

BIOMARKERS AND TOXICITY MECHANISMS 11 – BIOMARKERS of EXPOSURE and SUSCEPTIBILITY

Luděk Bláha, PřF MU, RECETOX www.recetox.cz

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









Biomarkers of Exposure

Biomarkers of internal and effective dose

depends on toxicokinetics

Biomarkers of internal dose (short / long term)

- examples: Cd in urine, DDE in fat tissues
 - should be easy to sample (urine, breath)
 - instrumental analytical methods (analyses of toxicant)

Biomarkers of effective dose

- the chemical interacted with the biological target
 - → analyses of ADDUCTS

Two types of adducts: selective and non-selective



SELECTIVE ADDUCTS OF TOXICANTS with BIOMOLECULES

SELECTIVE = CHEMICAL-SPECIFIC

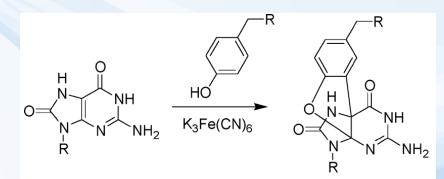
Adducts with DNA

styrene-oxide-O6-guanine N7-guanyl-aflatoxin B1

Hemoglobin-pesticides adduct

Methods of analyses:

- analytical chemistry
 - extraction from biological sample
 - chemical determination by HPLC or GC



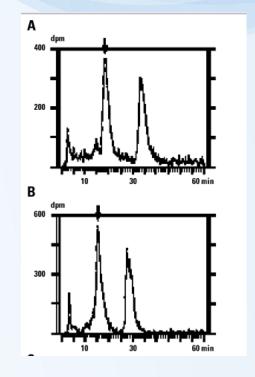




 Table 1 Reported human haemoglobin adduct levels for various xenobiotics

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|---------------------------------------|---|--------------------------------------|--|
| Chemical (type of exposure) | Adduct/analyte | Method | Adduct level (nmol g - haemoglobin) |
| N, N-Dimethylformamide (occupational) | 3-Methyl-5-isopropylhydantoin | Hydrolysis; GC-MS | 75–1000 (exposed) 4–12 (control) |
| Epichlorohydrin (occupational) | N- (2, 3-Dihydroxypropyl)valine | Modified Edman; GC-MS | 0.020 (exposed smokers) 0.007 (exposed non-smokers) 0.013 (control smokers) 0.007 (control non-smokers) |
| Acetaminophen (drug overdose) | 3-(Cystein-S-yl)acetaminophen | Immunoassay | 100-4100 |
| PAHs (occupational) | BPDF-Hb | Spectrofluorimetry | 0.005-0.139 |
| Ethylene oxide (occupational) | N- Hydroxyethylvaline | Modified Edman; GC-MS | 5–20 (exposed) 0.1–0.5 (control smokers) 0.01–0.1 (control non-smokers) |
| Ethene (occupational) | N- Hydroxyethylvaline | Modified Edman; GC-MS | 0.02 |
| Propylene oxide (occupational) | N- Hydroxypropylvaline | Modified Edman; GC-MS | 0.05-3.5 (exposed) < 0.02 (unexposed) |
| Acrylonitrile (smoking) | N- Cyanoethylvaline | Modified Edman; GC-MS | 0.09 |
| NNK (smoking) | 4- Hydroxy-1-(3-pyridyl) butan-1-one | Hydrolysis; GC-MS | 0.0015 (smokers) 0.0005 (non-smokers) |
| 4-ABP (smoking) | 4-ABP-cysteine | Hydrolysis; GC-MS | 0.00025–0.0025 (smokers) 0.00005–0.0005 (non-smokers) |
| Acrylamide (occupational, smoking) | N- (2-Carbamoylethyl)valine | Modified Edman; GC-MS | 9.5 (production workers) 0.054 (laboratory workers) 0.116 (smokers) 0.031 (non-smokers) |
| Butadiene (occupational) | N- (2,3,4-Trihydroxybutyl)valine | Modified Edman; GC-MS | 0.010–0.014 (exposed) 0.002–0.003 (control) |
| Styrene (occupational) | 2-Phenylethanol | Cleavage with Raney nickel, GC-MS | 3.7–8.0 (exposed) 2.0–8.6 (control) |

