

# BIOMARKERS AND TOXICITY MECHANISMS 05 – Mechanisms - DNA

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Tento projekt je spolufinancován Evropským sociálním fondem a státním rozpočtem České republiky.









INVESTICE DO ROZVOJE VZDĚLÁVÁNÍ

# DNA

- principal molecule for life
- structure and function carefully checked
- changes rapidly repaired
- irreversible changes → cell death (physiologically by apoptosis)

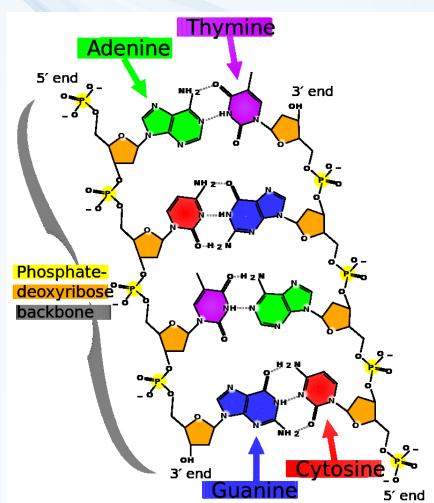
#### Mutagenesis → MUTATIONS

→ variability and evolution or → damage to DNA (structure or coding)

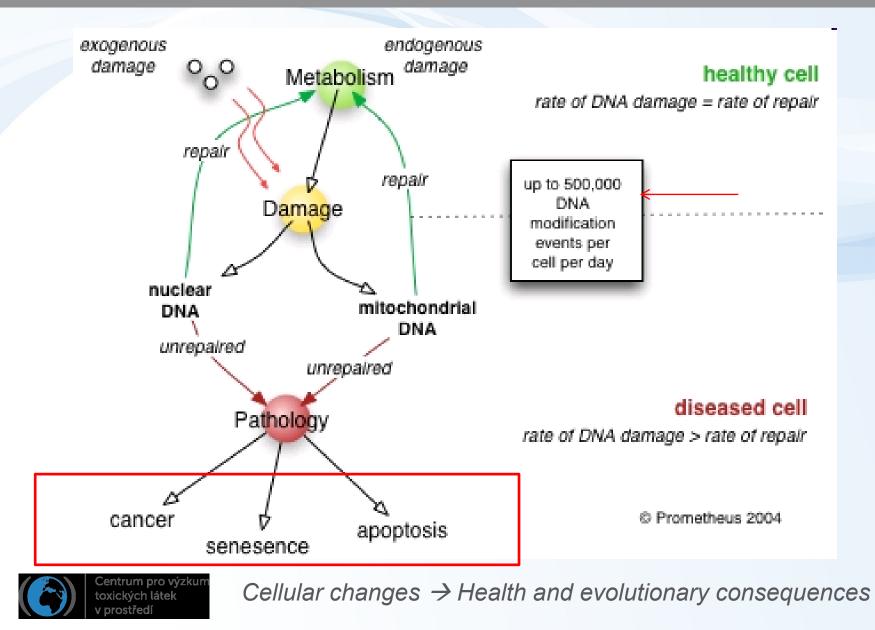
#### ... naturally

billions of nucleotides/day → most are repaired ... stress-induced → toxicity





## **DNA damage and its effects**



# **DNA** repair

# Damage of DNA is carefully controlled constitutively expressed repair systems

# Sudden changes in DNA

# Induction of additional repair enzymes (e.g. "SOS-repair" in bacteria - biomarker of DNA damage)



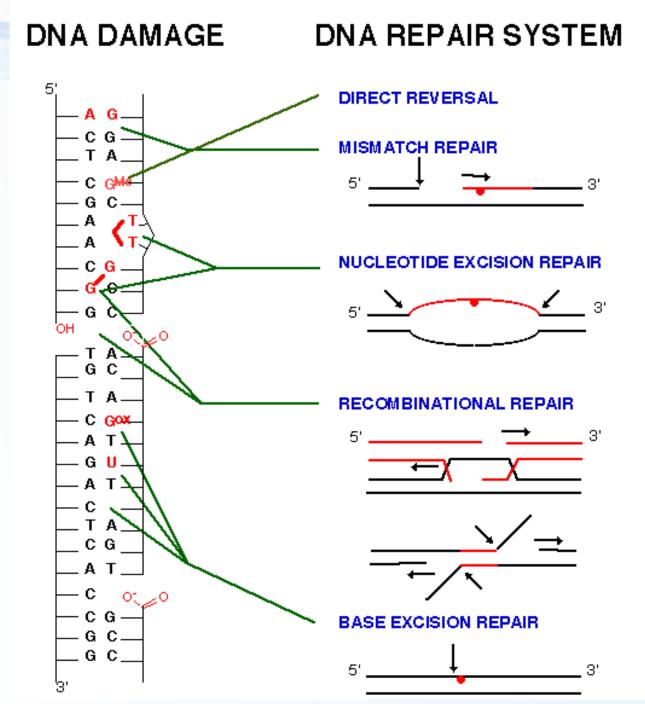
