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About the course: www.life.umd.edu/classroom/BSCI410-Liu/BSCI410/

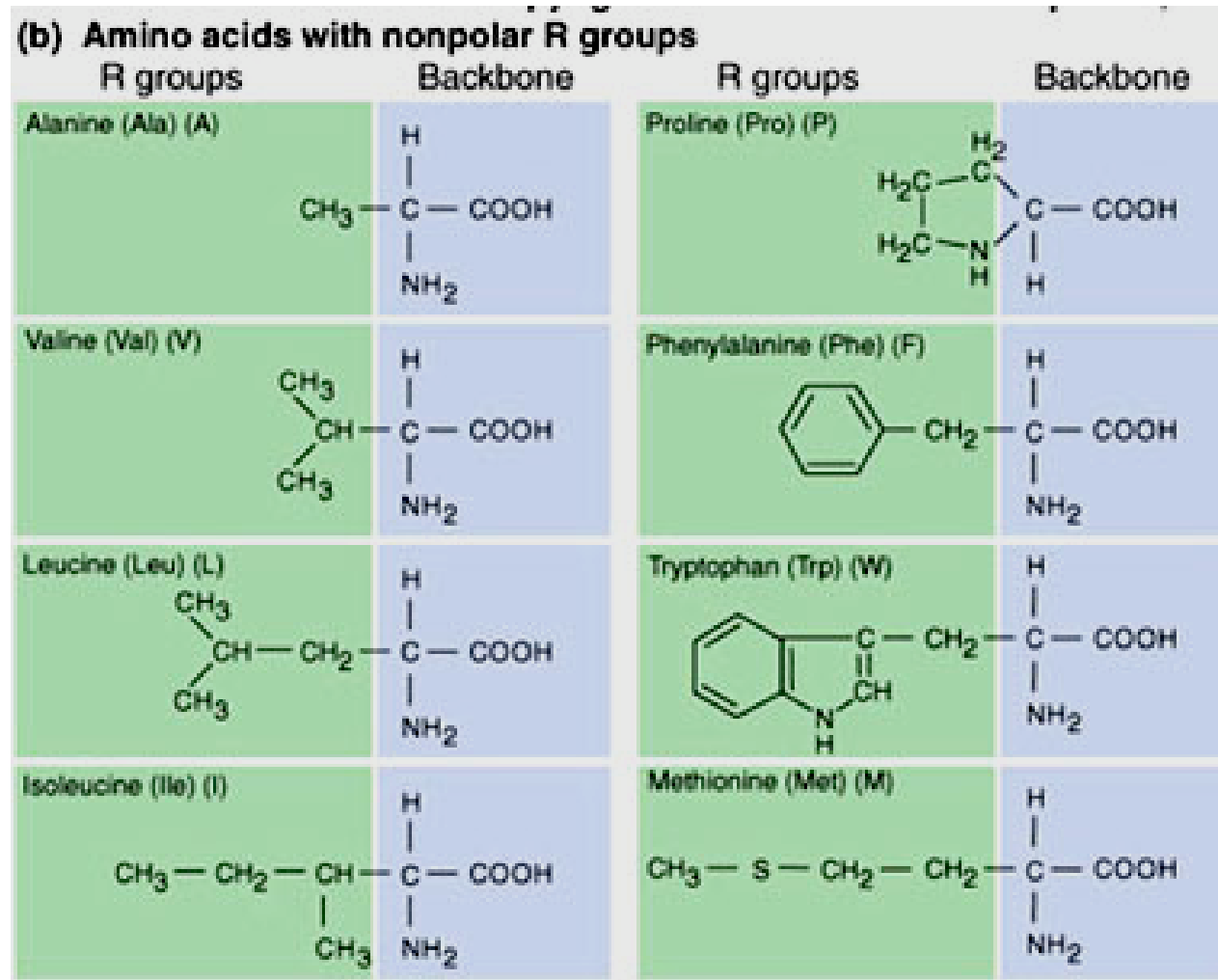
Lecture 2: Mutation and its effect

- Mutation type
- Mutational effect
- Spontaneous Mutation
- Mutagens

Read: Ch 7 p192-193; 196-198; 200-205

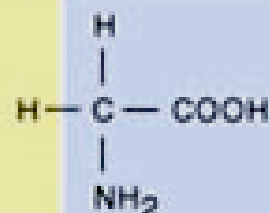
Figs: 7.2; 7.6; 7.7; 7.8; 7.12; 7.21;
7.22; 8.15; 8.16;

Fig. 7.21

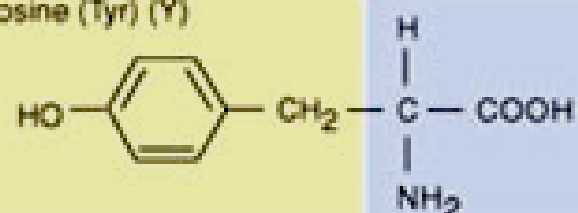


Amino acids with uncharged polar R groups

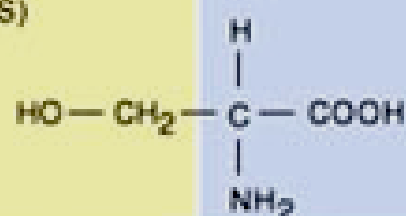
Glycine (Gly) (G)



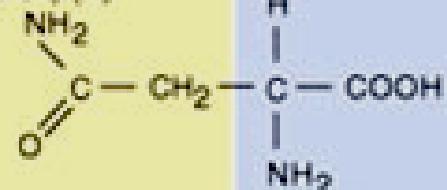
Tyrosine (Tyr) (Y)



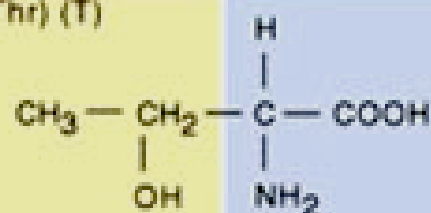
Serine (Ser) (S)



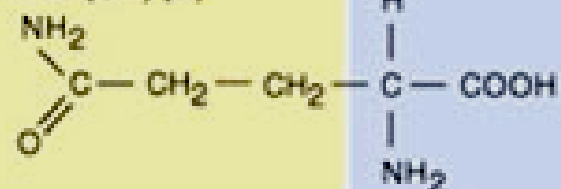
Asparagine (Asn) (N)



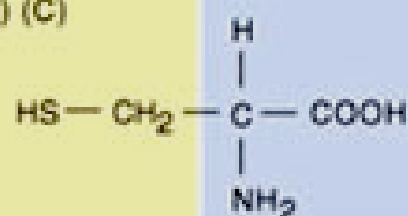
Threonine (Thr) (T)