Non-selective adducts

 binding with macromolecules (DNA, proteins) with no further information on the structure of actual adduct (i.e. causative agent not clear)

Typical nonselective biomarker methods

- ³²P-postlabelling assay
- DNA-strand breaks comet assay
- identification of oxidized DNA: 8-hydroxy-2'-deoxyguanosine



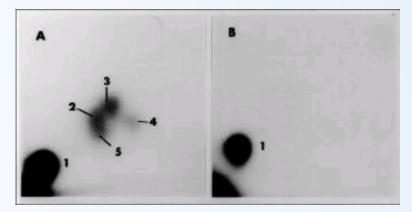
³²P-postlabelling assay principle

- Digestion of NA
- •Enzymatic labelling with 32P (kinase)
- •TLC or HPLC analyses of products

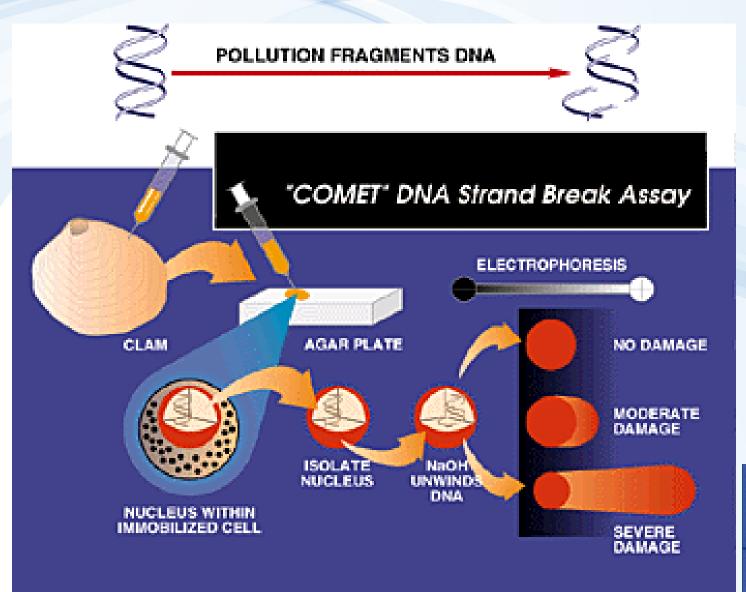
TLC result (thin layer chromatography)

