

Carcinogenesis

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Cancer is a result of a disruption of the normal cellular proliferation.

What are the factors behind this disruption?

Factors are:

- 1. Biological factors**
- 2. Chemical factors**
- 3. Physical factors**

Biological Carcinogens:

1. Viruses

2. Bacteria e.g. *Helicobacter pylori* –*Stomach Cancer*

3. Parasites e.g. *Schistosoma trematodes* –
Bladder Cancer

Viruses

Groups of viruses: DNA and RNA groups

1.DNA viruses which include

- a - Papilloma viruses –Urogenital cancer,Larynx,Cervical
8Kb+10 overlapping genes+E5,E6 are transforming genes.
- b – Adeno viruses ???
- c - Hepatitis viruses\ Hepatocellular carcinoma-
Hepatoma
- d - Herpes viruses
 - Epstein-Barr \ Burkitts lymphoma(t(8 to 2,22,14) +
Nasopharyngeal carcinoma
 - Cytomegalovirus\ Kaposi's sarcoma, ??Prostate
cancer, Colon cancer, Cervical carcinoma
 - Herpes 1 &2 ??

2.RNA viruses which include:

- a - Complete transforming viruses- v-Oncs, High Oncogenic + induce tumor rapidly at high frequency.
- b - Slow transforming viruses- Weak LTR
- c - Defective transforming viruses- v-Oncs+ another virus

3. Viral Cycles

- a. Virulent or lytic cycle:
New virus particles are made and released-burst- from the host cell.

- b. lysogeny cycle or integration cycle

Viral Carcinogenesis Mechanisms

a. DNA viruses

--T-antigens, large, middle and small;
LTR and E genes

Functions:

- 1.Necessary for host cell transformation
- 2.Stimulates the host cell to replicate its DNA
- 3.Binds to cellular DNA
- 4.Binds to p53 protein

b. RNA viruses

- v-Oncogenes--- gag-pol-env-onc or LTR
- Rous sarcoma virus+ src gene
- avian leukemia virus+ LTR
- MC29 virus + Myc oncogene
- Harvey sarcoma virus+ ras genes
- Mouse sarcoma virus + mos gene
- St-Fe SV + fes gene