Glutamine (Gln) (Q)

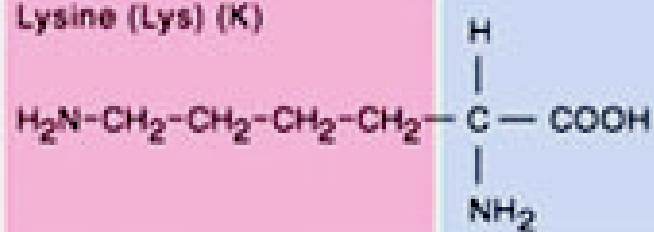


Cysteine (Cys) (C)

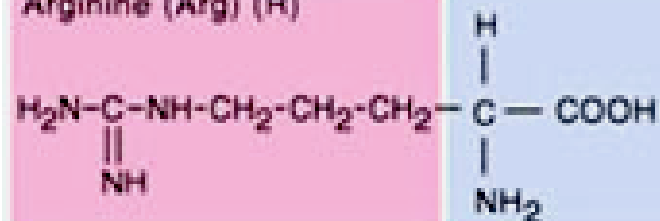


Amino acids with basic R groups

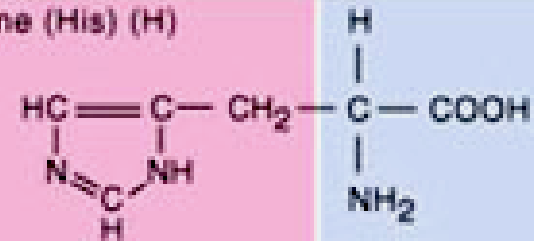
Lysine (Lys) (K)



Arginine (Arg) (R)

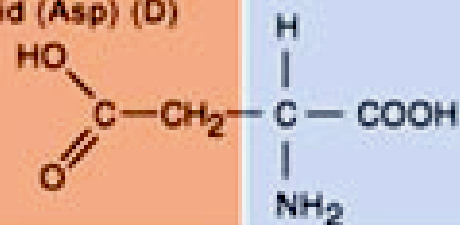


Histidine (His) (H)

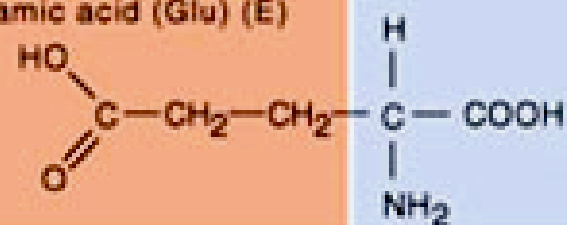


Amino acids with acidic R groups

Aspartic acid (Asp) (D)



Glutamic acid (Glu) (E)



Mutations

1. Substitution-1 base --> one of the three other bases

Transition: purine --> purine or pyrimidine --> pyrimidine

A--> G or G--> A T--> C or C--> T

Transversion: purine --> pyrimidine or *vice versa*

A--> T, C; G -->T,C; T-->A, G; C-->A,G

causes **missense, nonsense, silent, neutral** or
splicing mutational effects

2. Deletion or insertion-often causes **frameshift** mutation

3. Chromosomal rearrangement

inversion or translocation can change multiple genes

Effects of point mutations

tyrosine TAT, TAC

TAT -> CAT	tyr -> his	missense
TAT -> TAA	tyr -> stop	nonsense
TAT -> TTT	tyr -> phe	neutral in many cases
TAT -> TAC	tyr -> tyr	silent

Fig. 7.22

(a) From mutation to phenotype

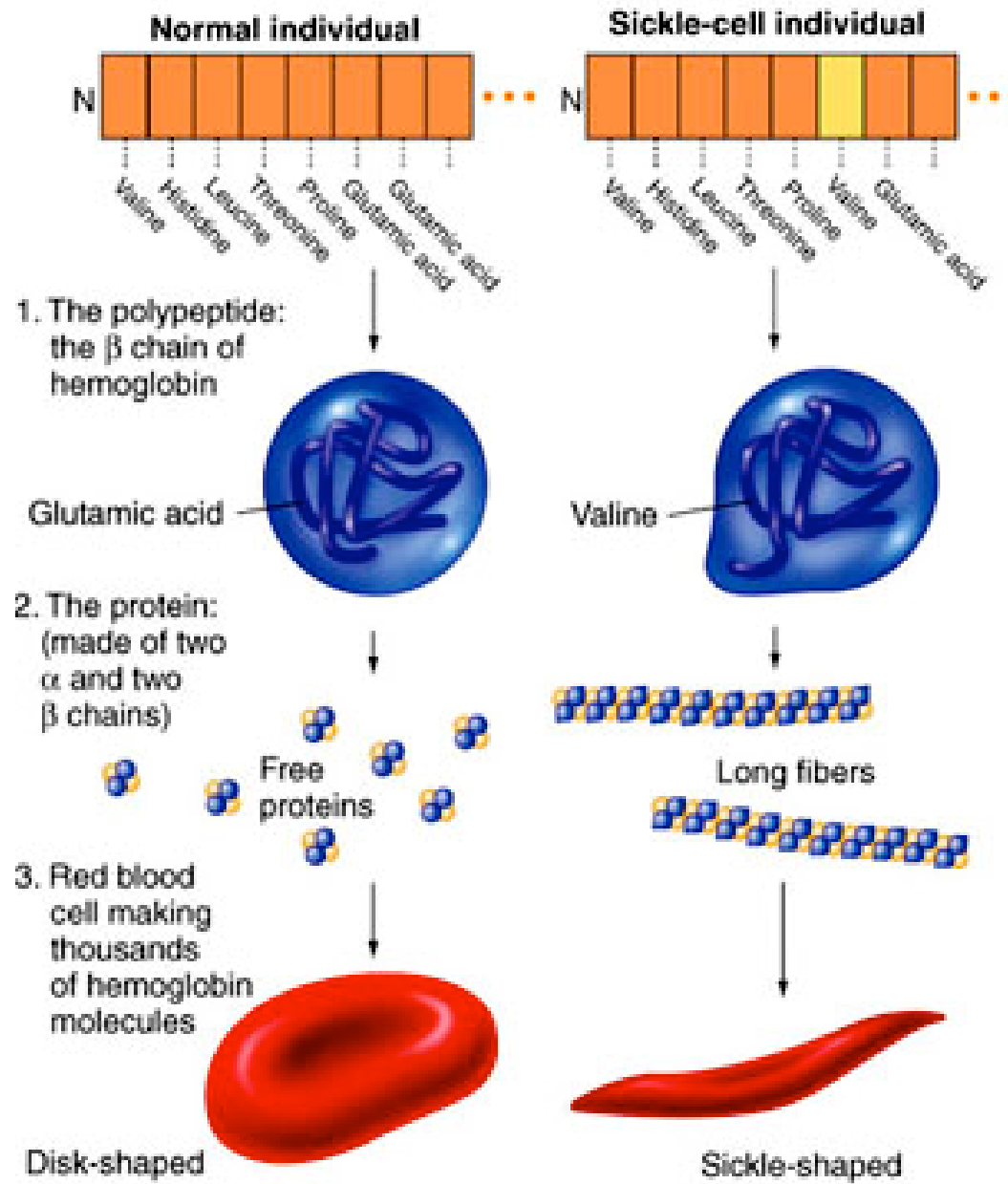


Fig. 8.15

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Splicing removes introns from a primary transcript.