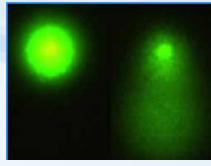
A - 2-5 = various adducts

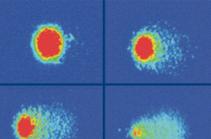
B - controls



Comet assay - principle

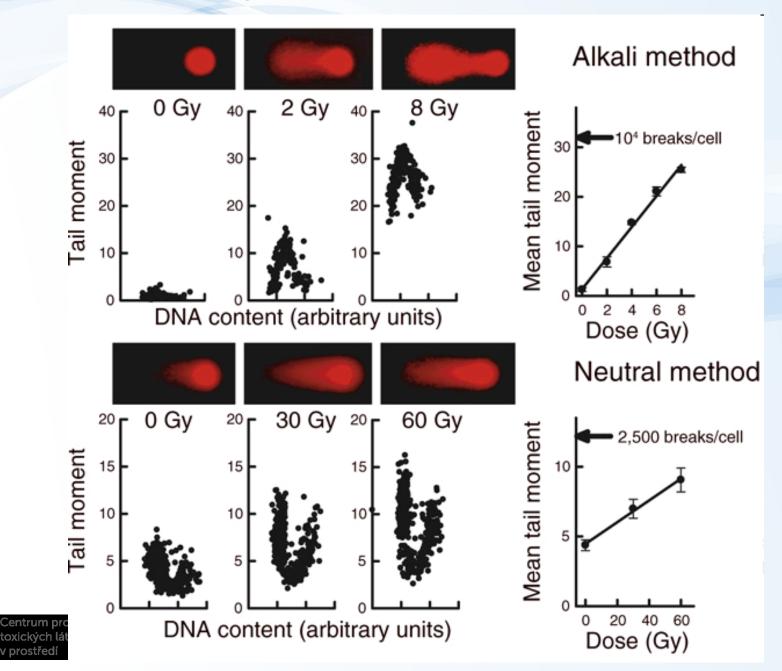








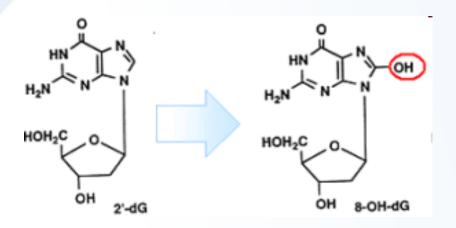
Example results - Comet assay vs. radiation



8-hydroxy-2 -deoxyguanosine analysis

Oxidative damage to DNA

- many causes → 8-OH-dG is the most common marker of DNA oxidation



Analysis: analytical chemistry methods

- HPLC
- immunochemistry (ELISA)

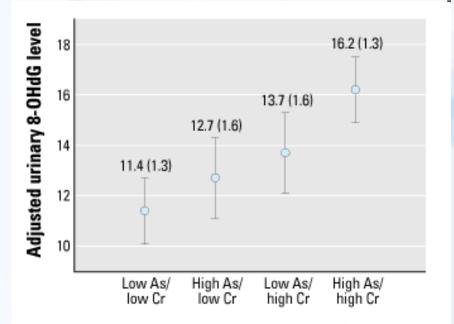
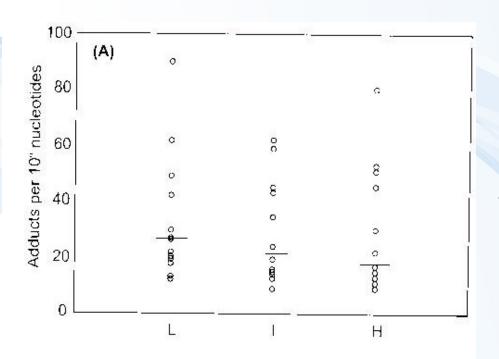


Figure 1. Adjusted urinary 8-OHdG level (ng/mg creatinine) by urinary arsenic and urinary chromium concentrations. Values shown are mean \pm SE. Cut points were determined according to medians (arsenic, 7.7 µg/g creatinine; chromium, 2.0 µg/g creatinine) of urinary creatinine-adjusted levels among all subjects.

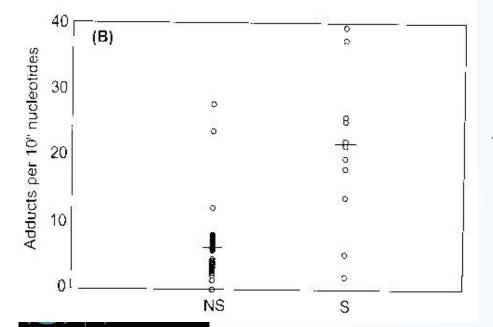




PAH-DNA adducts

* often high variability* may have difficult interpretation

Occup. exposure (Low / Intermed. / High)



Occupational
Non-exposed (NS)
vs.
Exposed (S)

Biomarkers of susceptibility



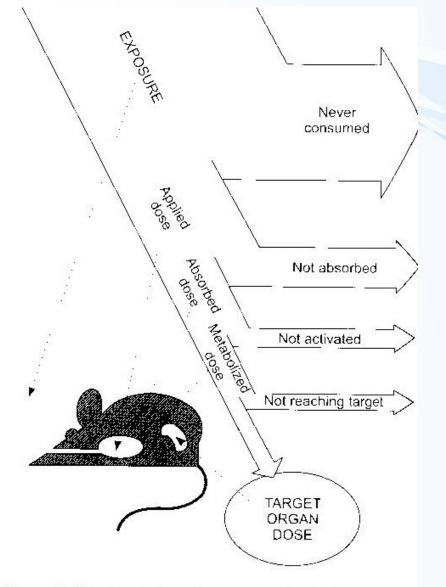


Figure 2 Representation of the relationships between ambient exposure and critical target dose and the progressive decrease in effective exposure due to various biological barriers. Source: *Low-Dose Extrapolation of Cancer Risks: Issues and Perspectives*, p. 188. Used with permission. c 1995 International Life Sciences Institute, Washington, DC, U.S.A.

