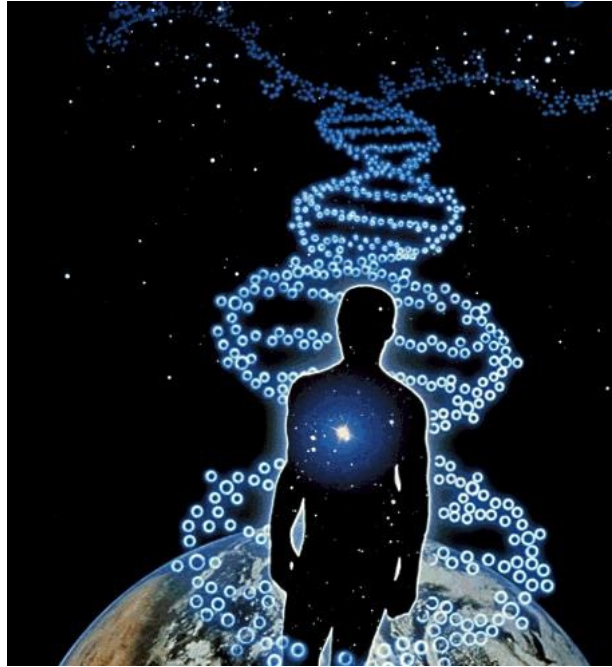
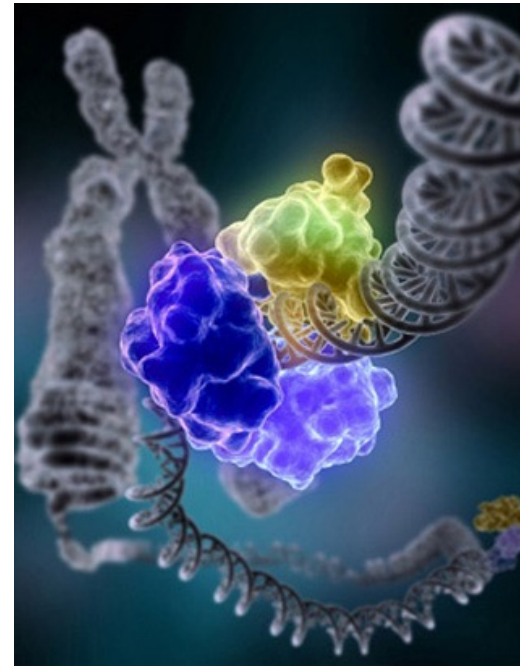


# DNA (damage)



# Repair



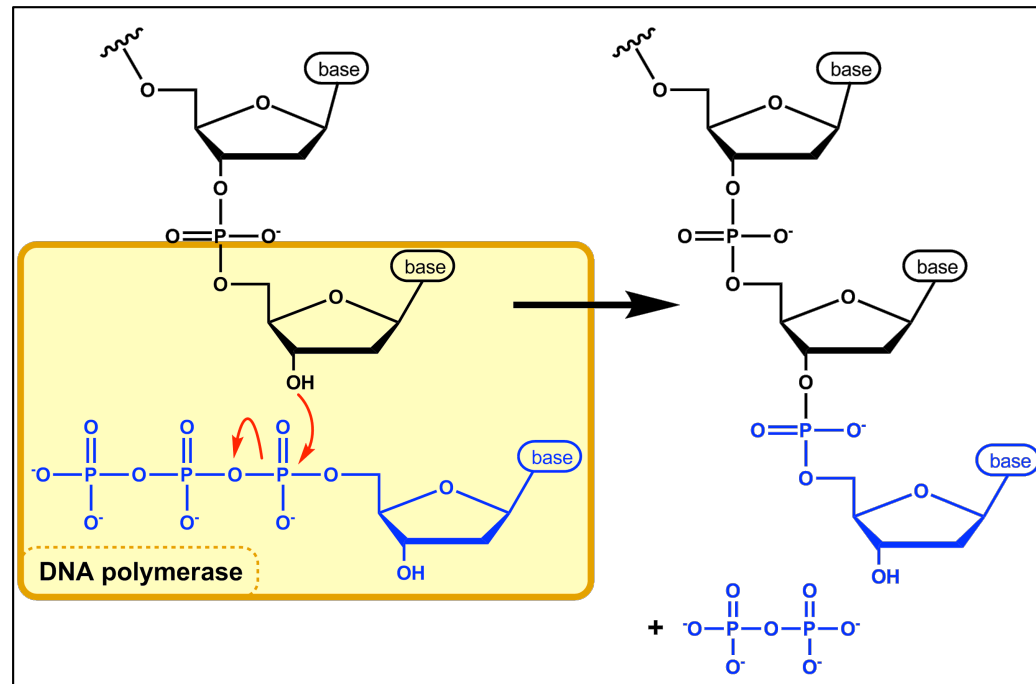
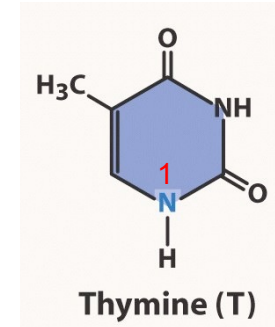
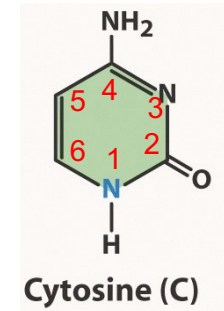
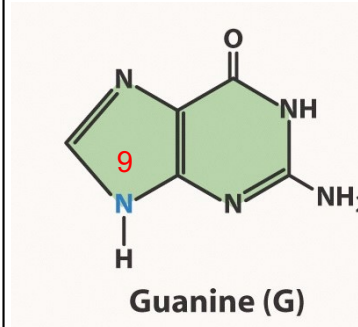
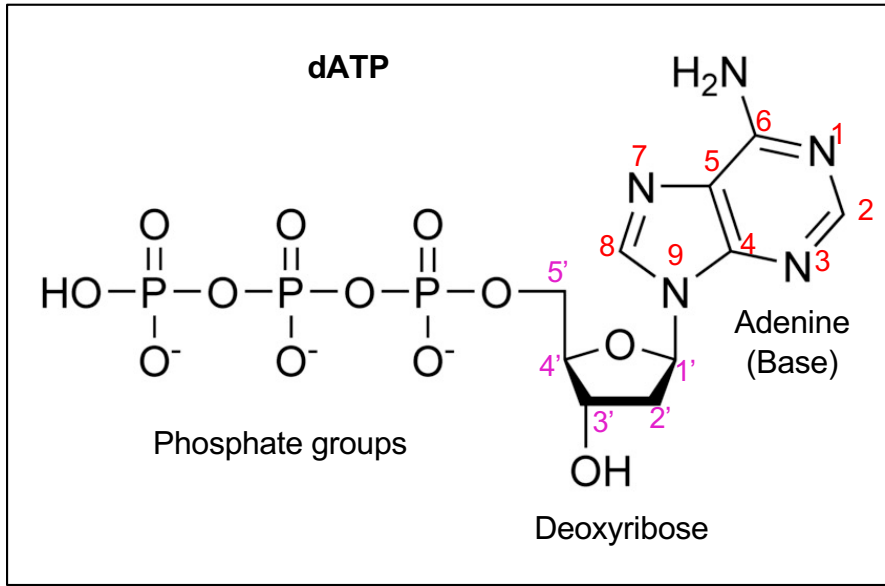
Part 1: DNA is **biochemically unstable**. Large **numbers** & many **types** of DNA lesions occur daily.