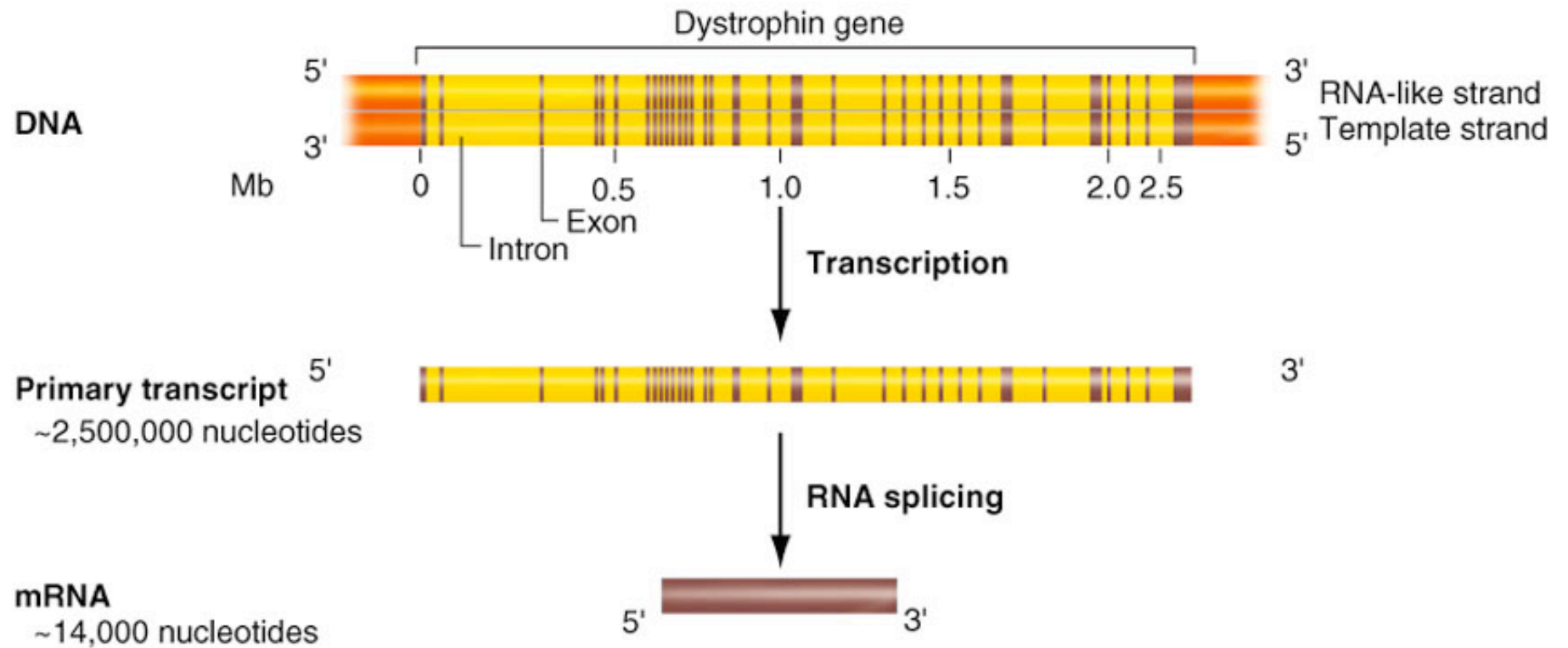
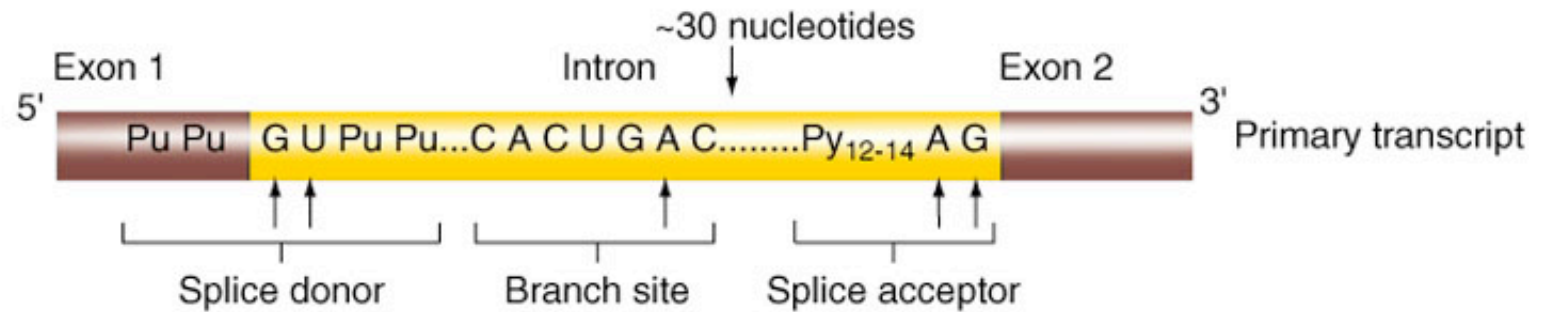