Chemical Carcinogenesis

- Sources**
- Ames test**

Group of chemicals

a. Basis analogs

-5-bromouracil , 2-amino purine

b. Nitrous acid group

c. Alkylating agents

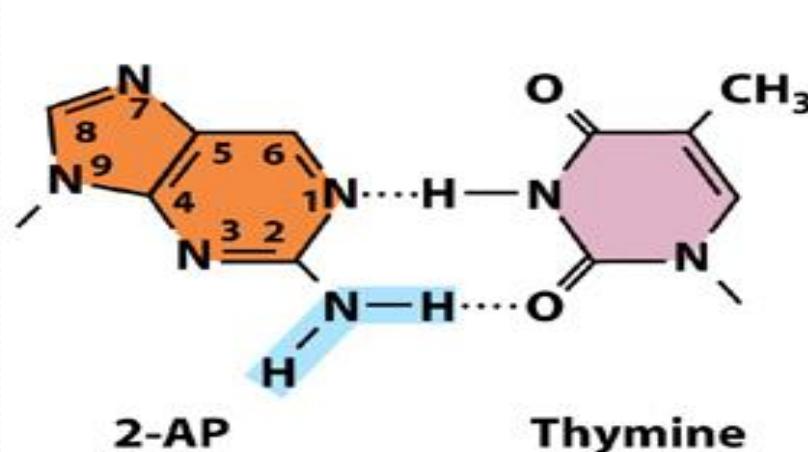
d. Acridine dyes

A. Basis analogs .a

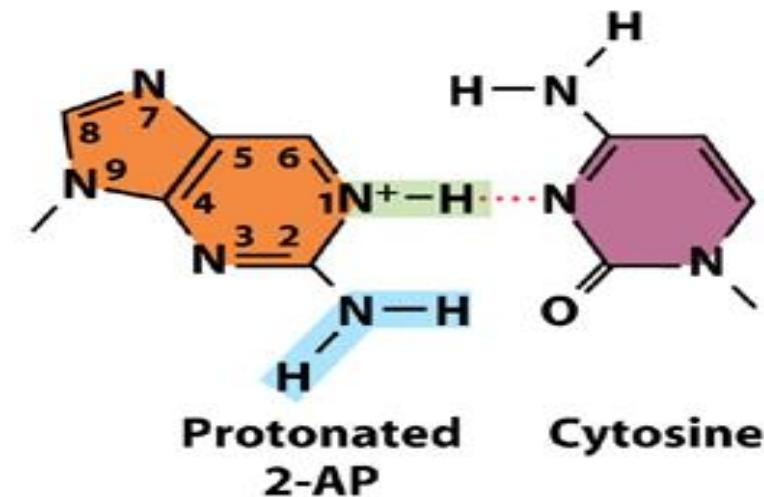
5-bromouracil ...analogue to U..T..C

2-amino purine...analogue to G..A

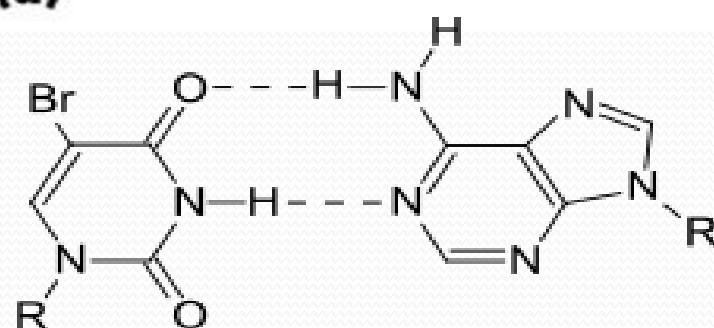
.b



(a)

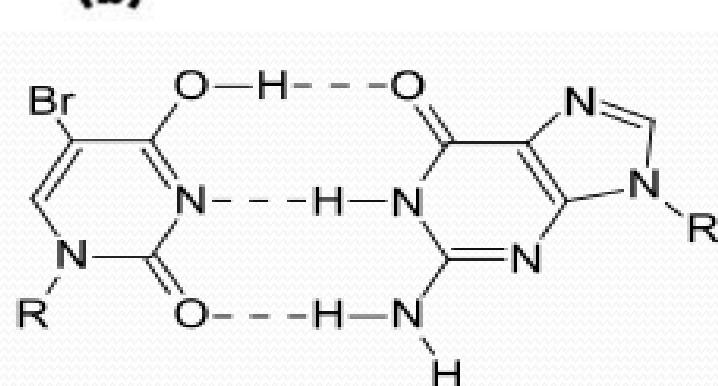


(b)



