

BIOMARKERS AND TOXICITY MECHANISMS 07 – Mechanisms Metabolism & Detoxification

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



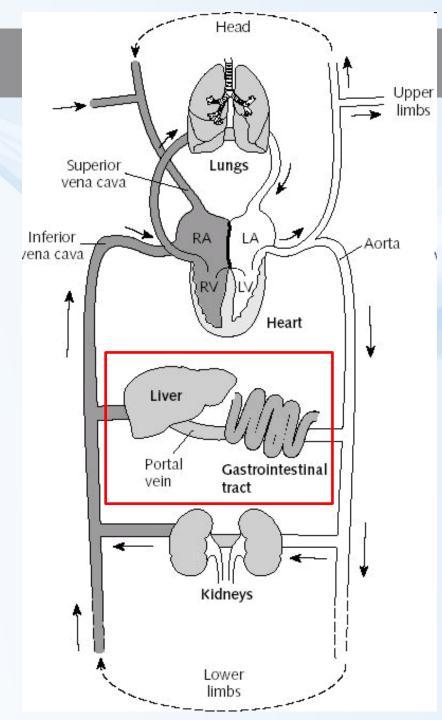






Metabolism and detoxification

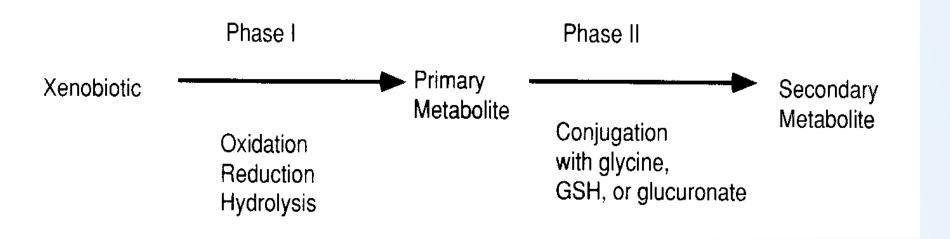
- Chemicals enter body
 ... mostly via food
- Pass directly through liver
 - → main metabolism organ





Detoxification

- Basic principle of detoxification
 - elimination of hydrophobic compounds from body→ formation of more polar & soluble products
- Two principal phases in metabolism (Phase I & II)
 - well studied in vertebrates (mammals)
 - liver: major organ involved in detoxification
- Plants
 - similar oxidating enzymes as described (cytochrom oxidase, phenol oxidase, peroxidase...)
- Phase III elimination both from cell & body



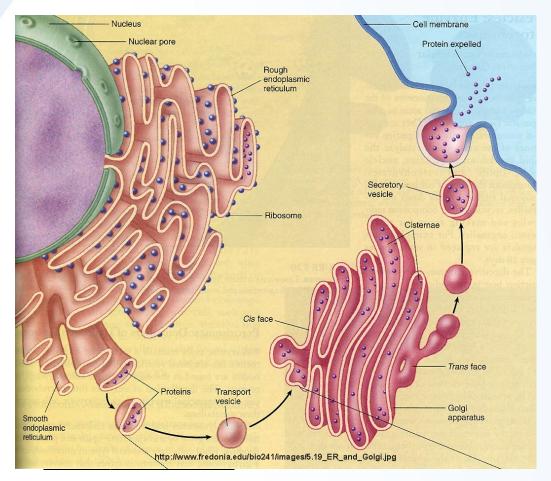
Importance of nutrients and vitamins in detoxification

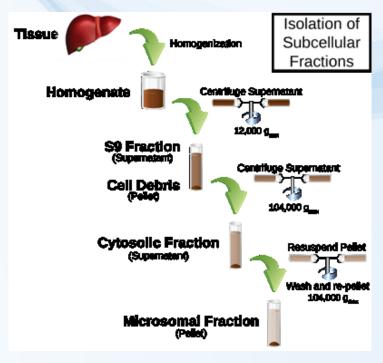
Detoxification Pathways



Phase I

- Key enzymes MFOs = mixed function oxidases / oxygenases
- Membrane bound to Endoplasmic Reticulum
 - membrane vesicles "microsomes" = S-9 fraction can be extracted from cells