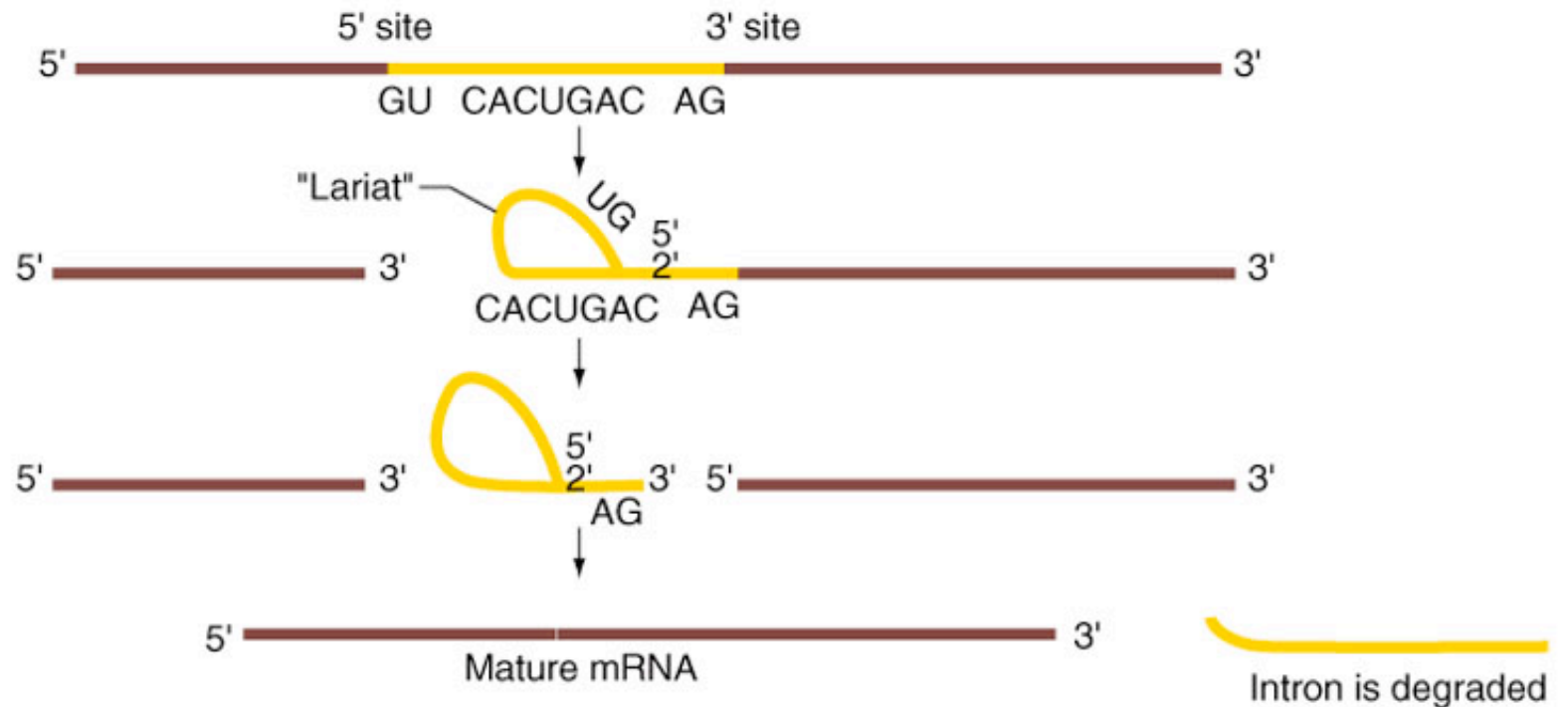
Fig. 8.16

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(a) Short sequences dictate where splicing occurs.

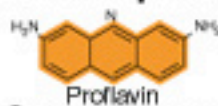


(b) Two sequential cuts remove the intron.



Frameshift mutations

(a) The mutagen proflavin can insert between two base pairs



Molecule of proflavin inserted between stacked base pairs

(b.1) Consequences of exposure to proflavin

rII⁺ Wildtype

Exposure to proflavin

FC0

rII⁻

Exposure to proflavin

FC0 FC7

rII⁺ revertant

Original mutation

Second mutation

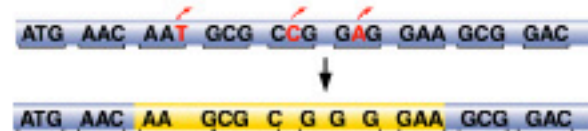
(b.2) Crossing *rII*⁺ revertant with wildtype yields *rII*⁻ recombinants



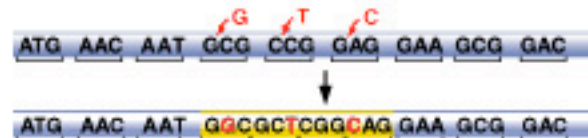
(c) Different sets of mutations generate either a mutant or a normal phenotype

Proflavin-induced mutations (+) insertion (-) deletion	Phenotype
- or +	Mutant
-- or ++	Mutant
---- or ----	Mutant
- +	Wildtype
--- or ---- or +++ or +++++	Wildtype

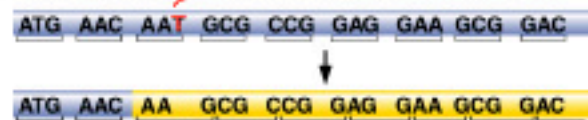
(d) Three single base deletions (---)



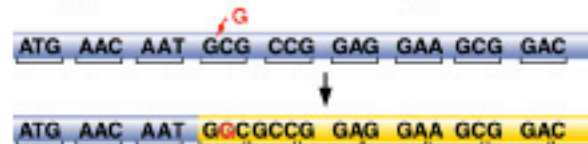
Three single base insertions (+++)



(e) Single base deletion (-)



Single base insertion (+)



■ correct triplet
■ incorrect triplet

Fig. 7.2

Type of mutation and effect on base sequence

(a) Substitution

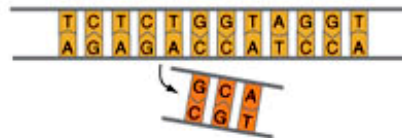
Transition: Purine for purine, pyrimidine for pyrimidine



Transversion: Purine for pyrimidine, pyrimidine for purine



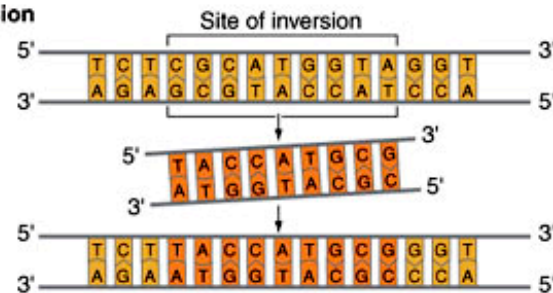
(b) Deletion



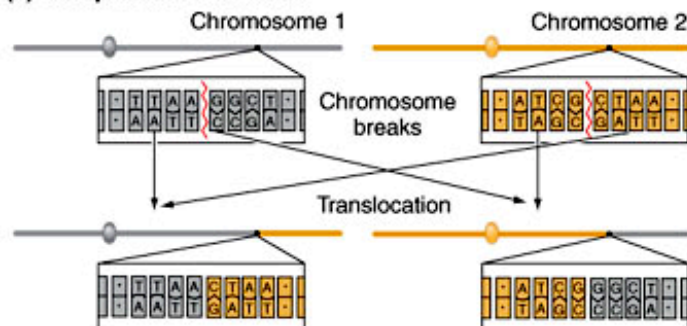
(c) Insertion



(d) Inversion



(e) Reciprocal translocation



Spontaneous mutations

Spontaneous mutation is rare: $2-12 \times 10^{-6}$ (per generation per gene)