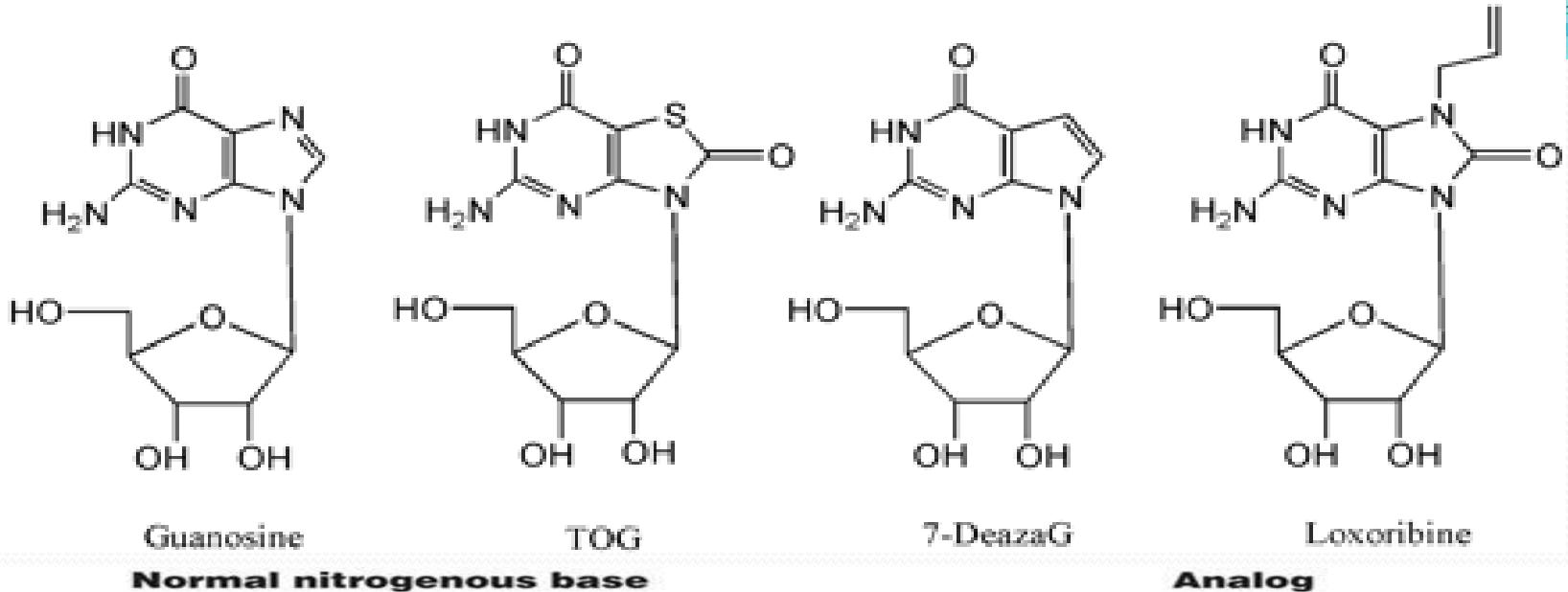
5-BrU (keto)

Adenine



5-BrU (enol)

Guanine



Guanosine

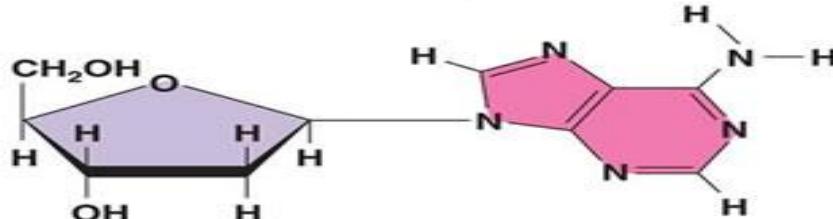
TOG

7-DeazaG

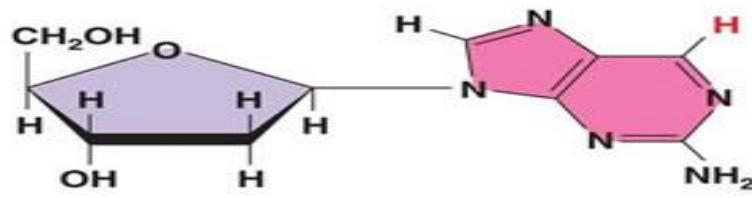
Loxoribine

Normal nitrogenous base

Analog

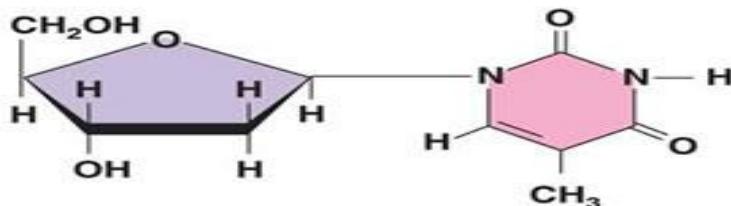


Adenine nucleoside

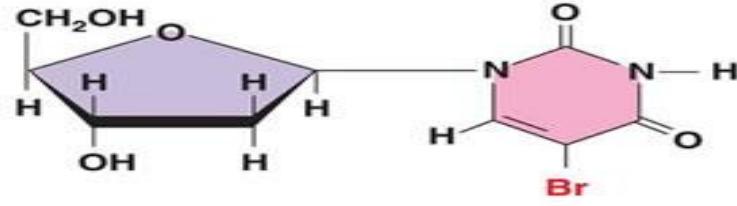


2-Aminopurine nucleoside

(a) The 2-aminopurine is incorporated into DNA in place of adenine but can pair with cytosine, so an AT pair becomes a CG pair.