**Part 2: Multiple repair mechanisms** cope with different types of DNA lesions.

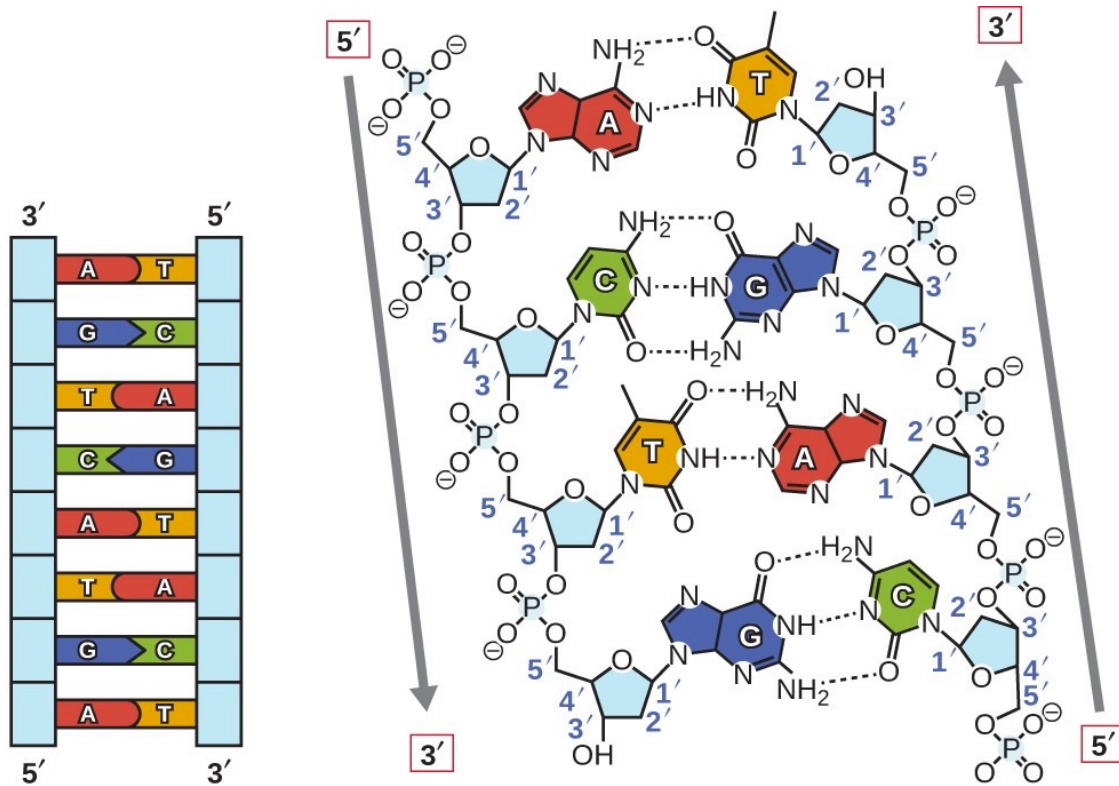
- Overview of four classical DNA repair pathways
- Three examples of DNA repair pathways.
- Outstanding questions.

- ❖ DNA repair is intimately **connected with** many other processes (replication, transcription, epigenetics, cell cycle, cell types, signaling, development, innate immunity, metabolism.....).
- ❖ DNA repair defects cause **hundreds of human diseases**.
- ❖ DNA repair is both **cancer-suppressive** and **treatment target**

# A quick recap: chemical composition of DNA



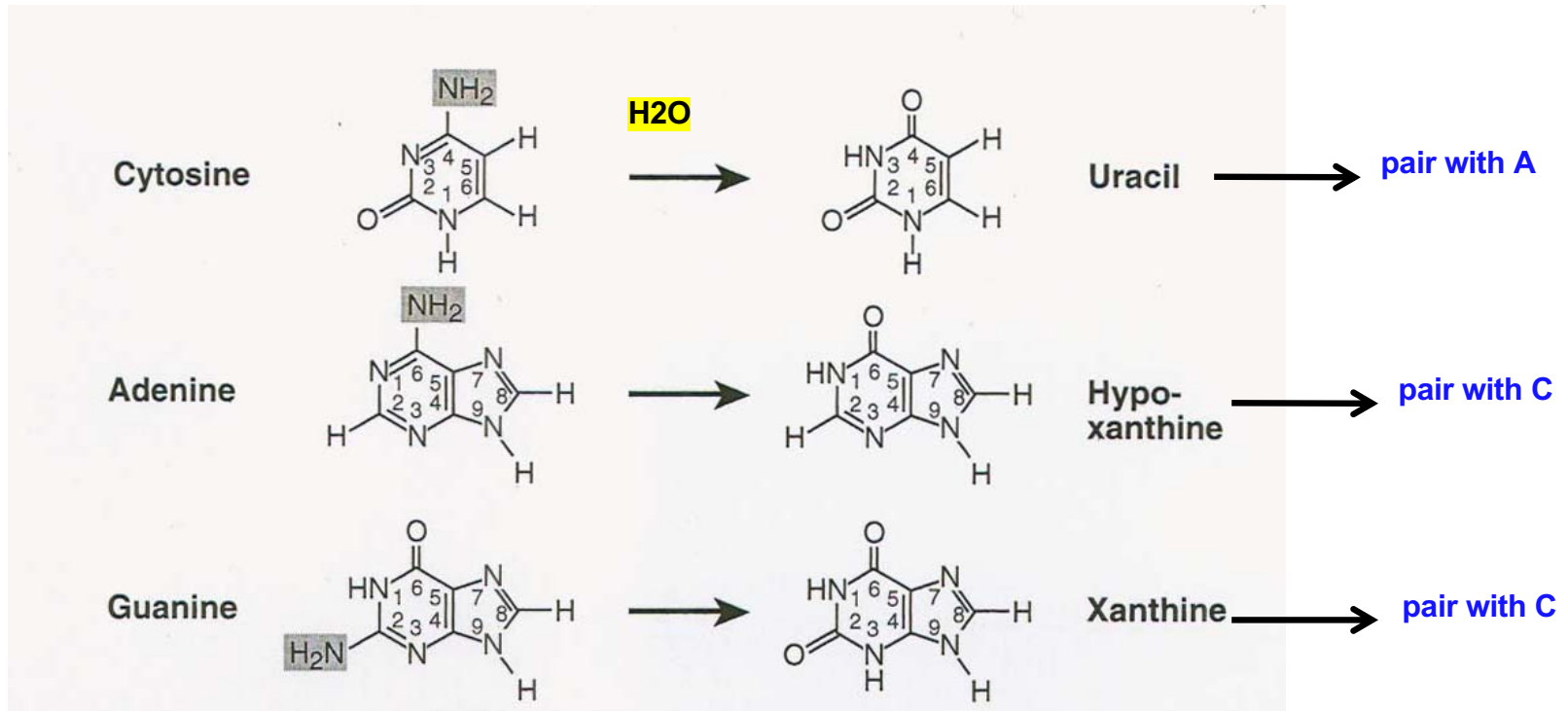
# Where do lesions occur on DNA?



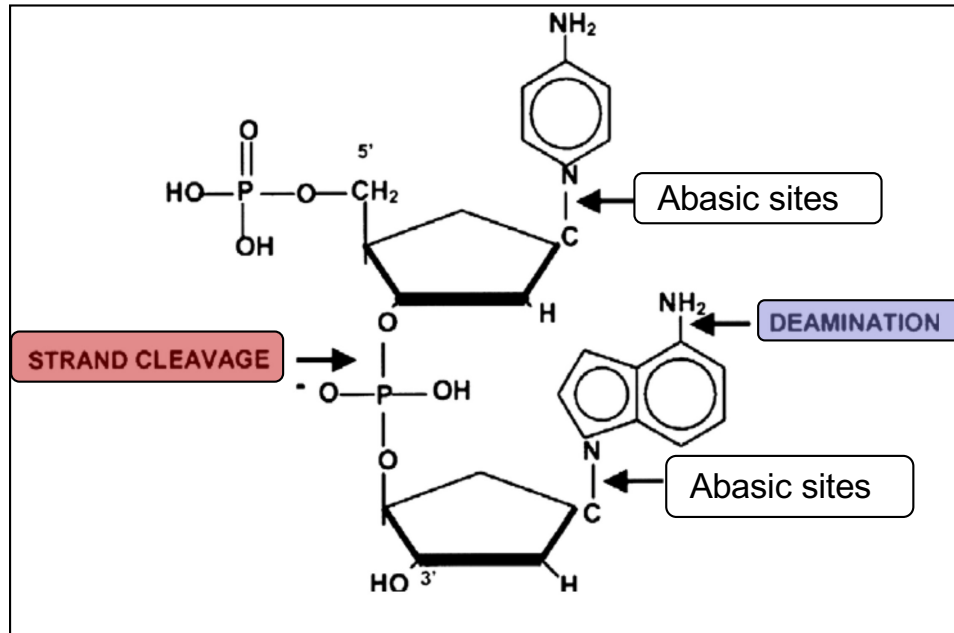
3 billion base pairs of DNA in human genome