Spontaneous mutations can be caused by

a. mistakes made during DNA replication (error rate 10^{-9})

b. environmental effect:

UV light: thymidine dimer

X-ray: break sugar-phosphate DNA back bone

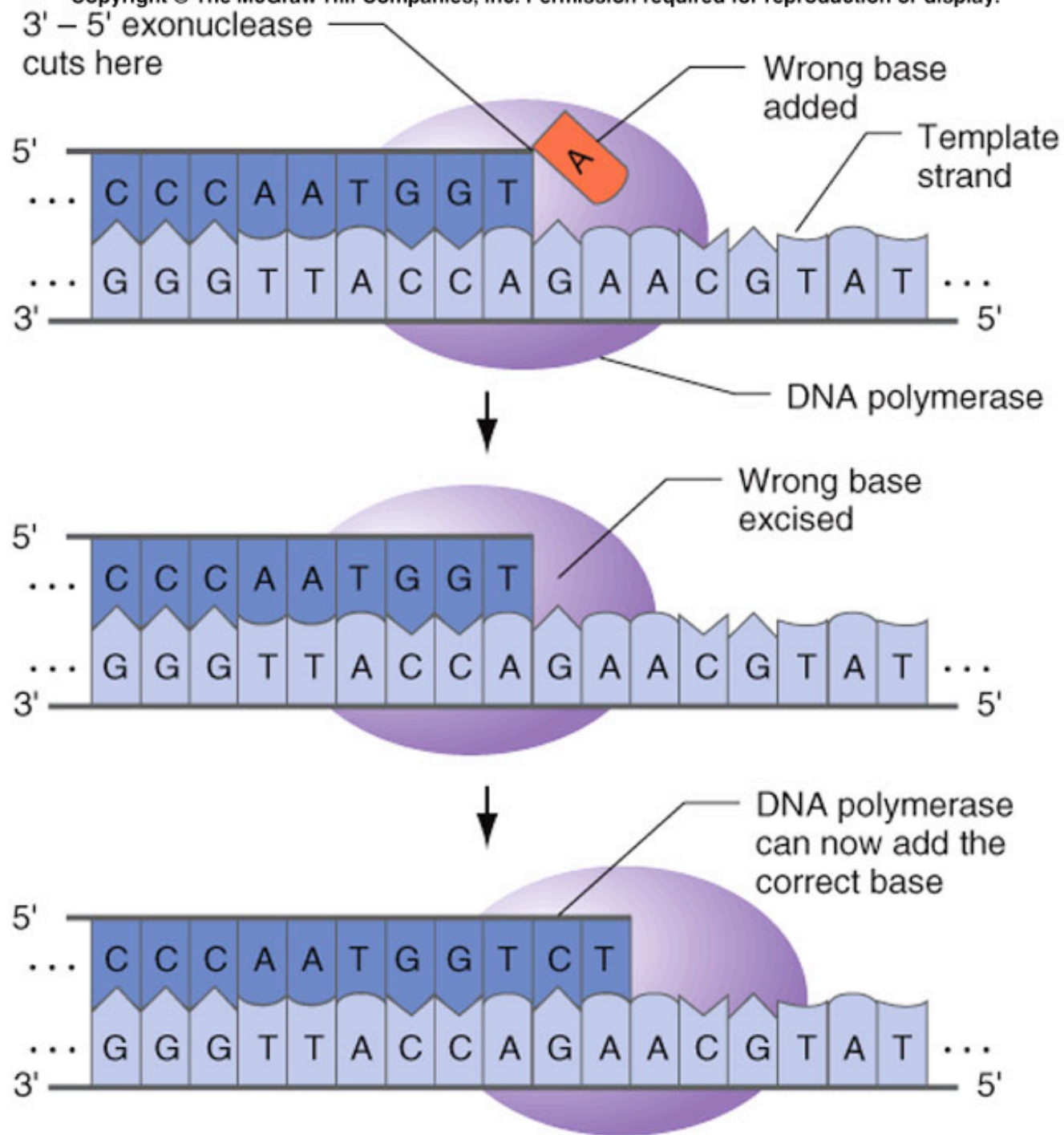
Oxidative damages: $G \rightarrow 8\text{-oxodG}$ (pair with A)

c. chemical changes (hydrolysis):