Thymine nucleoside



5-Bromouracil nucleoside

(b) The 5-bromouracil is used as an anticancer drug because it is mistaken for thymine by cellular enzymes but pairs with cytosine. In the next DNA replication, an AT pair becomes a GC pair.

Original base

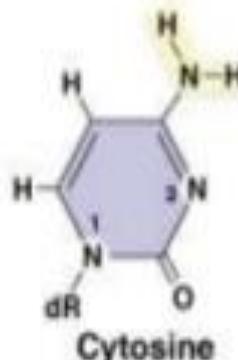
Mutagen

Modified base

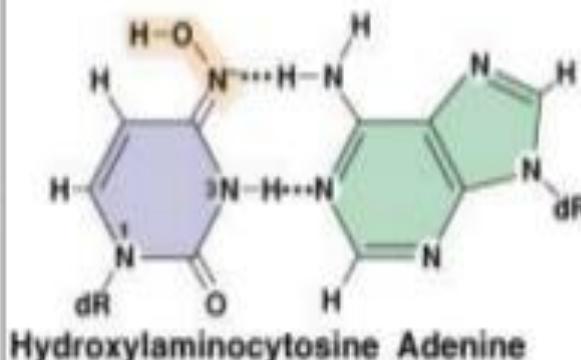
Pairing partner

Predicted transition

b)

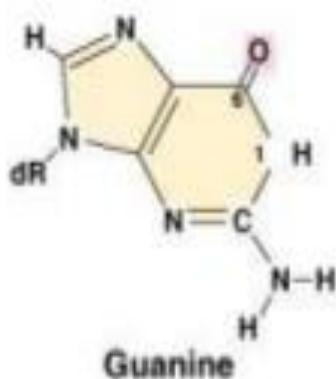


Hydroxylamine
(NH_2OH)

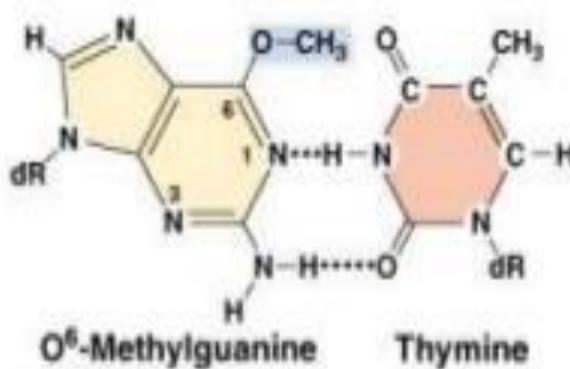


$\text{CG} \longrightarrow \text{TA}$

c)



Methylmethane
sulfonate (MMS)
(alkylating agent)



$\text{GC} \longrightarrow \text{AT}$

b. Nitrous acid group

Oxidation the bases to remove amine group of Carbone atom 6

-Adenine change to hypoxanthine to bind to cytosine instead of thymine
-Cytosine change to uracil to bind to adenine instead of guanine
-Guanine change to xanthine to bind to cytosine