# Hydrolysis - loss of small chemical groups

**Deamination** can occur at 3 bases



## Hydrolysis - Loss of entire base or ssDNA break

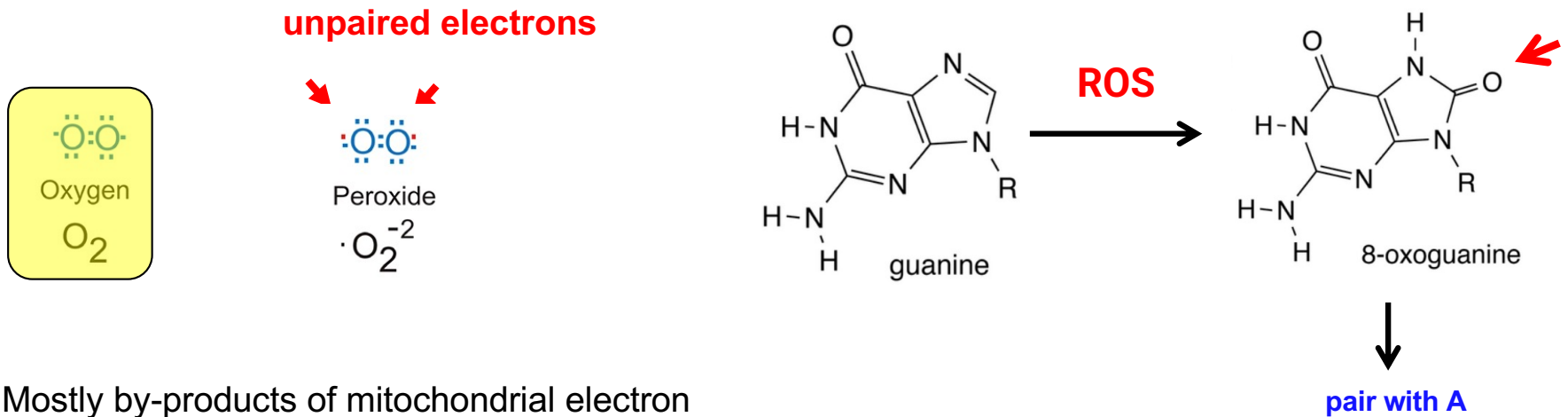


**Abasic sites are generated when a base is lost by** Hydrolysis of the N-glycosyl bonds by water.

Abasic sites are not stable and can lead to ssDNA breaks.

# Oxidation: react with small chemical groups & alter base structures

## Oxygen and Reactive oxygen species (ROS)

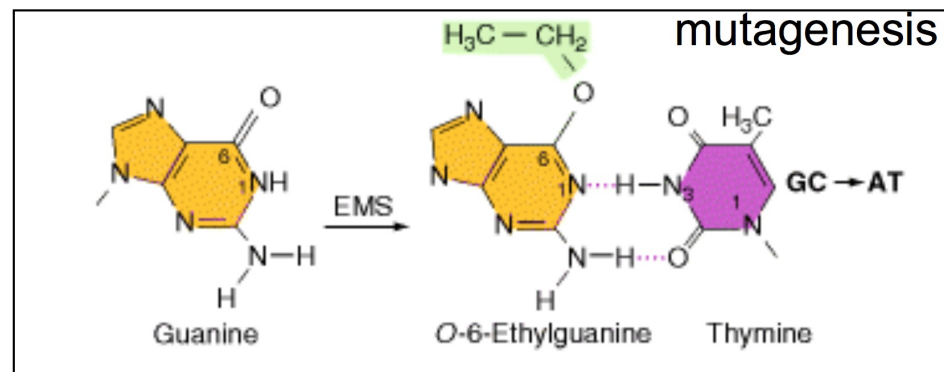


Mostly by-products of mitochondrial electron transport & metal catalyzed oxidation reactions.

# Methylation – gain of small chemical groups on the bases

(mediated by chemicals or enzymes )

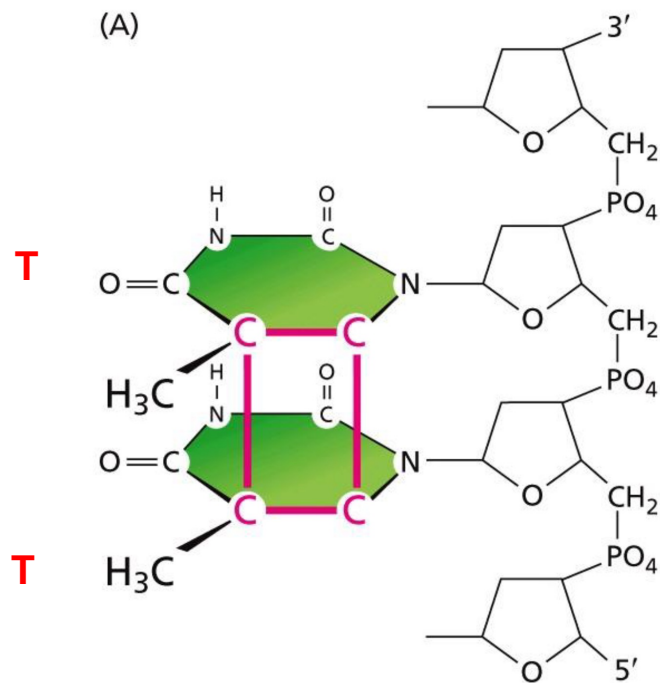
**Methylation** can occur at several places on the 4 bases



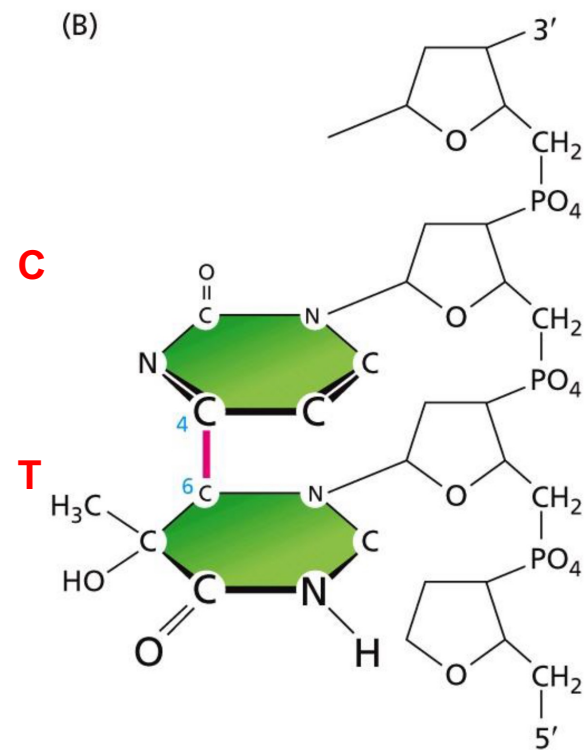
# Covalent linkages between bases on the same strand

## UV photoproducts

The most frequent photoproducts are covalent linkages between adjacent Pyrimidines: cytosine and thymine



CPD  
(cyclobutane pyrimidine dimer)