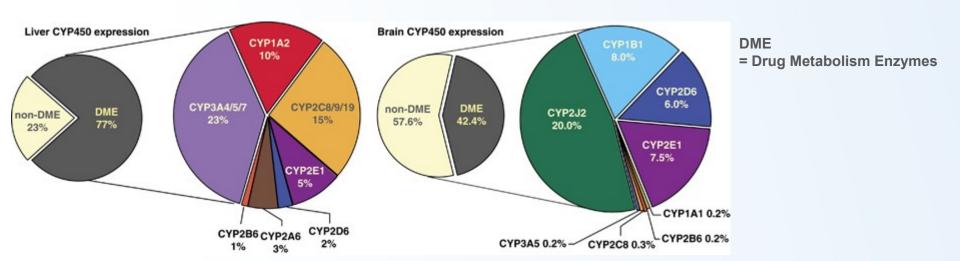




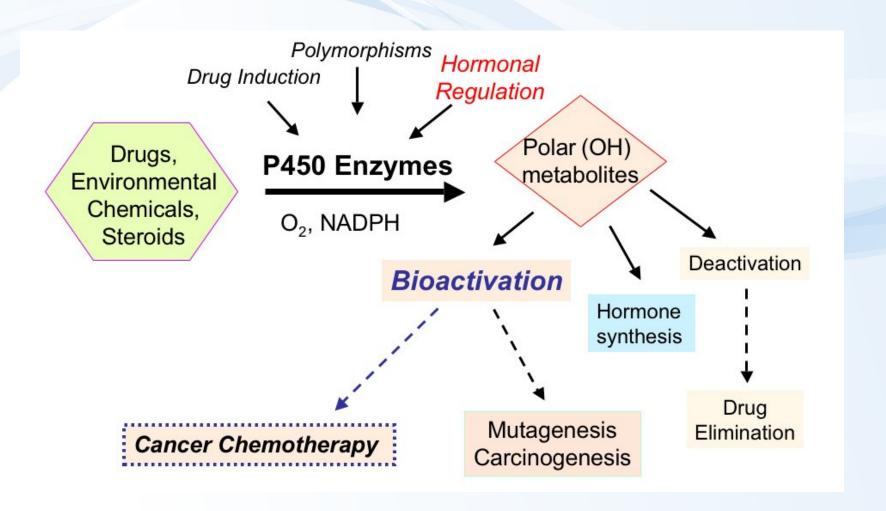
S9 microsomes used for in vitro metabolization (e.g.during genotoxicity testing)

Detoxification - Phase I

- Key principle enzymes are cytochromes P450 (CYPs)
 - Haem (porfyrin) containing enzymes
 - superfamily of more than 150 genes several classes and subclasses
 - different substrate specificity; structure ...
- Some examples ... Diverse functions
 - Cytochrome P450 1A (CYP1A)
 - basic for detoxification of hydrophobic environmental contaminants
 - Cytochrome P450 19A (CYP19)
 - "aromatase" involved in synthesis of estradiol (aromatization of testosterone)