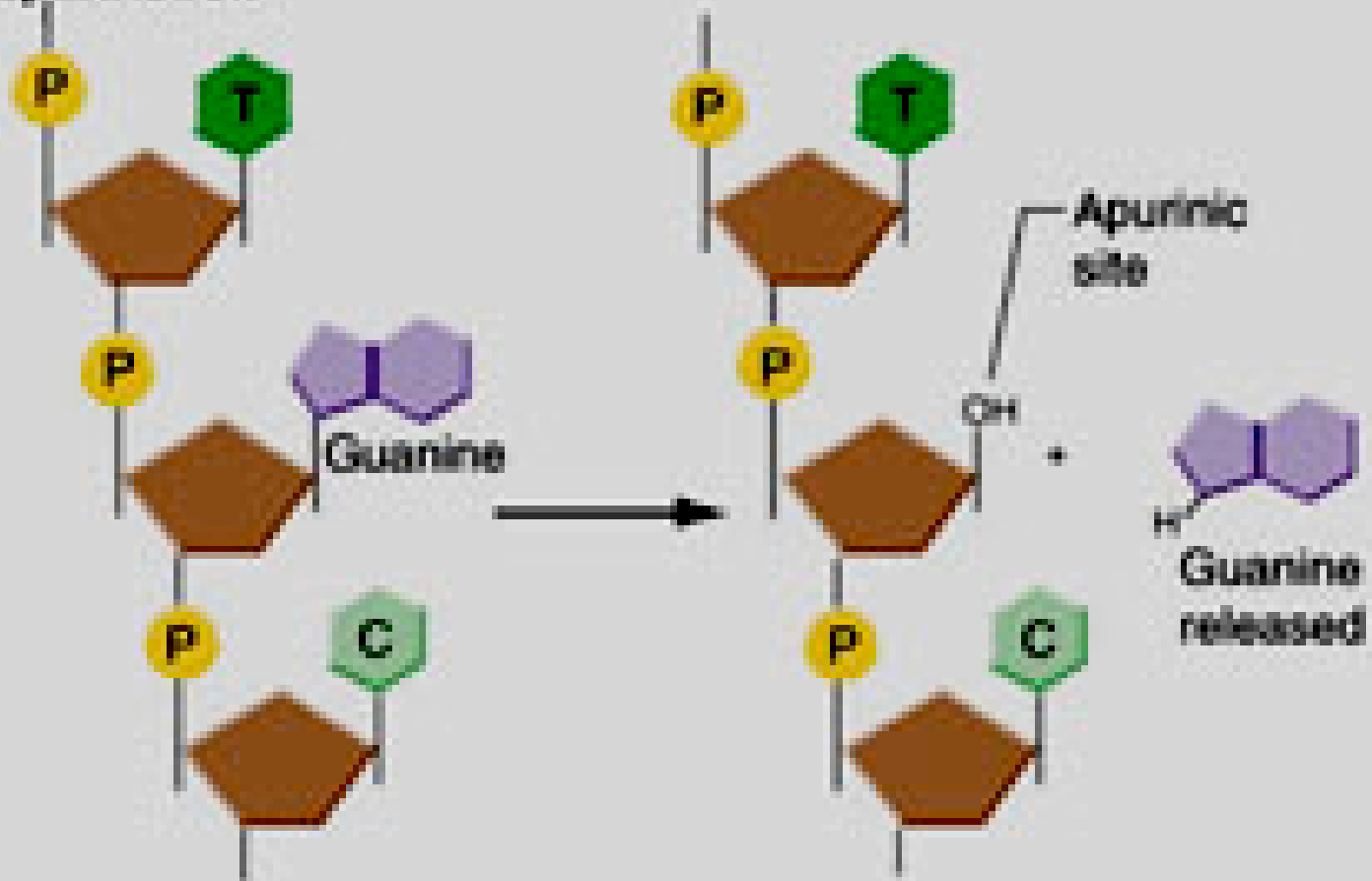
depurination; $A, G \rightarrow O$

deamination: $C \rightarrow U$

Fig. 7.8

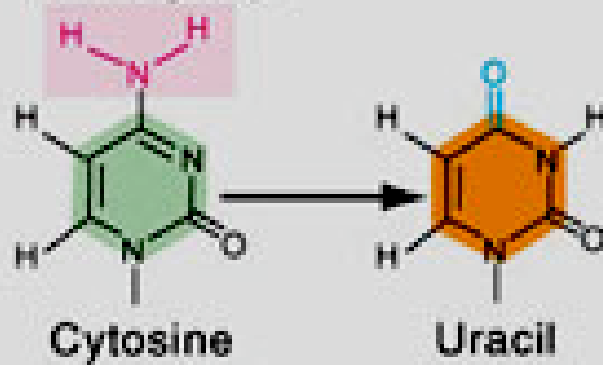


(a) Depurination

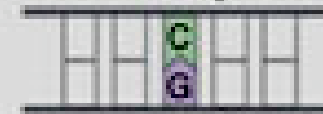


(b) Deamination

Amino group



Normal sequence



↓ Deamination



Replication

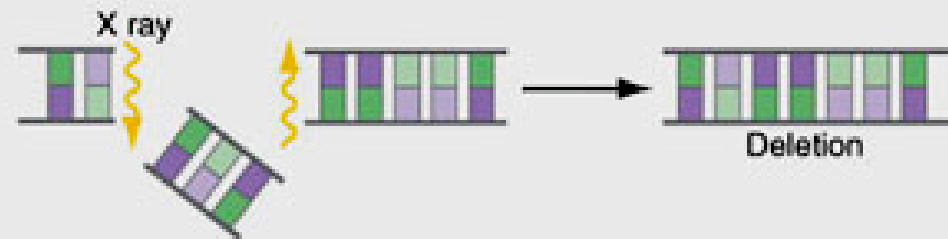


Replication

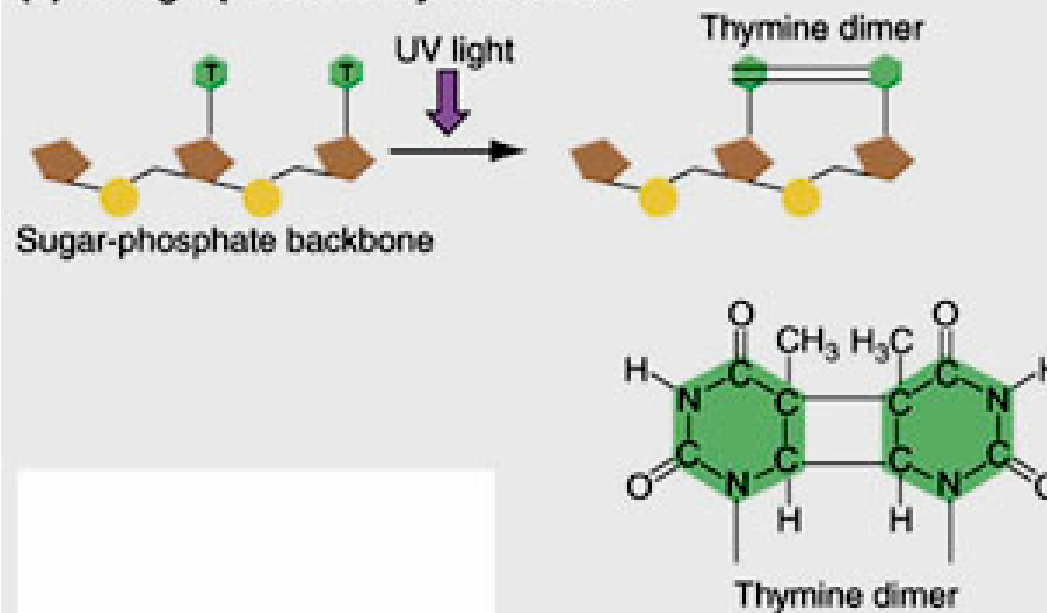


Mutant sequence

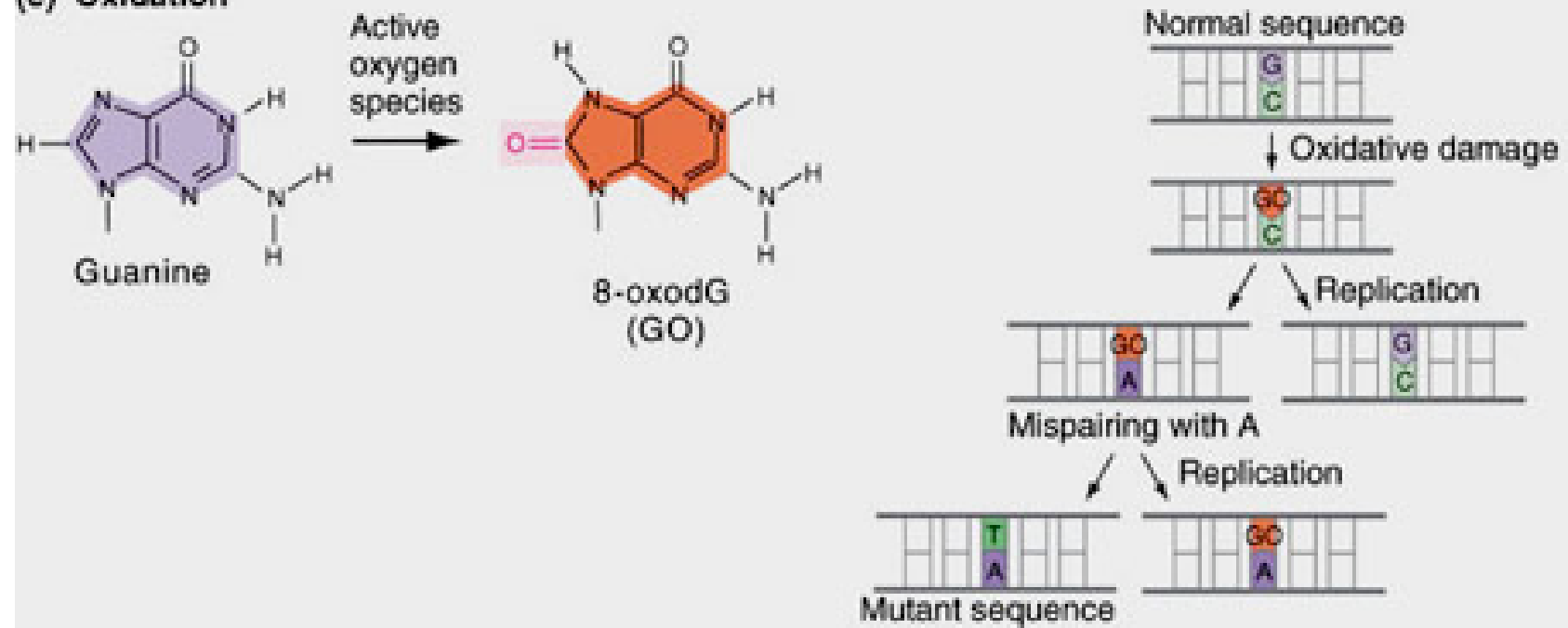
(c) X rays break the DNA backbone



(d) UV light produces thymine dimers



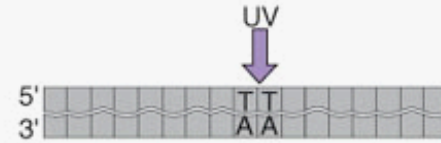
(e) Oxidation



(a) Excision repair

Fig. 7.7

1. Exposure to UV light.



2. Thymine dimer forms.



3. Endonuclease nicks strand containing dimer.



4. Damaged fragment is released from DNA.



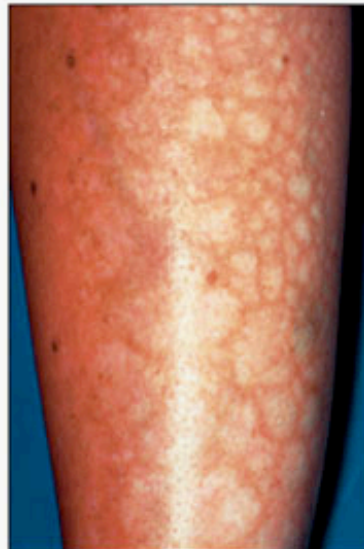
5. DNA polymerase fills in the gap with new DNA (yellow).



6. DNA ligase seals the repaired strand.



(b) Xeroderma pigmentosum



Mutagens