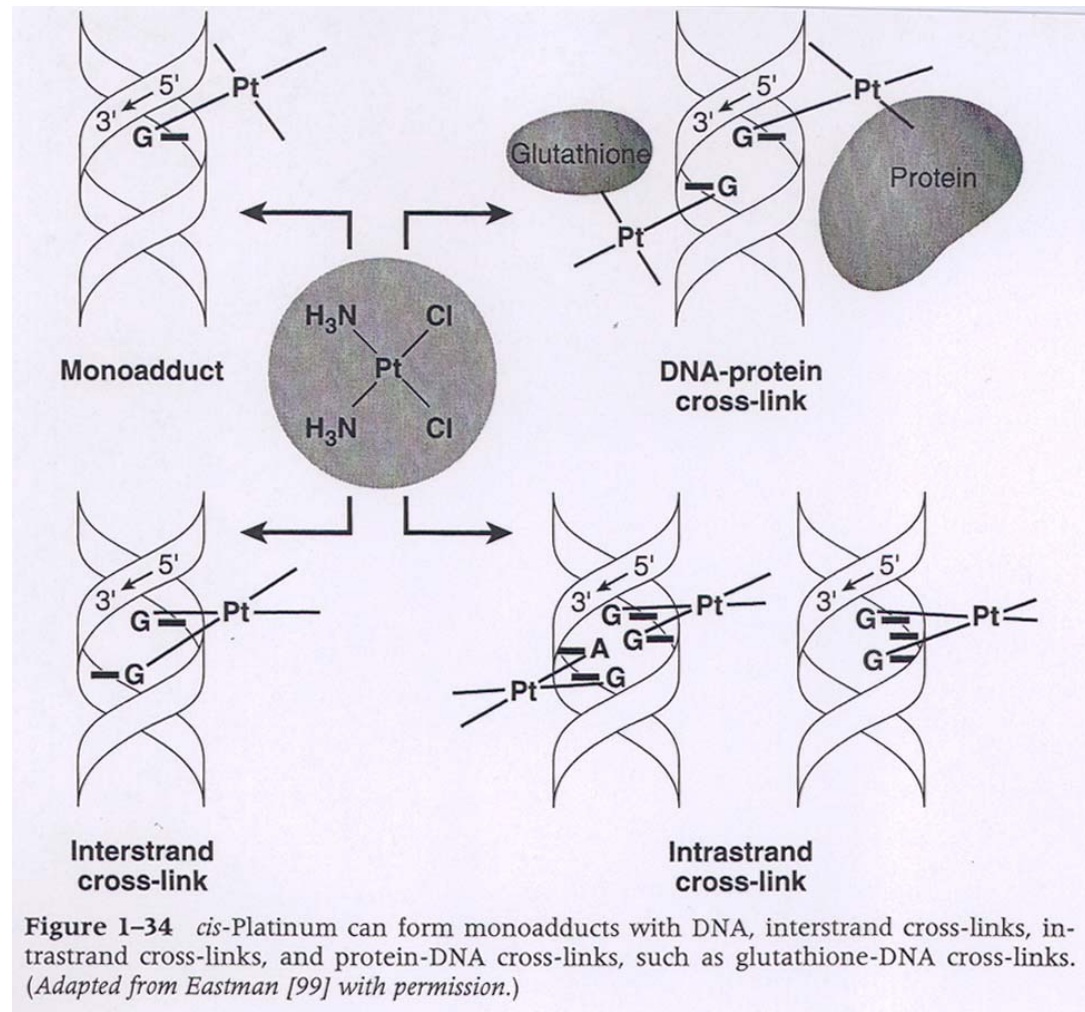


(6-4)-PP  
(6-4)-photoproduct



# Crosslink agents and different types of DNA lesions

**cis-Platinum leads to several types of crosslinks on DNA**



# Estimated amounts of base lesions

**Table 2-1** Endogenous DNA lesions arising and repaired in a diploid mammalian cell in 24 h

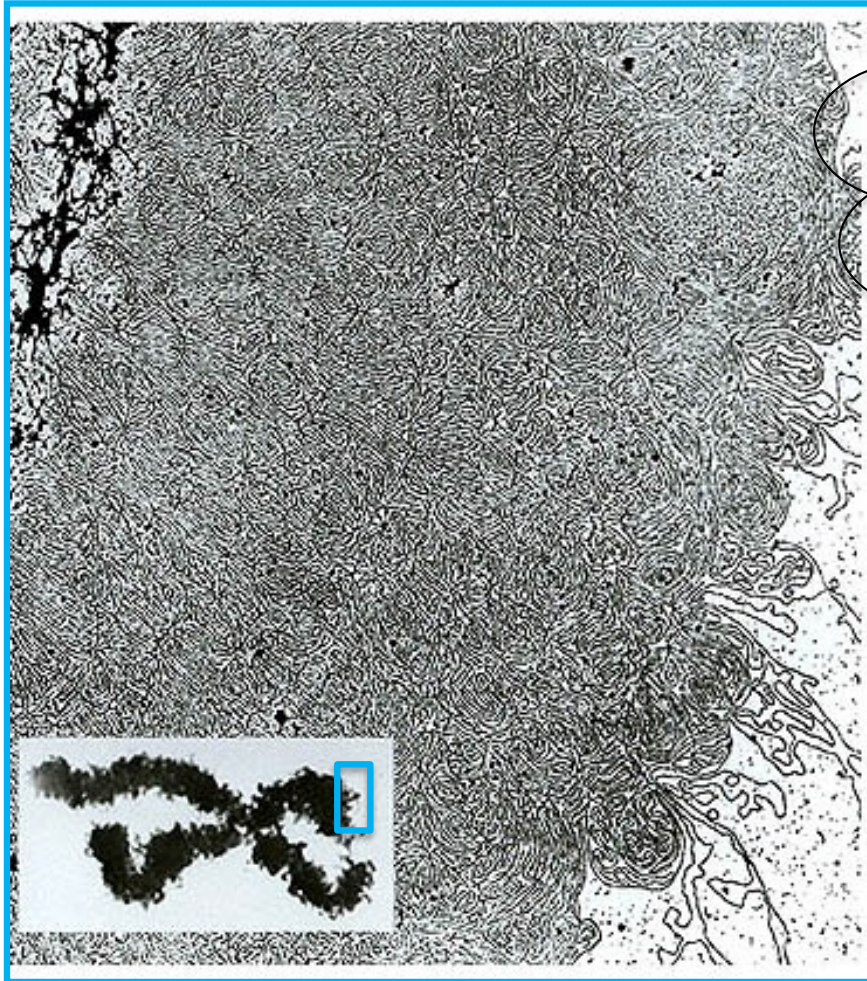
Endogenous source		No. of lesions
		100% double-stranded DNA
<b>Hydrolysis</b>		
Depurination	] Abasic site	18,000
Depyrimidination		600
Cytosine deamination		100
5-Methylcytosine deamination		10
<b>Oxidation</b>		
8-oxoG		~1,000–2,000
Ring-saturated pyrimidines (thymine glycol, cytosine hydrates)		~2,000
Lipid peroxidation products (M <sub>1</sub> G, etheno-A, etheno-C)		~1,000
<b>Nonenzymatic methylation by S-adenosylmethionine</b>		
7-Methylguanine		6,000
3-Methyladenine		1,200
1-Methyladenine and 3-methylcytosine		ND <sup>c</sup>
<b>Nonenzymatic methylation by nitrosated polyamines and peptides</b>		
O <sup>6</sup> -Methylguanine		20–100

<sup>a</sup>Data from reference 297.

<sup>b</sup>Estimates are for two  $3 \times 10^9$  -bp genomes per cell.