CYPs and their functions





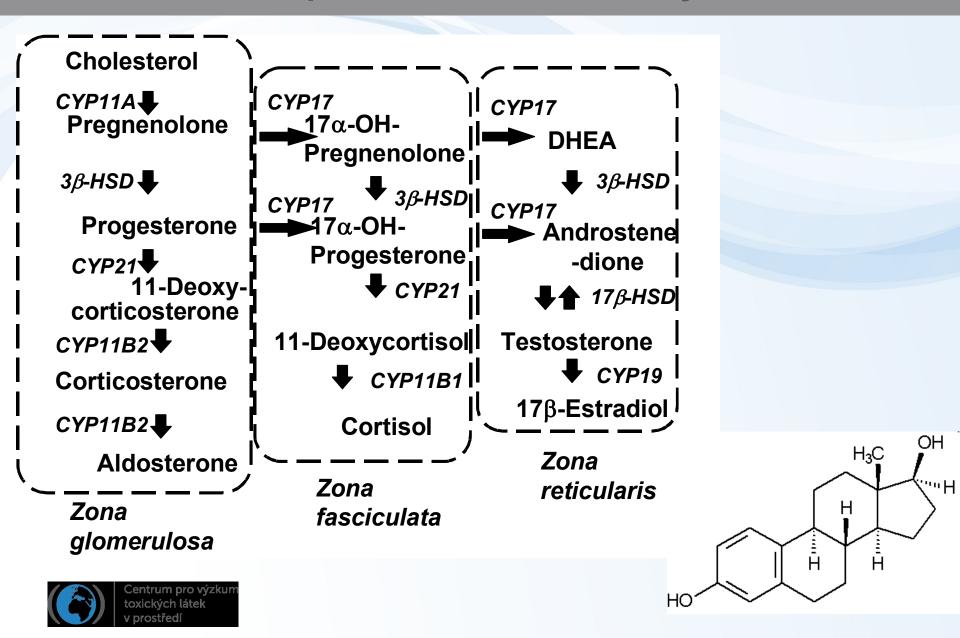
Types of reactions catalyzed by CYPs (and Phase II enzymes)

Phase	Туре	Reaction (gene)	Substrate C
ī	MFO	O-Deethylase (CYP1A1)	7-Ethoxycoumarin
1	MFO	Aryl hydrocarbon hydroxylase (CYP1A1)	PAH
1	MFO	Hydroxylase (CYP3A7)	Cortisol
1	MFO	Aromatase (CYP19)	Androgens
1	MFO	Cholesterol side-chain cleavage (CYP11A)	Cholesterol
I	MFO	Estrogen catechol formation, 2-Hydroxylation (<i>CYP1A1</i>) 4-Hydroxylation (<i>CYP1B1</i>)	Estrogens
	MFO	25-Hydroxycholecalciferol hydroxylase	25-Hydroxycholecalciferol
1	Oxidoreductase	17β-Hydroxydehydrogenase	,
		Type 1	Estrone to estradiol
		Type 2	Estradiol to estrone
	Oxidoreductase	11β-Hydroxydehydrogenase	Cortisol/cortisone
	Oxidation	Dehydrogenase	Alcohol/acetaldehyde
1	Oxidation	Monoamine	Norepinephrine
	Sulfatase	Sulfate cleavage	Steroid sulfates
II	Conjugation	GST	Epoxides
II	Conjugation	Catechol-O-methyltransferase	Catecholamines, catechol estrogens



Highlighted = will be discussed also later

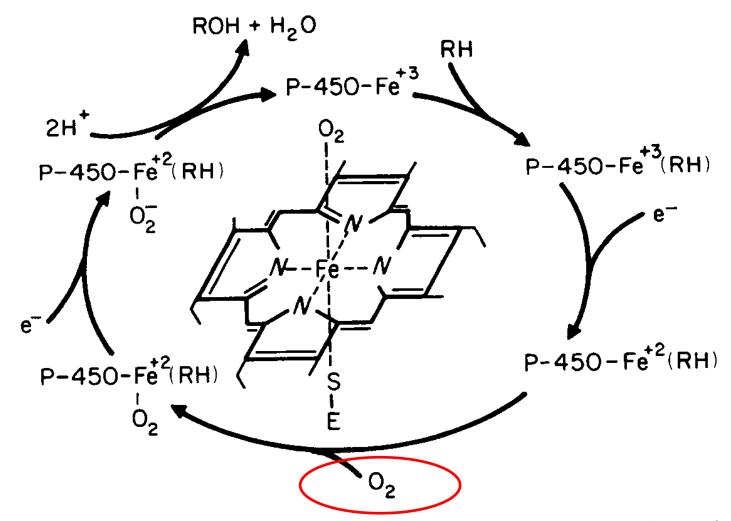
CYPs - example: steroid hormone synthesis