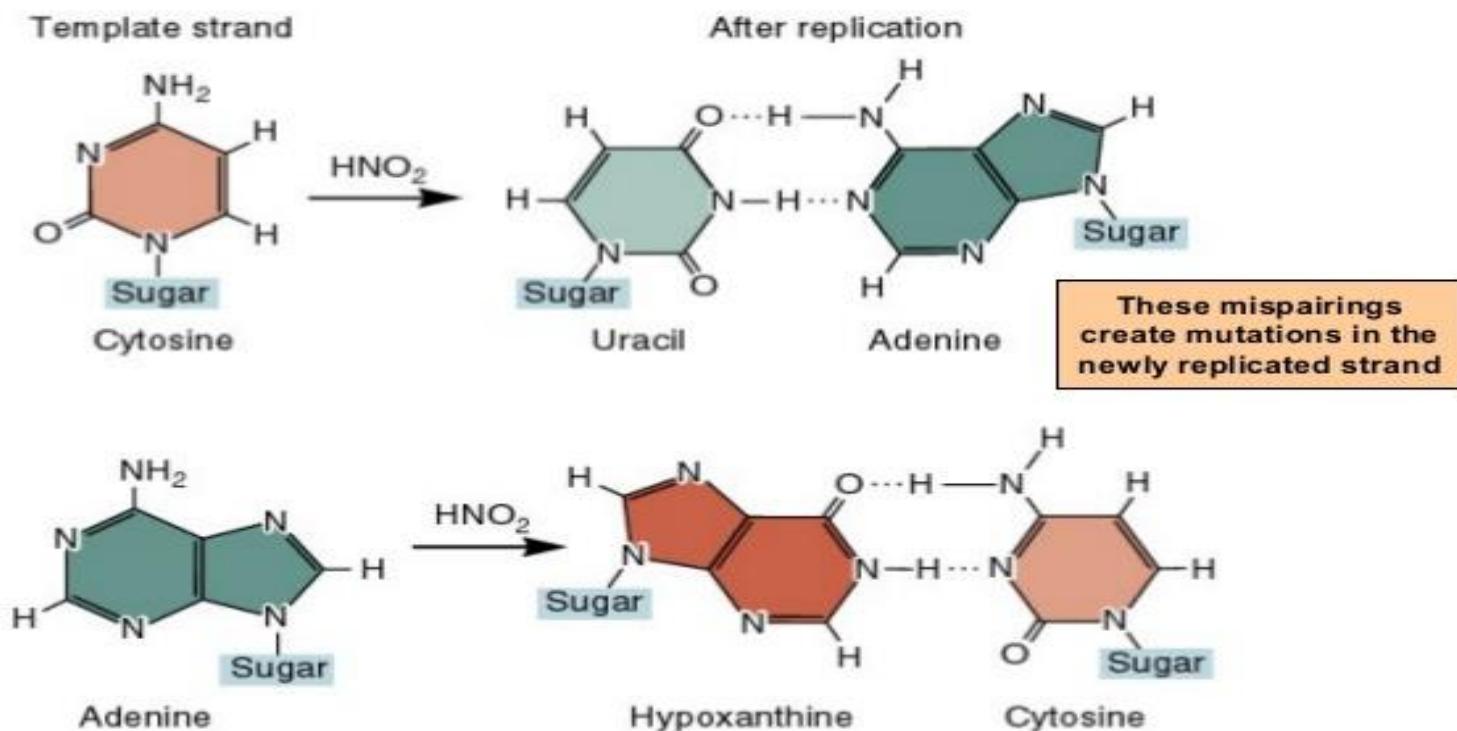
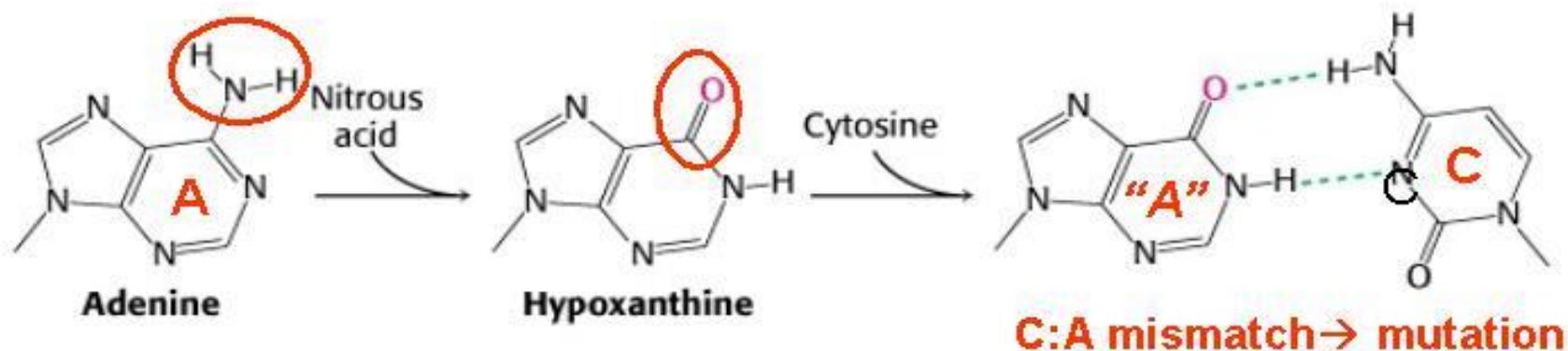