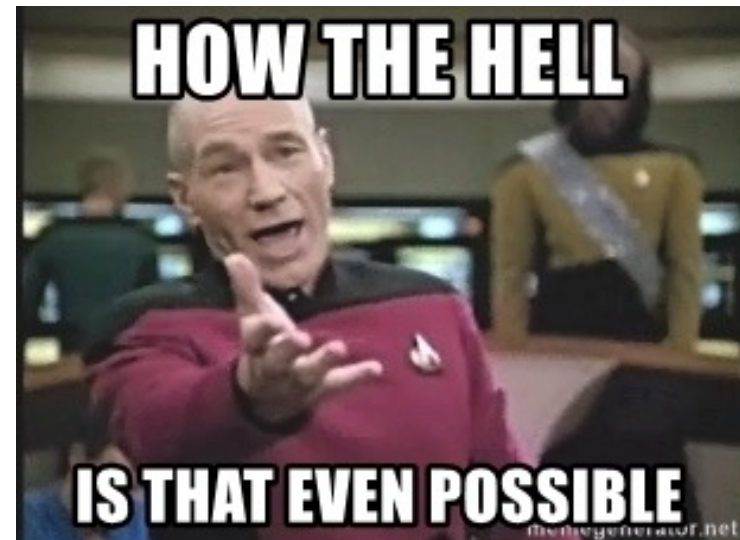
<sup>c</sup>ND, none detected

## Repair of eukaryotic genome is challenging

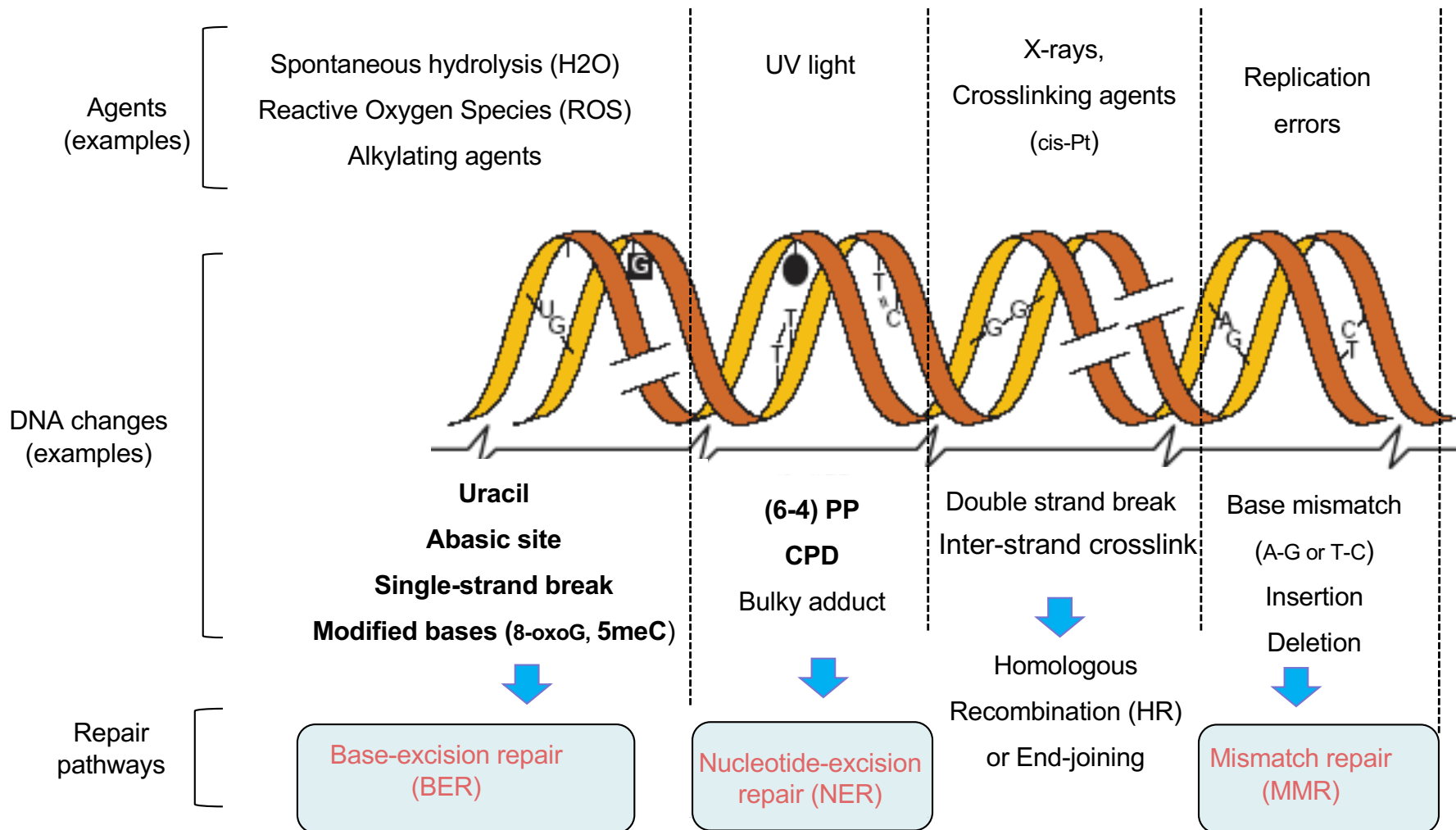


Electron micrograph of DNA from a small part of a chromosome

How can cell fix the huge numbers of DNA lesions in the human genome to minimize deleterious mutations that can lead to hundreds of types of diseases?!



# DNA lesions and repair pathways (partial)



Several new types of DNA repair pathways

# Common features of DNA repair pathways

## Stage I

### Detecting DNA lesions

- Proteins recognize specific modified bases (BER)
- Proteins recognize DNA strand distortion (NER)
- Proteins bind to mismatched bases (MMR)
- Proteins recognize broken ends (HR, EJ)

## Stage II

### Removing lesions and clean-up

- Different types of **nucleases & other enzymes** to remove different types of lesions
- lesion recognizing proteins + scaffold proteins: recruit lesion removal enzymes

## Stage III

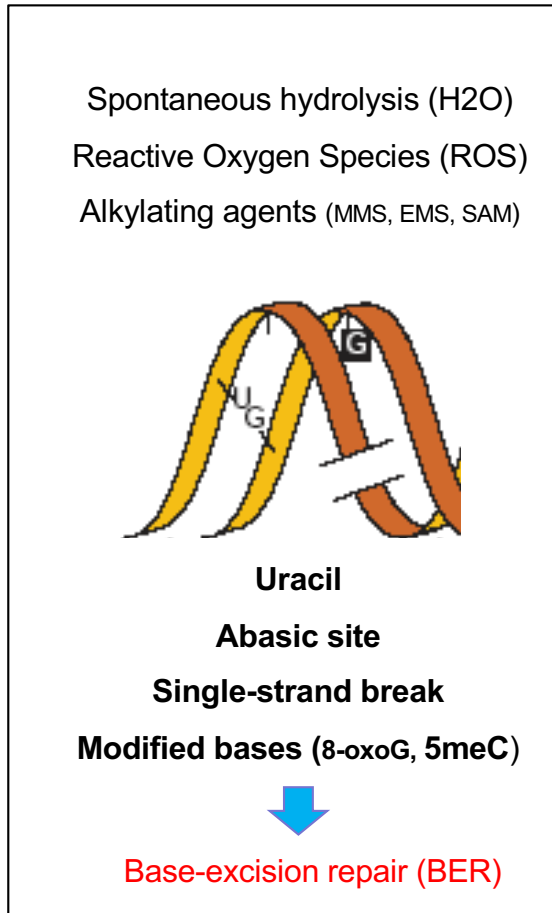
### Synthesizing new DNA

- More than a dozen of DNA polymerases to be deployed for repair tasks
- These polymerases are less precise and require tight regulations.
- Other proteins required: DNA ligase and more clean-up enzymes.