CYP450 overview

Family	Function	Members	Names	
CYP1	drug and steroid (especially estrogen) metabolism	3 subfamilies, 3 genes, 1 pseudogene	CYP1A1, CYP1A2, CYP1B1	
CYP2	drug and steroid metabolism	13 subfamilies, 16 genes, 16 pseudogenes	CYP2A6, CYP2A7, CYP2A13, CYP2B6, CYP2C8, CYP2C9, CYP2C18, CYP2C19, CYP2D6, CYP2E1, CYP2F1, CYP2J2, CYP2R1, CYP2S1, CYP2U1, CYP2W1	
СҮРЗ	drug and steroid (including testosterone) metabolism	1 subfamily, 4 genes, 2 pseudogenes	CYP3A4, CYP3A5, CYP3A7, CYP3A43	
CYP4	arachidonic acid or fatty acid metabolism	6 subfamilies, 11 genes, 10 CYP4A11, CYP4A22, CYP4B1, CYP4F2, CYP4F3, CYP4F8, CYP4F11, CYP pseudogenes CYP4F22, CYP4V2, CYP4X1, CYP4Z1		
CYP5	thromboxane A2 synthase	1 subfamily, 1 gene	CYP5A1	
CYP7	bile acid biosynthesis 7-alpha hydroxylase of steroid nucleus	2 subfamilies, 2 genes	subfamilies, 2 genes CYP7A1, CYP7B1	
CYP8	varied	2 subfamilies, 2 genes	CYP8A1 (prostacyclin synthase), CYP8B1 (bile acid biosynthesis)	
CYP11	steroid biosynthesis	2 subfamilies, 3 genes	CYP11A1, CYP11B1, CYP11B2	
CYP17	steroid biosynthesis, 17-alpha hydroxylase	1 subfamily, 1 gene	CYP17A1	
CYP19	steroid biosynthesis: aromatase synthesizes estrogen	1 subfamily, 1 gene	CYP19A1	
CYP20	unknown function	1 subfamily, 1 gene	CYP20A1	
CYP21	steroid biosynthesis	2 subfamilies, 2 genes, 1 pseudogene	CYP21A2	
CYP24	vitamin D degradation	1 subfamily, 1 gene	CYP24A1	
CYP26	retinoic acid hydroxylase	3 subfamilies, 3 genes	CYP26A1, CYP26B1, CYP26C1	
CYP27	varied	3 subfamilies, 3 genes	CYP27A1 (bile acid biosynthesis), CYP27B1 (vitamin D3 1-alpha hydroxylase, activates vitamin D3), CYP27C1 (unknown function)	
СҮРЗЭ	7-alpha hydroxylation of 24-hydroxycholesterol	1 subfamily, 1 gene	CYP39A1	
CYP46	cholesterol 24 hydroxylase	1 subfamily, 1 gene	CYP46A1	
CYP51	cholesterol biosynthesis	1 subfamily, 1 gene, 3 pseudogenes	CYP51A1 (lanosterol 14-alpha demethylase)	

Hydroxylation (oxidation) mechanism – key in "detoxification"





Scheme 3.1. Outside: suggested sequence of hydroxylation reactions carried out by cytochrome P-450. Inside: schematic presentation of the configuration of the P-450 prosthetic group.