Various types of molecular changes in DNA ... and corresponding repair systems

#### Note!

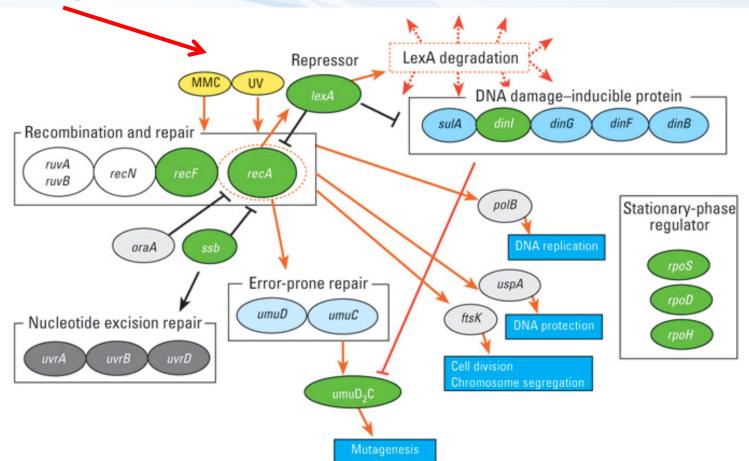
•Not all nucleotides are affected in the same rate (mutations occur only at specific sites due to physicochemical properties)

- G is commonly affected
- T=T at the same strand
- G=G crosslinks





Complex system of **SOS repair** proteins induced in *E. coli* by DNA damage



**Figure 3.** A literature-based linkage map between genes in the SOS response in *E. coli.* The map represents inducible genes/proteins in the SOS response for repair from DNA damage. Black lines indicate pathways in the normal repair process and red lines with arrows activation/induction due to an exposure to damaging agents. Recombination and repair, DNA damage–inducible protein, nucleotide excision repair, error-prone repair, and stationary-phase regulator have family molecules in each box. Green circles are genes used for the analysis.