### Important Considerations:

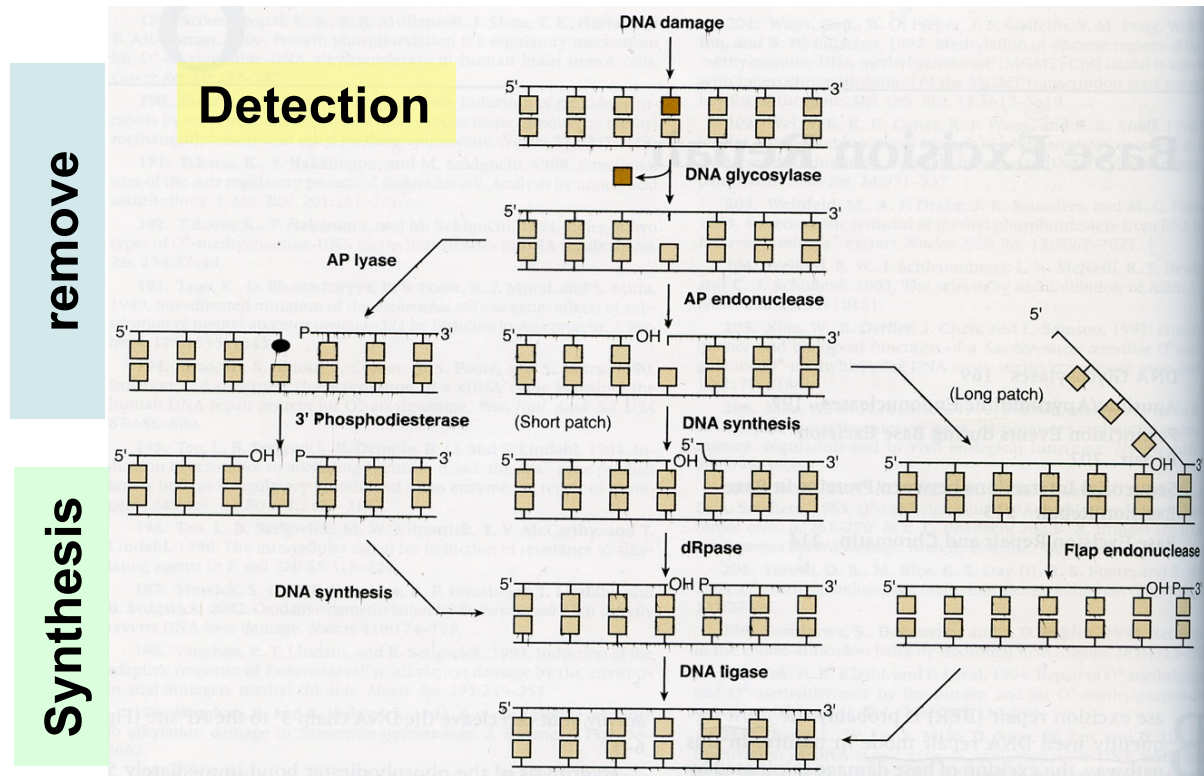
- Processivity of the multi-step repair processes.
- Cope with chromatin contexts and restore epigenetic states afterwards.
- DNA repair efficiency in response to genotoxins.
- Coordinate with other processes, such as transcription.

# Base Excision Repair (BER)



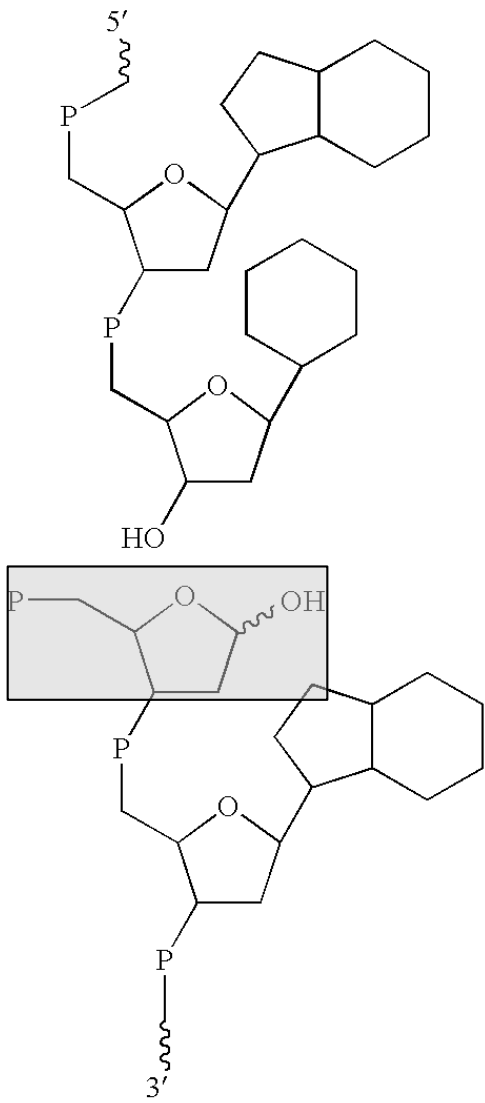
## Damage Recognized:

Base deamination,  
 Base oxidation,  
 Base methylations



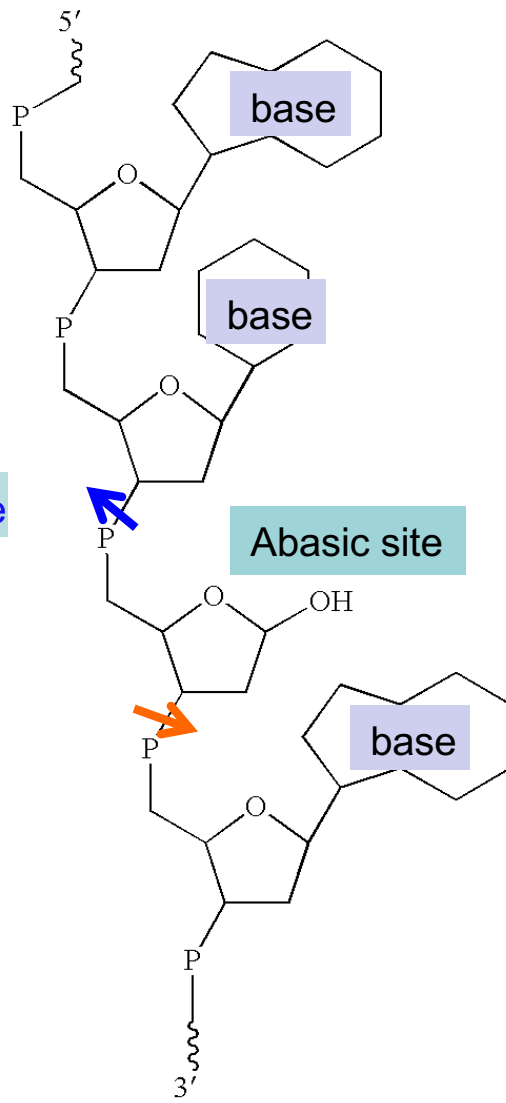
## Gene Products Required (7):

1. **Glycosylases (>11 types in human)**
2. **AP lyase + 3' Phosphodiesterase**
3. **AP endonuclease + dRpase (Deoxyribo-phosphodisterase)**
4. **DNA polymerase**
5. **Flap nuclease**
6. **DNA Ligase**



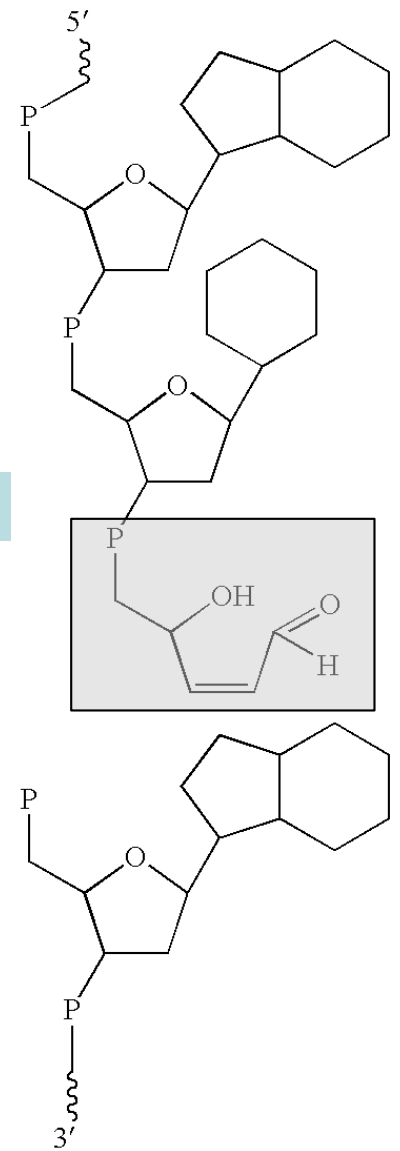
Deoxyribo-phosphodiesterase  
(dRpase)