Mutagen treatment greatly increases the mutation rate

Exposure to X-ray, UV light

Chemical treatment: base analogs 5'-bromouracil (=T or rarely C)

hydroxylating agent (add OH-group to C)

alkylating agent such as EMS (ethylmethane sulfonate)

deaminating agent such as nitrous acid

intercalating agent such as Acridine Orange

Transposons that insert into a gene and disrupt the normal reading frame

Chemical Mutagens

Fig. 7.12a1

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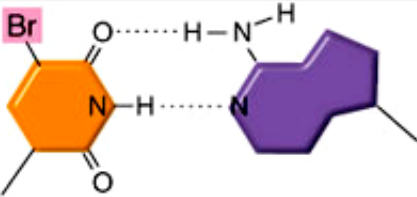
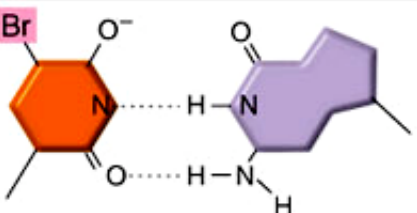
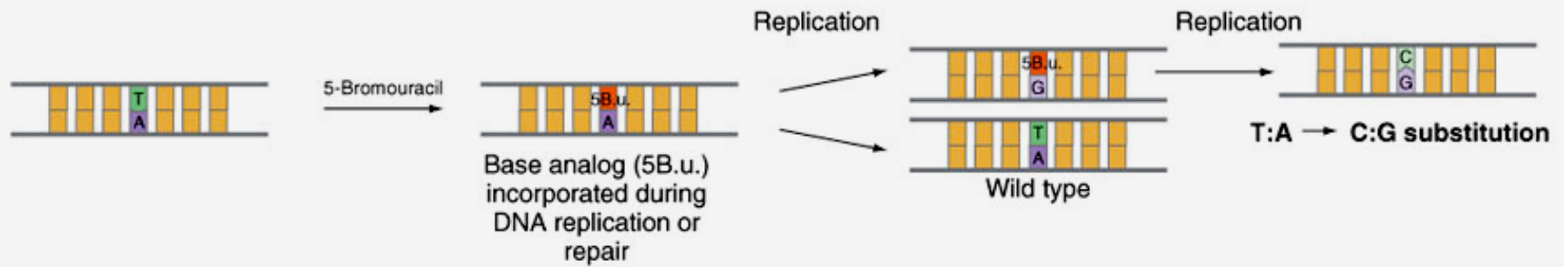
Type of mutagen	Chemical action of mutagen	
(a) Replace a base: Base analogs have a chemical structure almost identical to that of a DNA base.	 <p>5-Bromouracil—normal state, behaves like thymine</p> <p>Adenine</p>	 <p>5-Bromouracil—rare state, behaves like cytosine</p> <p>Guanine</p>
	5-Bromouracil: almost identical to thymine. Normally pairs with A; in transient state, pairs with G.	