Examples of CYP mediated reactions

Oxidation

Side Chain Oxidation

$$R-N$$
 CH_3
 CH_3
 CH_3
 CH_3
 CH_3
 CH_3
 CH_3
 CH_3

N-Dealkylation



Examples of CYP mediated reactions

$$R-O-CH_3$$
 $R-OH + HCHO$

O-Dealkylation

$$R-CH_2-CH-CH_2$$
 $R-CH_2-C-CH_2 + NH_3$
 $| NH_2 | O$

Deamination

Examples of CYP mediated reactions

$$O_2N$$
 O_2N
 O_2N

Desulfuration

Azobenzine

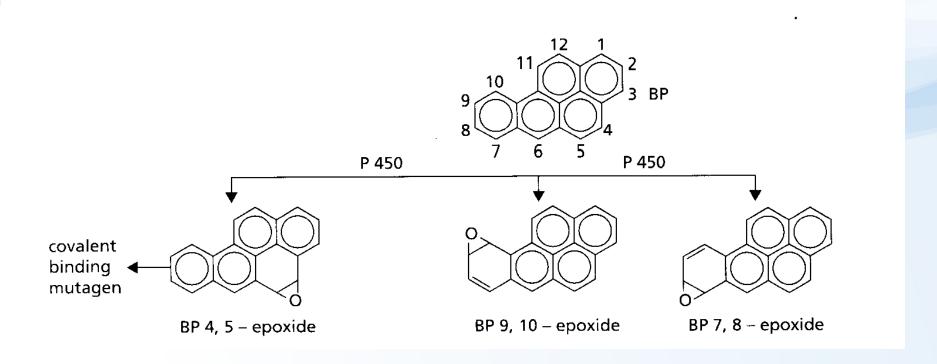
Reduction

Hydrolysis



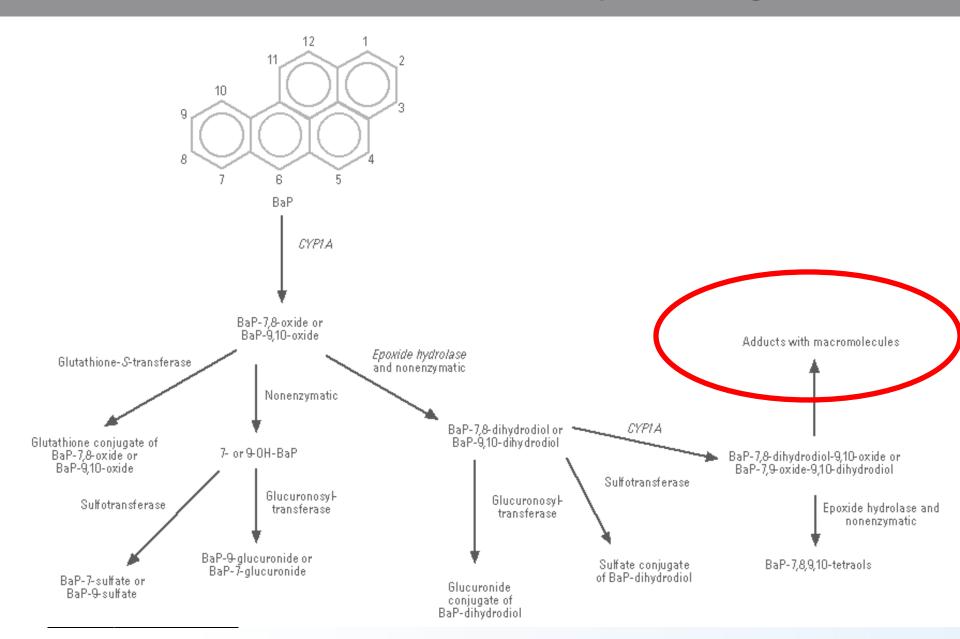
CYPs and BIOACTIVATION pro-mutagen (procarcinogen) → mutagen (carcinogen)