### **TYPES** of mutations

### **POINT** mutationts

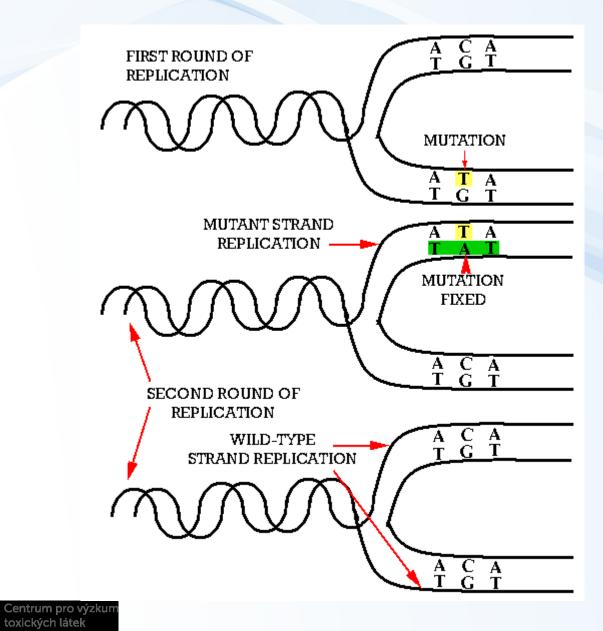
Base exchanges
Deletions / Insertions
→ Impacts of point mutations
(a) silent, (b) missense, (c) nonsense, (d) frameshift

### **CHROMOSOMAL** mutations

→ large scale impact



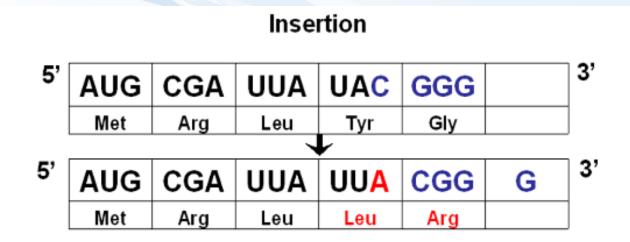
#### **BASE – EXCHANGE**





#### INSERTION DELETION

Reading frame shift

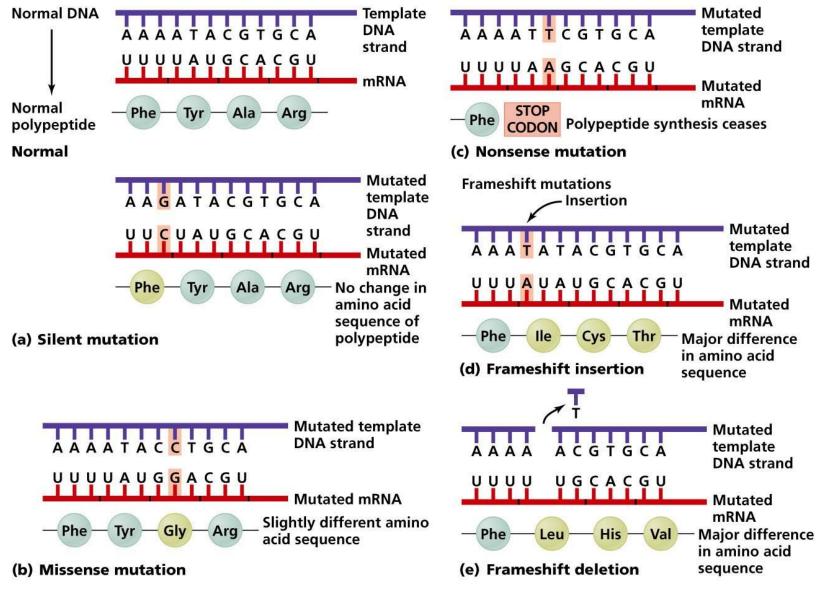


#### Deletion

5'	AUG	CGA	UUA	UAC	GGG	AAA	3'
	Met	Arg	Leu	Tyr	Gly	Lys	]
5'	$\checkmark$						
	AUG	CGA	UUA	UAG	GGA	AA	3'
	Met	Arg	Leu	Stop			]