Toxicokinetics

determines susceptibility of an individual at various levels

→ Biomarkers of susceptibility

Will the individual be sensitive? Will patient respond to a drug?

Importance of susceptibility biomarkers

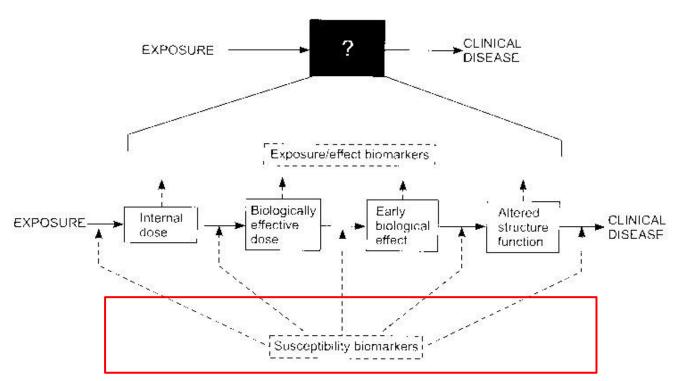


Figure 1 The biomarker paradigm linking exposure with disease and showing expansion of the classical epidemiological 'black box' to reveal discrete mechanistic stages. Reprinted with permission from *Environ, Sci. Technol.* (1997) **31**, pp. 1837–1848. Copyright 1997 American Chemical Society.



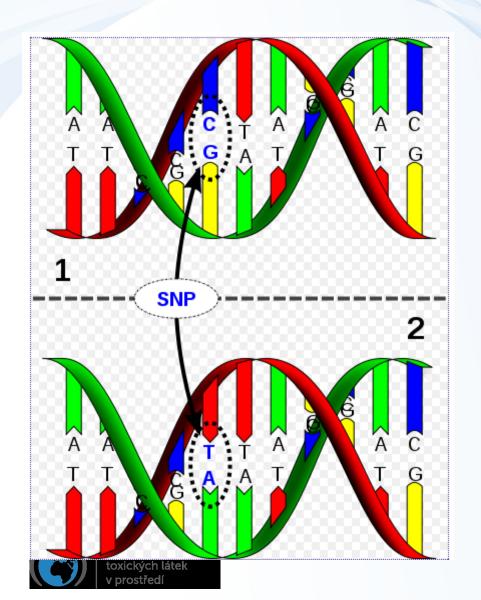
Biomarkers of susceptibility

Susceptibility depends on genotype and metabolism

- genetic polymorphism in detoxification enzymes
- variability in specific isoenzymes
- → susceptibility to "activate" toxicants: example: N-acetylation of arylamines NAT2
 - → susceptibility to genotoxins
- → family cancers
- → susceptibility to drugs (including anticancer drugs)