Figure 16.13 Mispairing of modified bases

Chemical Mutagen : Nitrous acid (HNO_2)

Deamination causes A:T to G:C transitions



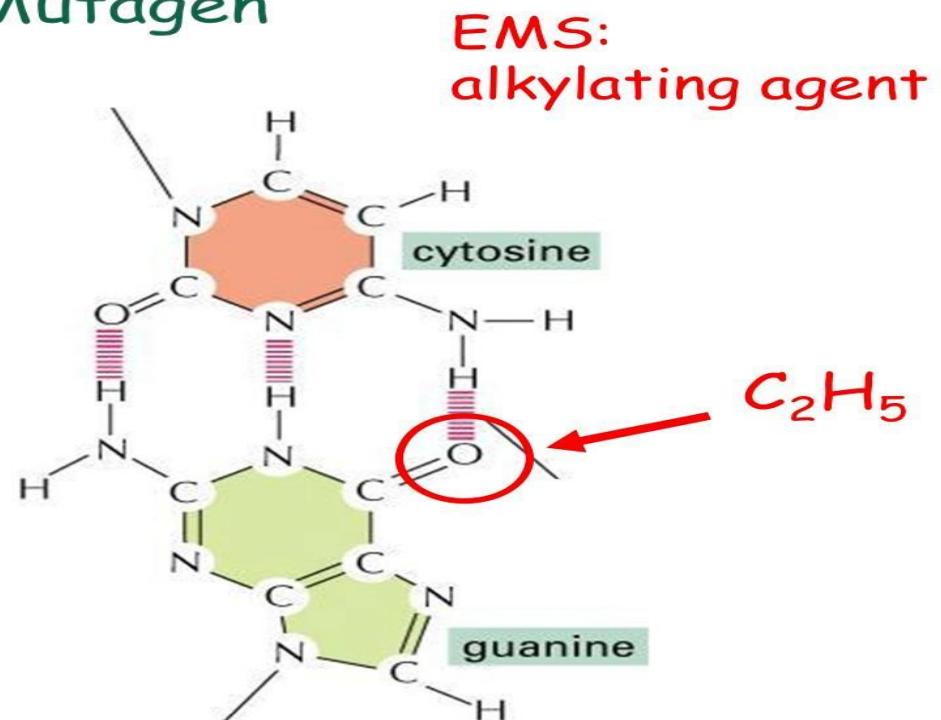
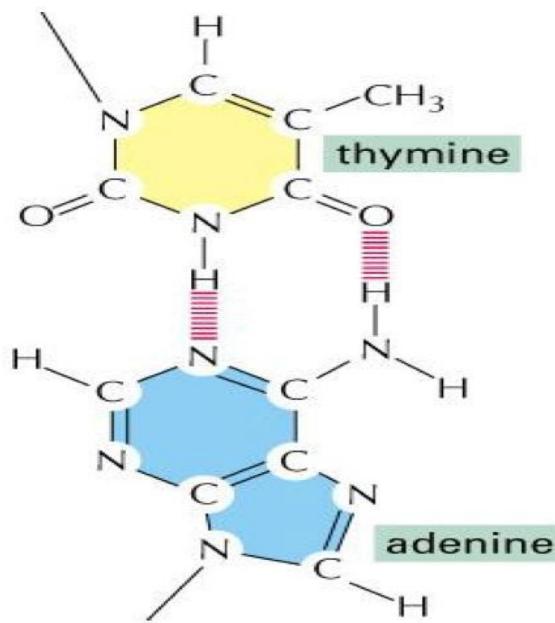
* **HNO_2 also deaminates C to U:
causes G:C to A:T transitions**