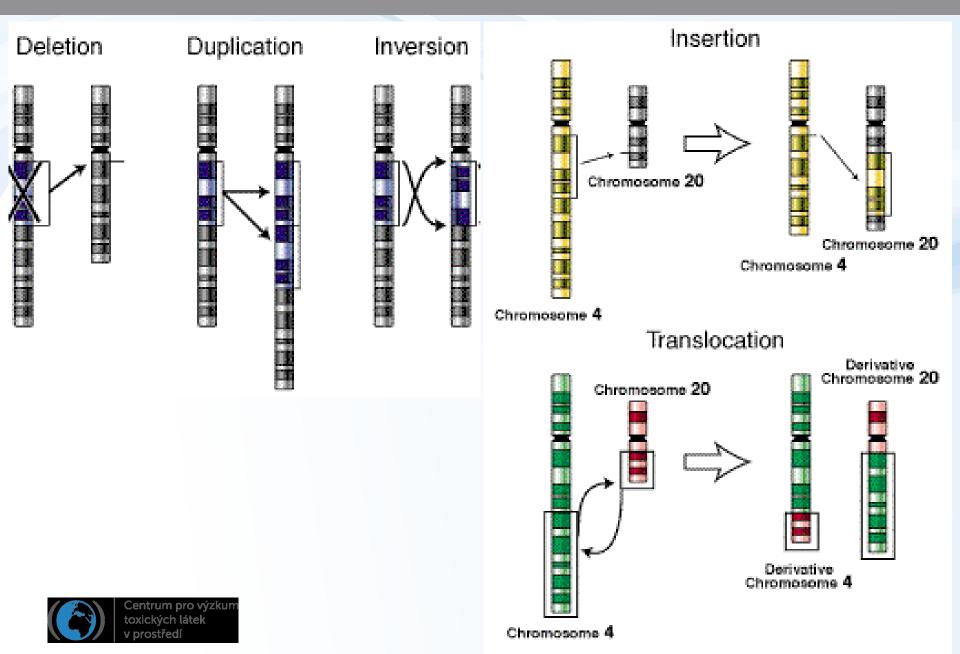


# Impacts of point mutations $\rightarrow$ (a) silent, (b) missense, (c) nonsense, (d) frameshift



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#### Large – chromosomal mutations



## What are the agents inducing mutations? MUTAGENS

# **PHYSICAL FACTORS**

## **Ionizating radiation**

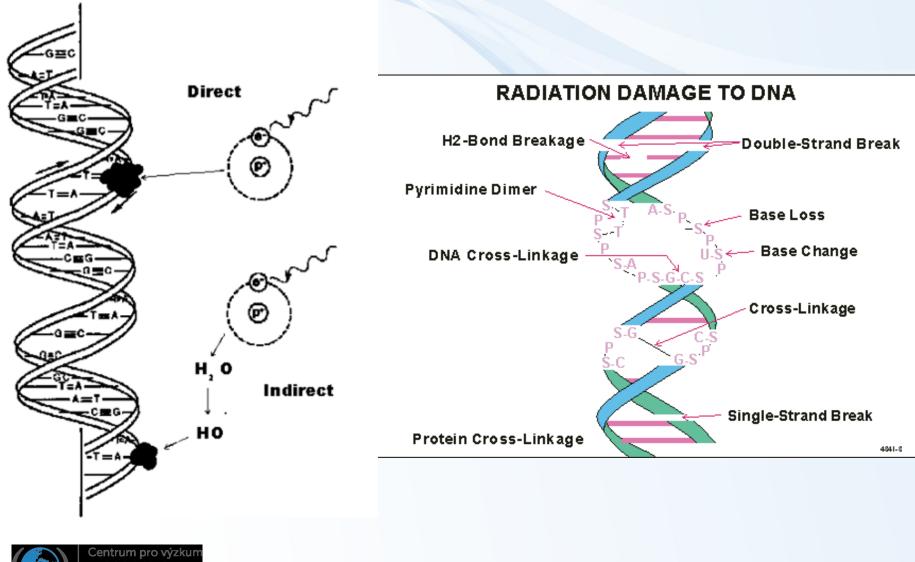
- direct interactions with NA
- interactions with water
  - $\rightarrow$  formation of OH\*
    - (and other oxygen radical species ROS)
- → Various impacts on bases and strands