Example: genetic polymorphism SNPs - single nucleotide polymorphism



SNPs

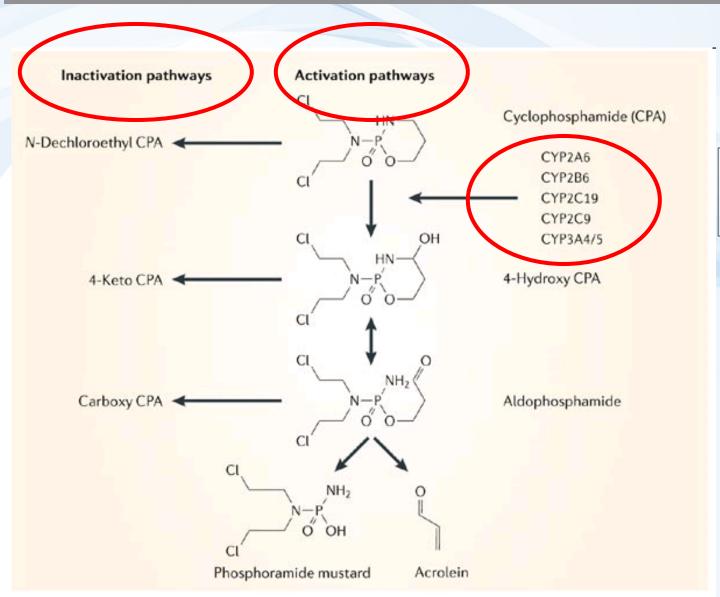
- → affects protein functions
- → in specific cases (see example) some SNPs identified

→ PERSONALIZED MEDICINE

To identify SNP as a biomarker

Many **genotypes** (from many individuals) must be sequenced and compared with **phenotype** (e.g. responsiveness to certain drug)

Cyclophosphamide (anticancer drug) and its toxicity





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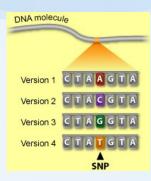
Example: genetic polymorphism

CYP450 Enzymes and Polymorphisms

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| Enzyme | Fraction of drug metabolism | Major polymorphisms |
|---------|--------------------------------|-------------------------|
| CYP3A4 | 40-45% | Rare |
| CYP2D6 | 20-30% | '2xn, '4, '10, '17, '41 |
| CYP2C9 | 10% | 2, 3 |
| CYP2C19 | 5% | °2, °3 |
| CYP1A2 | 5% | *1K |
| CYP2B6 | 2-4% | - |
| CYP2E1 | 2-4% | - |
| CYP2A6 | 2% | '4, '9 |
| CYP2C8 | 1% | *3 |
| CYP3A5 | <1% | .3 |

Alleles known to be involved in polymorphism



The CYP 2D6 gene is extremely polymorphic with more than 70 allelic variants described so far ¹

Ingelman-Sundberg, TRENDS in Pharmacological Sciences, Vol. 25 No.4 April 2004 ¹ Dahl, Clin. Pharmacokinet 2002; 41 (7); 453-470

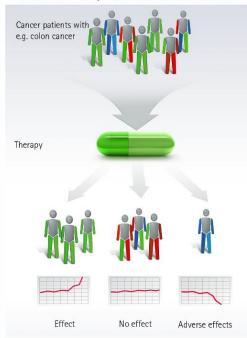
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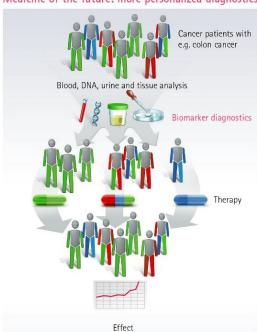


Personalized medicine

Personalized medicine: tailored treatments

Medicine of the present: one treatment fits all Medicine of the future: more personalized diagnostics



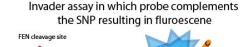


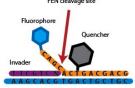
Different people respond differently to the same therapy: while one treatment brings about the desired success in one group of patients with e.g. colon cancer, it does not change the condition of other groups at all, or even leads to adverse effects (left). The reason: the genetic makeup and metabolic profile of each individual patient influences the effect of a drug. Personalized medicine takes these individual patterns of cellular and metabolic products into account in the diagnostic phase: biomarker diagnostics separates patients into groups with similar characteristics, and provides information on the best individual treatment. This should enable all patients to benefit from their own, "personal" therapy.

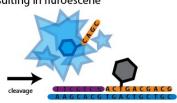


SNP diagnostics:

- 1) DNA isolation
- 2) Multiplication of specific gene eg. CYP
- 3) SNP identification
- ... Molecular biology methods such as
- * NA sequencing
- * Probe pairing ... number of variants







Invader assay in which probe mismatches at the SNP location preventing cleavage from occuring

