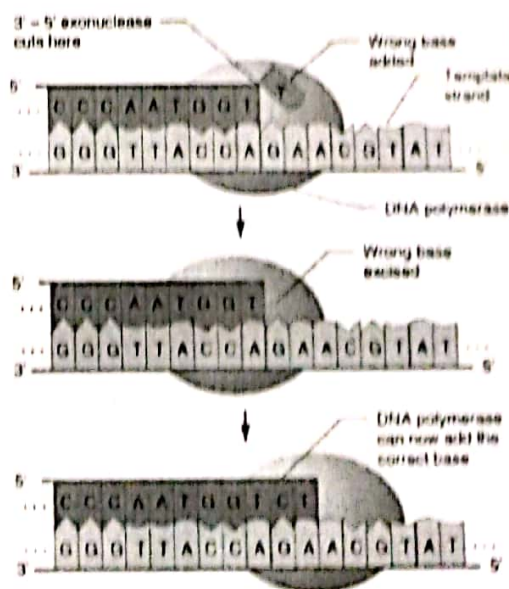


Strand-directed Mismatch Repair



*this repair occur when the error during DNA replication escaped from proofreading activity.

- Mismatch repair (**MMR**) corrects errors introduced during DNA replication (e.g. base substitution, deletions or insertions)
- Replication errors are rare due to high fidelity of DNA replication process
- DNA polymerases have proofreading 3'-5' exonuclease (reverse) activity which recognizes mismatched bases and excises them



Strand-directed Mismatch Repair

- Mismatch system recognizes and corrects errors that escaped from DNA polymerase proofreading machinery

• 3 steps process:

1. Mismatched base pair is recognized
2. Excision of DNA segment containing the mismatched nucleotide from the newly synthesized strand

③ Resynthesis of the excised segment

- It is called strand-directed MMR because MMR enzymes are selectively directed to the newly synthesized strand rather than to the old strand

(it can distinguish between old strand and new strand)



old strand

Double strand breaks repair (DSB)

- A dangerous type of DNA damage which can lead to chromosomes fragmentation and consequently loss of genes (**chromosomal aberration**) if left unrepaired

• Two types of repair mechanisms:

① Non-homologous End Joining (NHEJ): it is an error-prone mechanism of repair because it results in change of DNA sequence at the site of breakage

② Homologous recombination (HR) is an error-free mechanism of repair because the damage is accurately repaired using information from sister chromatid

not completely efficient

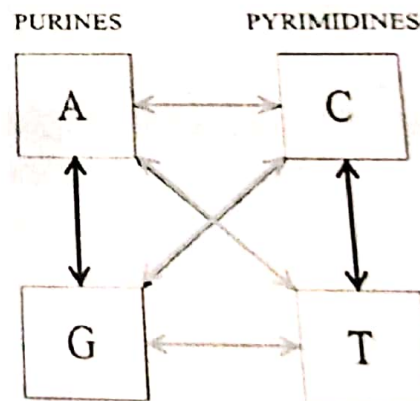
completely efficient

Types of mutations at the DNA level	Results at the molecular level
No mutation Wild type	<div style="display: flex; justify-content: space-around; align-items: center;"> <div style="text-align: center;"> <p>Thr Lys Arg Gly</p> <p>Codon 1 Codon 2 Codon 3 Codon 4</p> <p>A C A A G A G A G G T</p> </div> <div style="text-align: center;"> <p>Codons specify wild-type protein</p> </div> </div>
Base insertion Frameshift mutation	<div style="display: flex; justify-content: space-around; align-items: center;"> <div style="text-align: center;"> <p>Thr Glu Glu Arg</p> <p>A C A A A A G A G A G G T</p> </div> </div>
Base deletion Frameshift mutation	<div style="display: flex; justify-content: space-around; align-items: center;"> <div style="text-align: center;"> <p>Thr Arg Glu Val</p> <p>A C A A G A G A G G T</p> </div> </div>

Point Mutation



- Point mutation: an alteration in DNA sequence by a single nucleotide base and consequently a change in single base pair (substitution)
- Substitution at a point is called **Transition** if one purine is replaced with another purine or one pyrimidine with another pyrimidine and it is called **Transversion** if one purine is replaced with one pyrimidine or vice versa



اد transition لها اربع احتمالات للحدوث بينما transversion لها ثمن احتمالات

Point Mutation



خليك بالبيت

No mutation

Point mutations

Silent

Nonsense

Missense

DNA level
mRNA level
protein level

TTC
AAG
Lys



TTT
AAA
Lys



ATC
UAG
STOP

Conservative
TCC
AGG
Arg



non-conservative
TGC
ACG
Thr

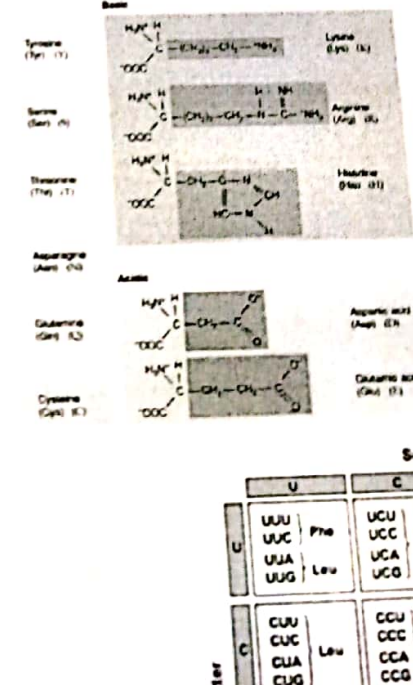
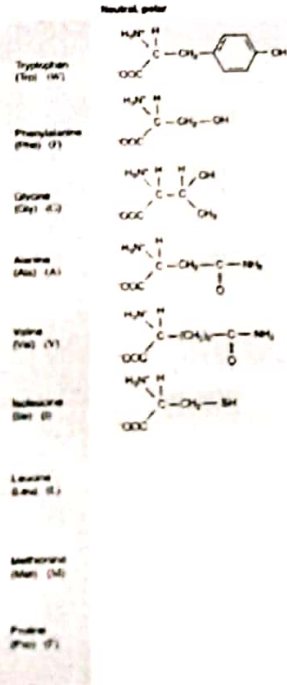
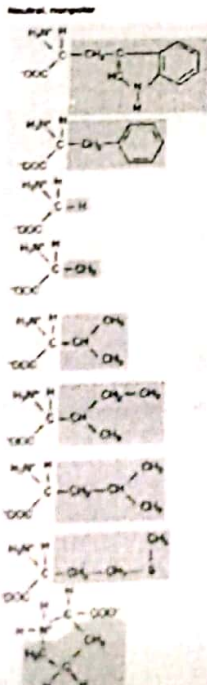


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Point Mutation



خليك بالبيت



		Second letter				
		U	C	A	G	
U	UUU	UCU	UAU	UGU	U C A G	
	UUC	UCC	UAC	UGC		
	UUA	UCA	UAA	UGA		
C	CUU	CCU	CAU	CGU	U C A G	
	CUC	CCC	CAC	CGC		
	CUA	CCA	CAA	CGA		
A	AUU	ACU	AAU	AGU	U C A G	
	AUC	ACC	AAC	AGC		
	AUA	ACA	AAA	AGA		
G	GUU	GCU	GAU	GGU	U C A G	
	GUC	GCC	GAC	GGC		
	GUA	GCA	GAA	GGA		

مطلوب معرفة كل amino acid
بأي مجموعة موجودة موجودة