## **UV** radiation

- interaction with aromatic cycles (bases)
- $\rightarrow$  base dimerization (T=T)



#### Ionizing radiation effects on DNA



toxických látek v prostředí

### What are the agents inducing mutations? MUTAGENS

## **CHEMICALS**

### 1) Small electrophilic molecules

(attracted by nucleophilic/basic sites ... e.g. in DNA)

### 2) Other reactive molecules

\* alkylating and arylating agents – covalent adducts
\* specifically intercalating agents

### 3) Base analogs

inserted during replication instead of nucleotides

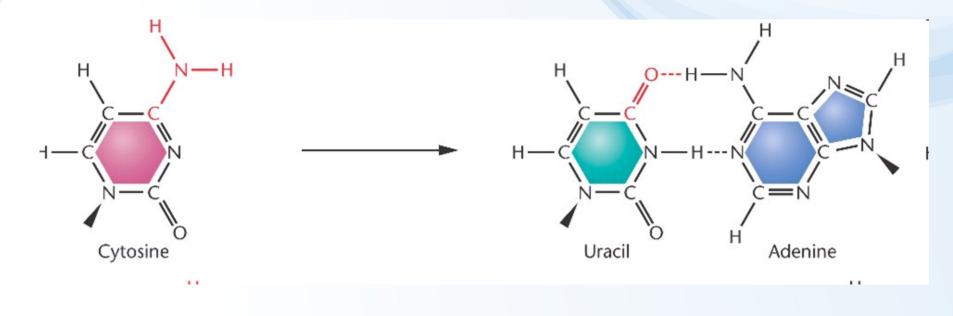
Some compounds may require "activation" by metabolism pro-mutagen (pro-carcinogen) → mutagen (carcinogen)



## Small molecules $\rightarrow$ deamination of bases

HNO<sub>2</sub>, HSO<sub>3</sub><sup>-</sup> Hydroxylamine (HO-NH2), Methoxyamine (CH3-O-NH2)

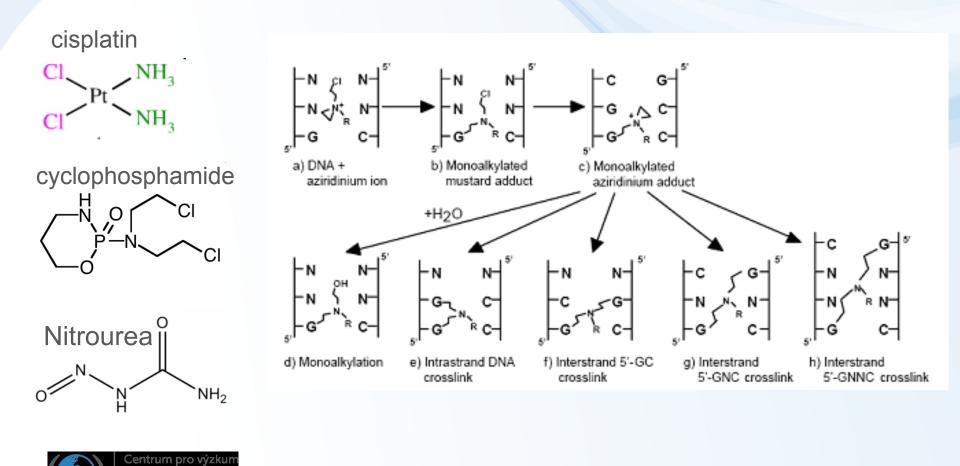
Example: deamination leading to GC  $\rightarrow$  AT shift





## ALKYLating compounds

Covalent binding to NA (alkylation of bases, crosslinks in dsDNA) Alkylsulphates, Nitro-urea, N-nitroso-alkyles, cis-platinum



toxických látek v prostředí

## **ARYLating compounds**

Covalent binding, aromatic "adducts" with bases (see also discussion at biomarkers)