c. Alkylating agents

Mustard gas, nitrogen gas, ethyl ethan sulfonate-EES,
ethyl methan sulfonate-EMS

Adding ethyl or methyl groups –CH₃,CH₂- to DNA base

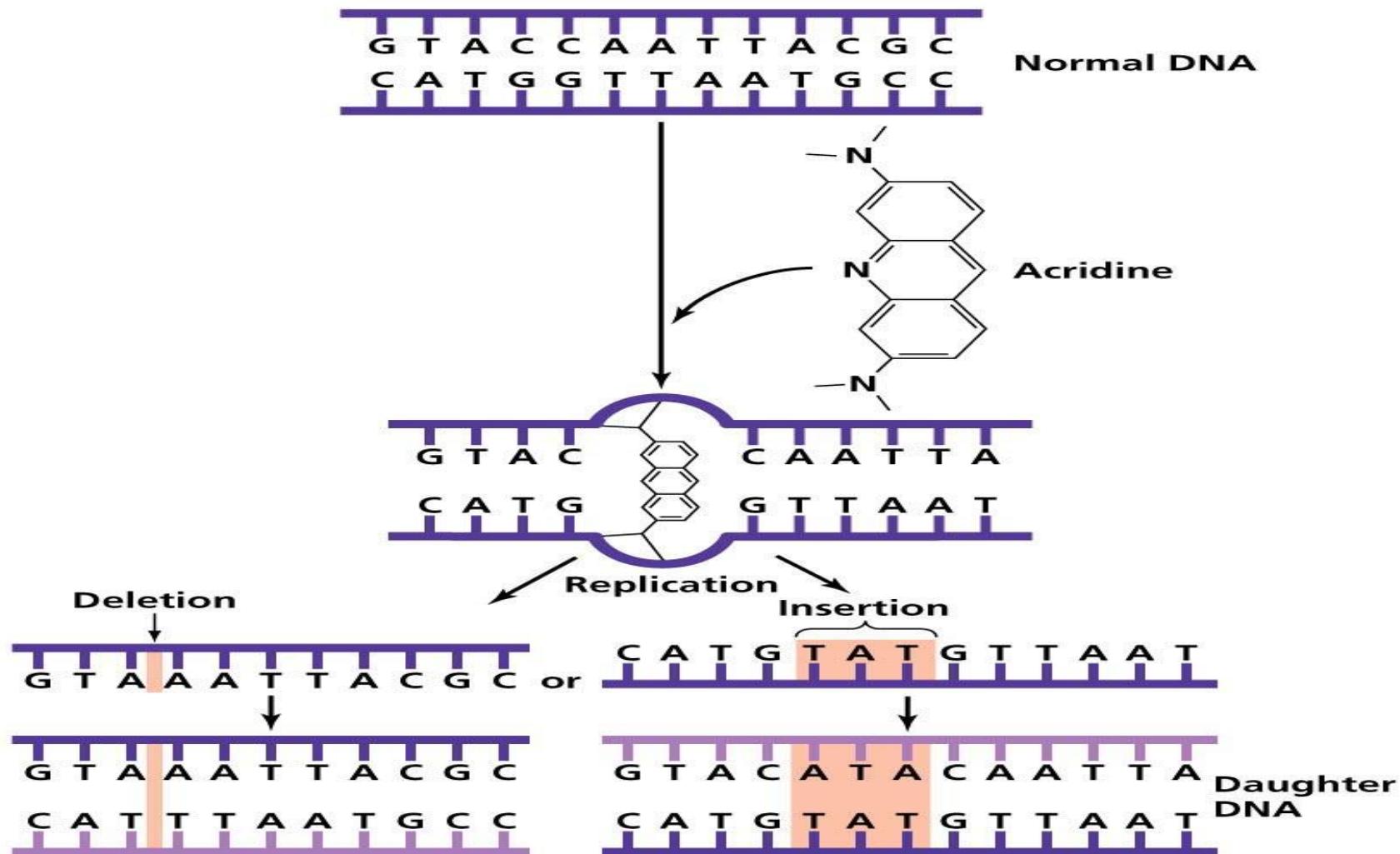
Induced mutations - Mutagen



alkylated G prefers to bind T

d. Acridine dyes

Cause DNA instability, Framshift mutation



Chemical agents (III)

- **Intercalating agents or acridine dyes**
 - * Proflavin, acridine orange, ICR-170, ICR-191
 - * Ethidium bromide
 - * แทรกระหว่างโครงสร้าง DNA → interfere DNA replication
 - * Frameshift mutation