AP Endonuclease



Abasic site

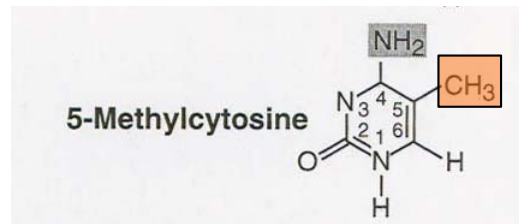
AP Lyase



3' Phosphodiesterase

# BER in development

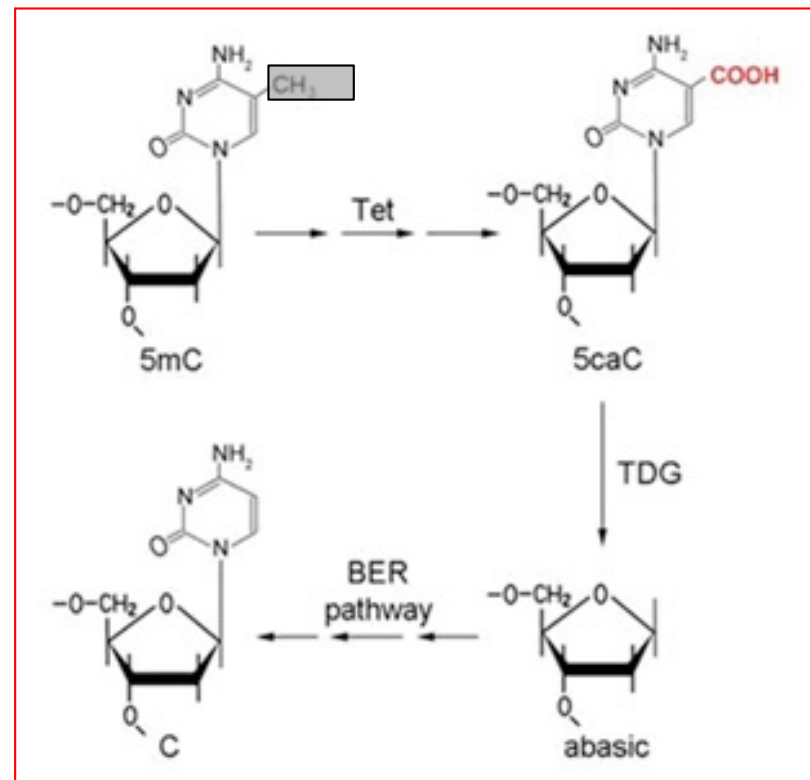
**Base modification: a means for functional variability, while maintaining genetic information**



- In mammalian genomes, 5mC exists mostly in CpG context (~80% CpGs methylated) by DNMTs.
- 5mC - transcription repression, used widely for control lineage-specific genes, X-chr & mobile elements inactivation.

**Demethylation is observed:**

1. early in development, some posterior epiblast cells become primordial germ cells wherein loss of 5mC and prepares them for germ-cell-specific processes.
2. At specific loci in response to stimuli.



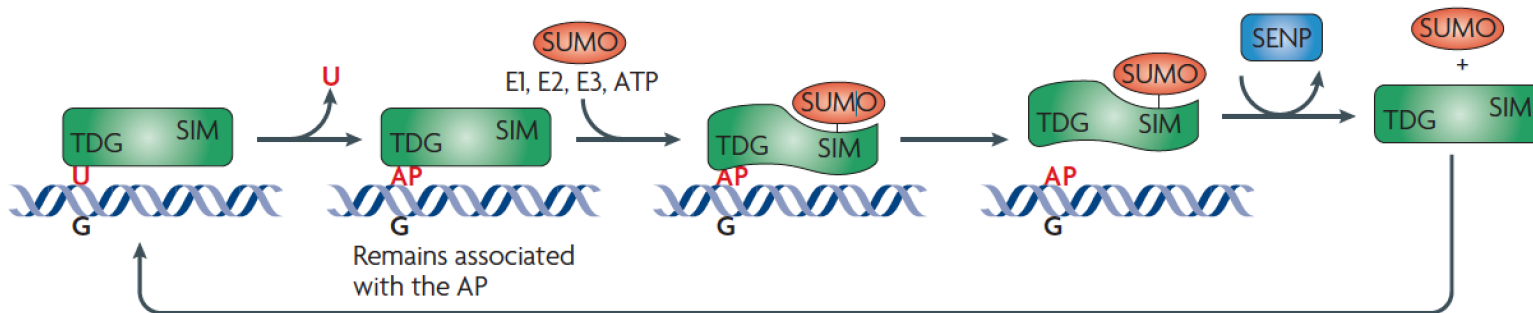


## How are BER enzymes coordinated?

*Hands-off Model: proceed by the sequential action of factors facilitated by pair-wise interactions*

- Protect reaction intermediates & ensure the completion of the reaction once initiated.
- Reaction accuracy and specificity.

**TDG requires sumoylation and desumoylation for each catalytic cycle.**



# Key proteins & steps in Nucleotide Excision Repair (NER)

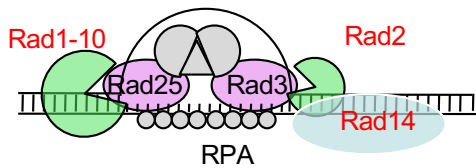
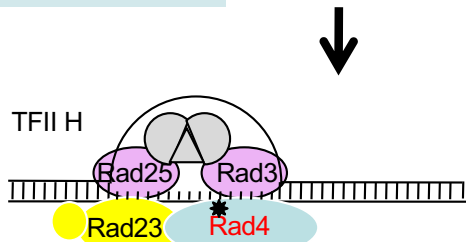
## Detection



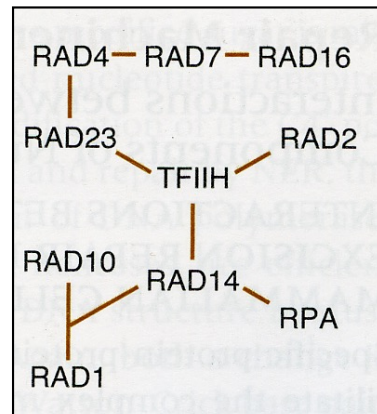
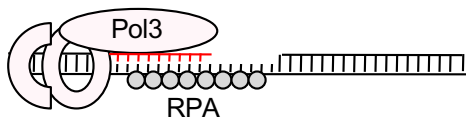
## Damage Recognized:

Large changes in the nucleotide – eg. CPD

## remove



## Synthesis



## Model of the Global Genomic NER (GG-NER) mechanism.

- (A) A lesion induces DNA helix distortion;
- (B) **Rad4-23** detects helix distortion & stabilizes DNA bend;
- (C) **Rad4-23** recruits **TFIIH** at the site of the lesion;
- (D) **TFIIH** (10 subunits, txc component, +ATP) unwinds DNA, until the Rad3 helicase encounters a modified base; the other helicase Rad25 goes on unwinding DNA to create a 20-bp "bubble" structure;
- (E) Rad14, Rad2, and RPA are recruited
- (F) Rad1-Rad10 joins the complex to enable dual incision (5' cut by Rad1-10 and 3' cut by Rad2).
- (G) RPA remains on ssDNA to facilitates transition to repair synthesis by Pol III with the help of RFC and PCNA; ligase I finally seals the nick.

# NER in human diseases

## Diseases associate with NER

XP (Xeroderma Pigmentosum): UV<sup>s</sup> , multiple skin disorders, skin *cancer*, neurological abnormalities.

XPC(**Rad4**)-XPD(Rad3)-XPE(DDB2)-XPF(Rad1)-XPG(Rad2); XPA (**Rad14**)-XPB (Rad25)

CS (Cockayne's Syndrome): UV<sup>s</sup>, mental and growth *retardation*.