Mycotoxins (Aflatoxins) – requires activation

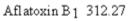
**PAHs (benzo[a]pyrene)** – requires activation **PAH** derivatives

> - 2-AA, 2-AF (grill produ - NQO – model mutagei in experiments

OCH<sub>2</sub>

O<sub>2</sub>N C-NH<sub>2</sub>

... many others

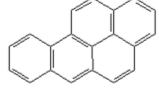


AF-2 (furylfur ami de) 248.19

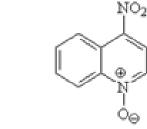
2-aminofluorene

(2-AF) 181.23

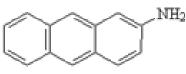
 $\rm NH_2$ 



benzo[a]pyrene (B[a]P) 252.31



4-nitroquinoline-1-oxide (NQO) 190.15

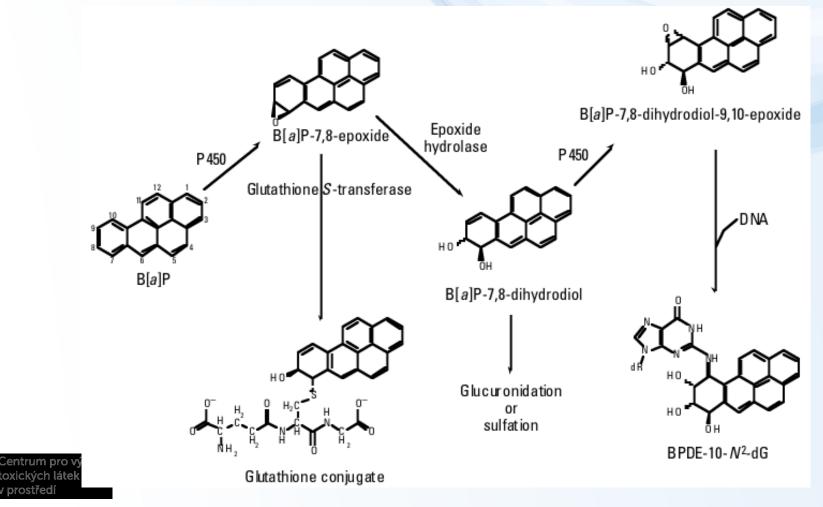


2-aminoanthracene (2-AA) 193.24



# Bioactivation of benzo[a]pyrene → genotoxicity

BaP is oxidized to epoxides and OH-derivatives during detoxification (CYP450) → increased reactivity (including binding to bases ... primarily G or A) (Similar bioactivation e.g. at aflatoxin)



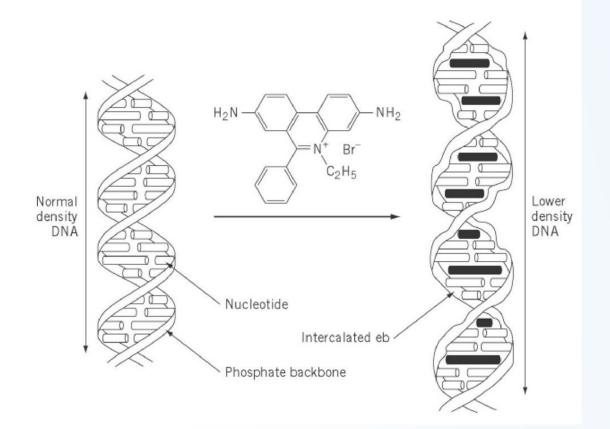
# Intercalating agents

Compounds with characteristic structures "fitting" into DNA → both noncovalent and covalent intercalation

#### Example 1 – ETHIDIUMBROMIDE

- experimental dye visualization of DNA
- intercalation  $\rightarrow$  sharing of electrones with bases  $\rightarrow$  high fluorescence