Fig. 7.12a2

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How mutagens induce mutations



Type of mutagen	Chemical action of mutagen
(b) Alter base structure and properties: <i>Hydroxylating agents:</i> add a hydroxyl (–OH) group	<p style="text-align: center;">Cytosine $\xrightarrow{\text{Hydroxylamine}}$ N-4-Hydroxycytosine (C*) Adenine</p> <p style="text-align: center;">Hydroxylamine adds –OH to cytosine; with the –OH, hydroxylated C now pairs with A instead of G.</p>
<i>Alkylating agents:</i> add ethyl (–CH ₂ –CH ₃) or methyl (–CH ₃) groups	<p style="text-align: center;">Guanine $\xrightarrow{\text{Ethylmethane sulfonate}}$ O-6-Ethylguanine (G*) Thymine</p> <p style="text-align: center;">Ethylmethane sulfonate adds an ethyl group to guanine or thymine. Modified G pairs with T above, and modified T pairs with G (not shown).</p>
<i>Deaminating agents:</i> remove amine (–NH ₂) groups	<p style="text-align: center;">Cytosine $\xrightarrow{\text{Nitrous acid}}$ Uracil Adenine</p> <p style="text-align: center;">Adenine $\xrightarrow{\text{Nitrous acid}}$ Hypoxanthine Cytosine</p> <p style="text-align: center;">Nitrous acid modifies cytosine to uracil, which pairs with A instead of G; modifies adenine to hypoxanthine, a base that pairs with C instead of T.</p>

Fig. 7.12b1

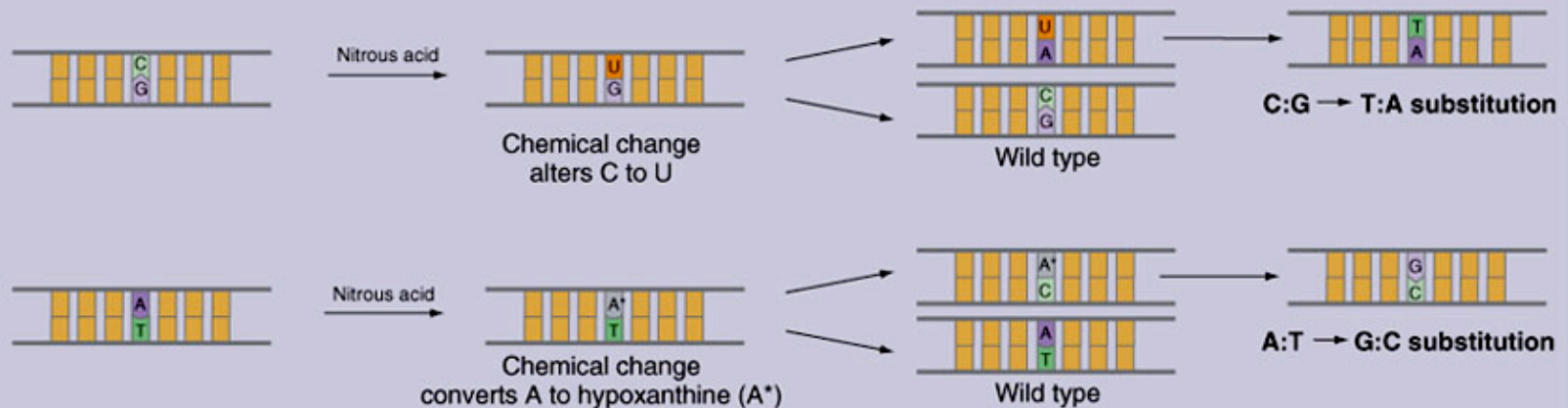
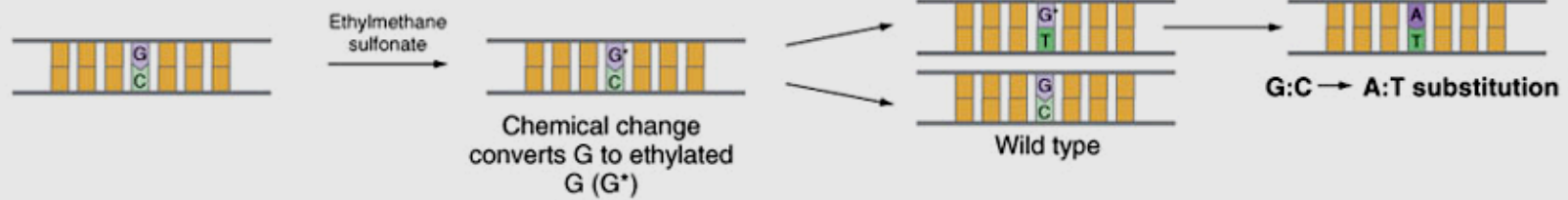
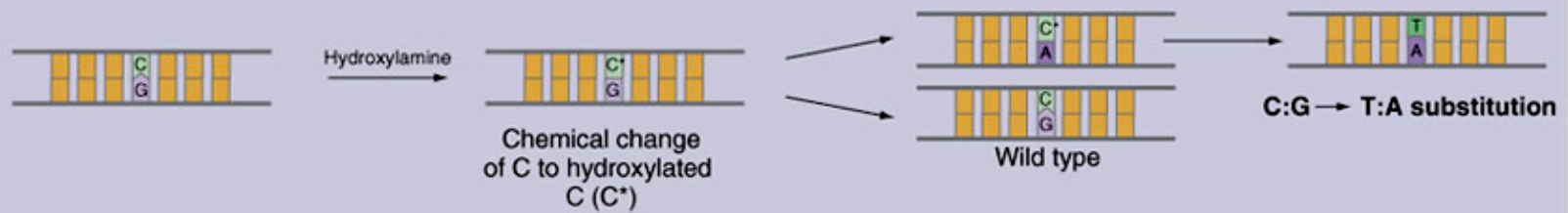


Fig. 7.12c1

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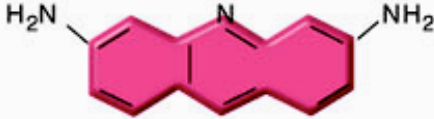
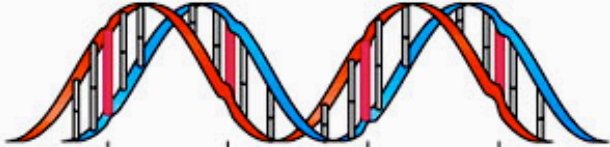
Type of mutagen	Chemical action of mutagen
(c) Insert between bases: Intercalating agents	<div data-bbox="653 630 1083 748"></div> <div data-bbox="821 776 936 805">Proflavin</div> <div data-bbox="1171 610 1776 756"></div> <div data-bbox="1339 776 1608 837">Intercalated proflavin molecules</div> <p data-bbox="598 870 1835 932">Proflavin intercalates into the double helix. This disrupts DNA metabolism, eventually resulting in deletion or addition of a base pair.</p>

Fig. 7.12c2

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