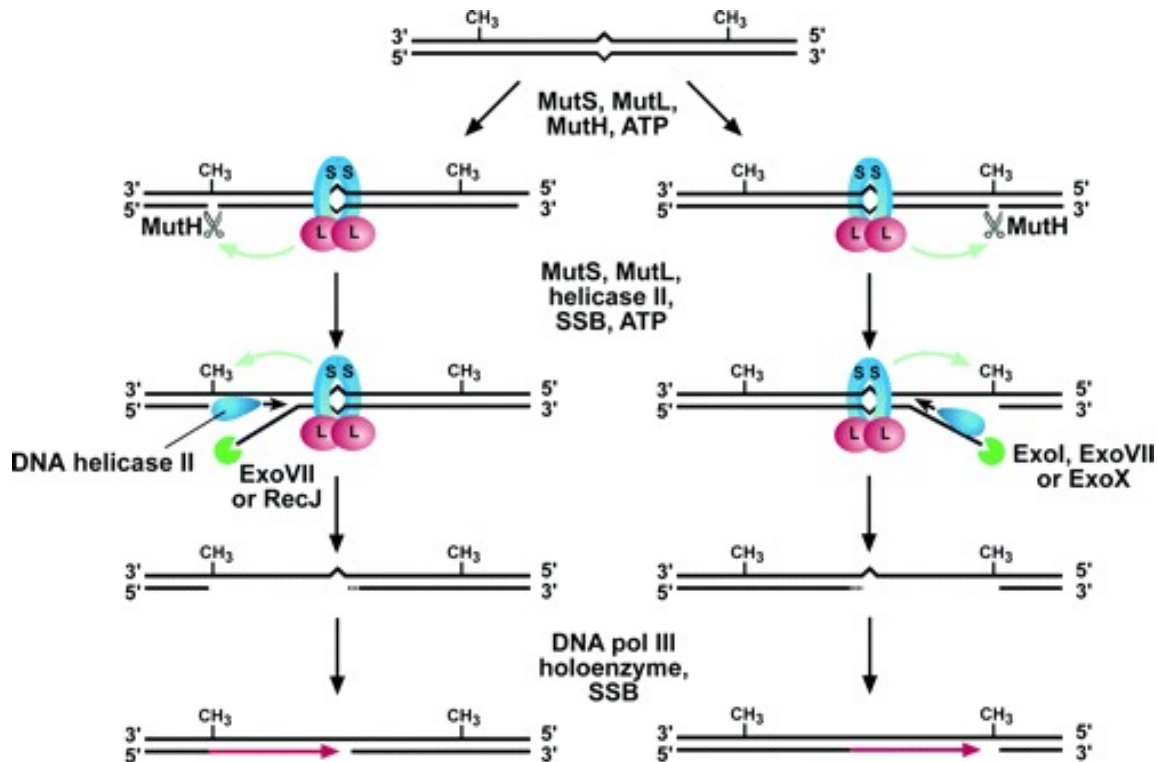
CSA (Rad28), CSB (Rad26) – Transcription coupled NER

TTD (Trichothio-dystrophy): mental and growth retardation, brittle hair.

XPB (Rad25), XPD(Rad3) – TFIIH subunits

Structural and genetic studies to understand the differences among the three syndromes are lacking.

## Mechanism of E. coli methyl-directed MMR



Green arrows = MutS- and MutL-dependent signaling between the 2 DNA sites involved in the reaction.

**Damage Recognized:**  
Base-base mismatch  
Small insertion/deletion loops

### MMR proteins:

- **MutS** (damage recognition)
- **MutL** (recruit of MutH and MutU)
- **MutH** (endonuclease cut at unmethylated strand at GTAC)
- **MutU** (DNA helicase II)
- **Exonucleases** (ExoI, etc)
- **DNA Pol III** -fills in the gap
- **SSB** (Single strand binding protein)
- **DNA Ligase**

# Mismatch Repair in Eukaryotic cells (yeast and human)

**MutS $\alpha$**  (Msh2/Msh6) - recognizes mismatch or 1 bp insertion or deletion

**MutS $\beta$**  (Msh2/Msh3) - recognizes 2-12 bp insertion or deletion

**MutL $\alpha$**  (Mlh1/Pms1) - “match maker” that recruit downstream factor

**Discrimination between parent and daughter strand** may be accomplished by the presence of nick in daughter strands or by ribonucleotides?

Molecular Cell

**Article**

## **Ribonucleotides Misincorporated into DNA Act as Strand-Discrimination Signals in Eukaryotic Mismatch Repair**

Medini Manohar Ghodgaonkar,<sup>1</sup> Federico Lazzaro,<sup>2</sup> Maite Olivera-Pimentel,<sup>1</sup> Mariela Artola-Borán,<sup>1</sup> Petr Cejka,<sup>1</sup> Martin A. Reijns,<sup>3</sup> Andrew P. Jackson,<sup>3</sup> Paolo Plevani,<sup>2</sup> Marco Muzi-Falconi,<sup>2</sup> and Josef Jiricny<sup>1,\*</sup>

## MMR in human diseases

- 
- Defects lead to **Lynch syndrome** or Hereditary Nonpolyposis Colon Cancer (HNPCC). A common cancer predisposition disease. It is nearly always associated with "microsatellite instability" (variations # repeat units of short tandemly repeats).
- MMR affects triplet repeat stabilities implicated in Huntington, fragile X etc.

### **Base Excision Repair**

- removal of oxidative and alkylating damage; also involved in SSB repair.
- damaged bases are removed as free bases.
- intimately linked with the SUMO system.
- have an important role in development and aging.

### **Nucleotide Excision Repair**

- removal of UV-induced damage and bulky adducts & ~ 20% oxidative damage
- damaged bases are removed as oligonucleotides (~20nt)
- Most proteins participate in other cellular processes.
- intimately linked with the ubiquitination system.
- deficient in human disorders (XP, CS and TTD)

### **Mismatch repair**

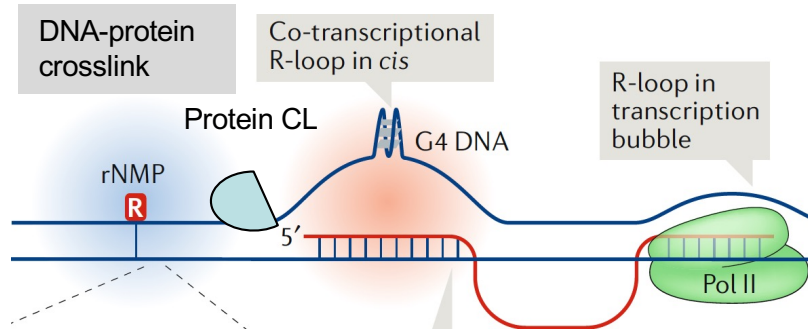
- removal of nucleotide errors or small insertion/deletion generated by DNA polymerases.
- wrong bases are removed by generating a ssDNA gap.
- Most proteins also affect other cellular processes (HR)
- deficient in human disorders: Lynch syndrome and neurodegeneration disorders.

## Many puzzles regarding DNA repair

1. **Mechanisms.** How each step & handover occur accurately; majorities of repair proteins are modified by SUMO, Ub, Phos - how modifications aid repair? New repair proteins are continuing to be discovered.
2. **Chromatin environments:** DNA repair foci - phase-separation; epigenetic markers - silenced vs activate chromatin regions?
3. **Cross-talks:** intimately linked with other cellular functions DNA replication, checkpoints, transcription, innate immunity, metabolism, etc.
4. **Disorders** - How defects in DNA repair processes can lead to different human disorders?



# New DNA repair pathways



## Examples:

- Protein-DNA **crosslink** repair
- RER – **R**ibonucleotide **E**xcision **R**epair
- R-loop repair
- G4 structure removal
- Repair during DNA replication
- Repair during mitosis (MiDAS)

# Connections between DNA repair & human diseases

## Review

### DNA Damage and Cancer Immunotherapy: A STING in the Tale

## Review

CellPress

### DNA damage and innate immunity: links and trade-offs

Georgia Chatzinikolaou<sup>1</sup>, Ismene Karakasilioti<sup>1,2</sup>, and George A. Garinis<sup>1,2</sup>

### Interplay between Cellular Metabolism and the DNA Damage Response in Cancer

## Feature Review

### DNA Damage Triggers a New Phase in Neurodegeneration

# DNA repair, damage tolerance, and signaling