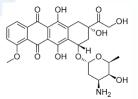
Concerned a



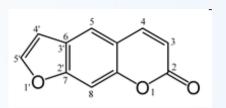
# Intercalating agents

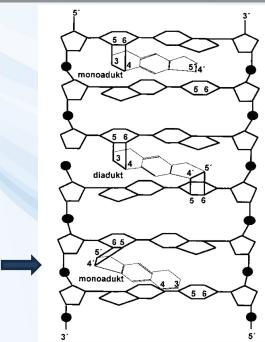
#### **Other examples**

-Anticancer drug - doxorubicin



- Psoriasis treatment – psoralen  $\rightarrow$ 





-Experimental research compnds (e.g. acriflavine)  $\rightarrow$ 

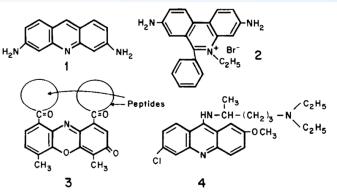


Chart 5.8. Examples of intercalating agents. Key: 1, acriflavine; 2, ethidium bromide; 3, actinomycin; 4, quinacrine.

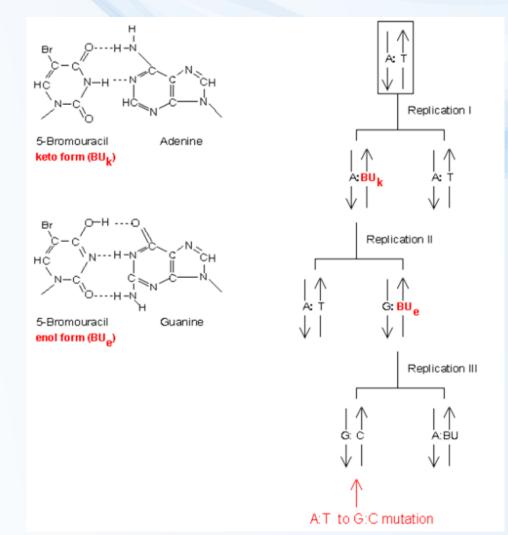


# **Base analogs**

## Structure similarity with natural bases

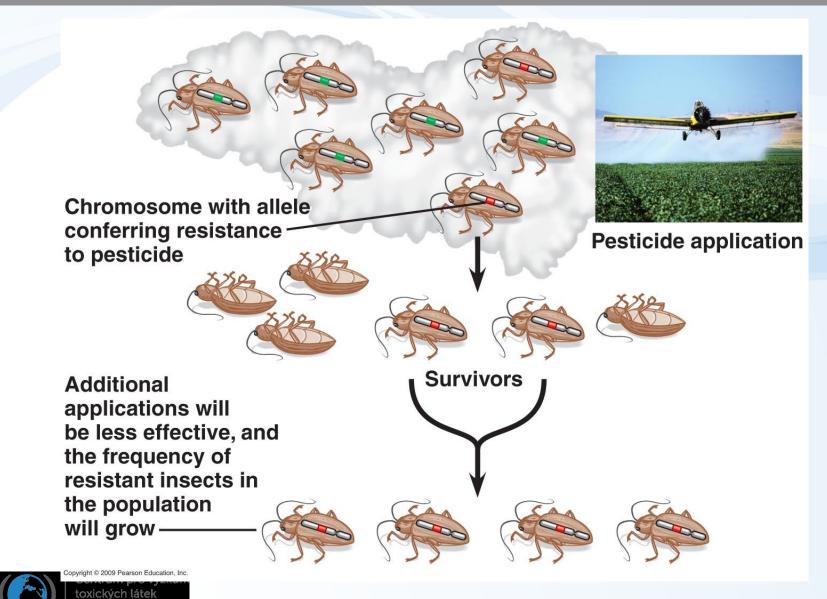
- $\rightarrow$  Incorporation into DNA during replication
- $\rightarrow$  Base exchange mutations

Example
5-Br-Uracil (anticancer drug)
AT → GC shift





# Mutations (alleles) and evolution



prostředí