Benzo[a]pyrene





CYPs and BIOACTIVATION of procarcinogen



CYPs and BIOACTIVATION – AFLATOXIN-A

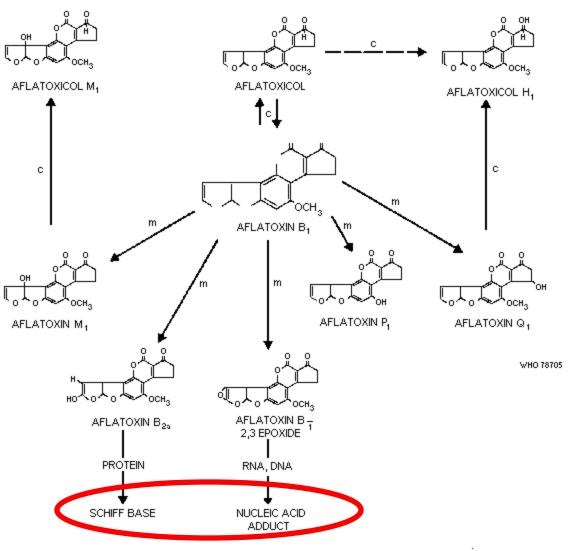
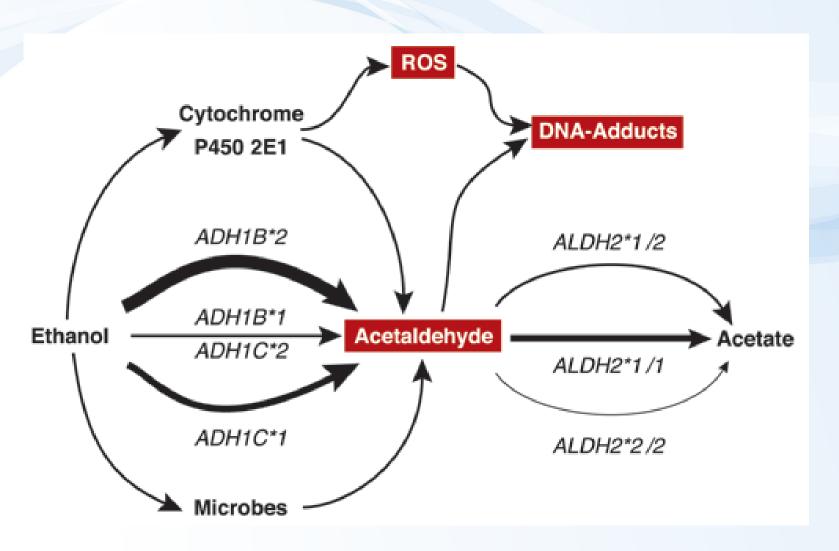




Fig. 2. Aflatoxin B₁ metabolism in the liver.

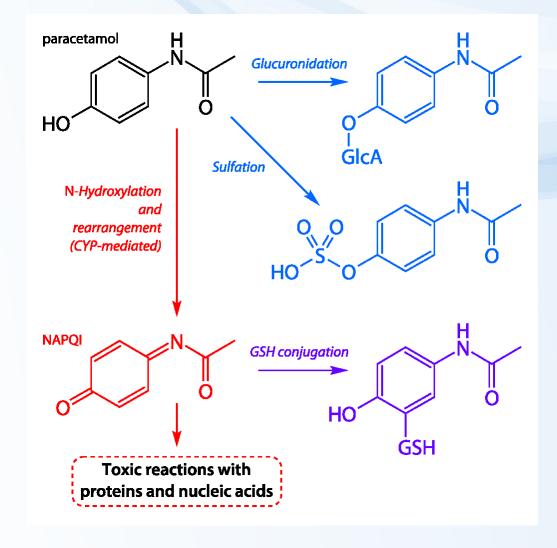
CYPs and BIOACTIVATION – ethanol





CYPs and toxicity of drugs

Example - PARACETAMOL toxicity





Detoxification - Phase II

- Key reactions = conjugations
 - Reactive xenobiotics or metabolites formed in phase I with endogeneous substrates
 - saccharides and their derivatives glucuronic acid,
 - aminoacids (glycine)
 - peptides: glutathione (GSH)
- Forming water soluble AND "nontoxic" products (conjugates)
- Phase II enzymes ("transferases"):
 - glutathion S-transferase (GST)
 - UDP-glucuronosyltransferase (UDP-GTS)
 - epoxid hydrolase (EH)
 - sulfotransferase (ST)

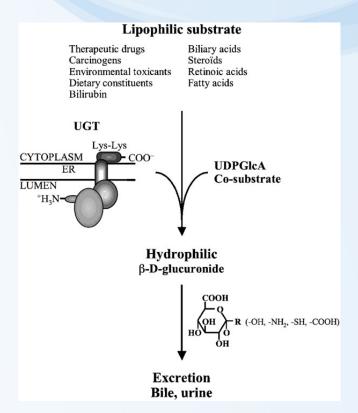




Table 3. Major phase II detoxification activities in humans

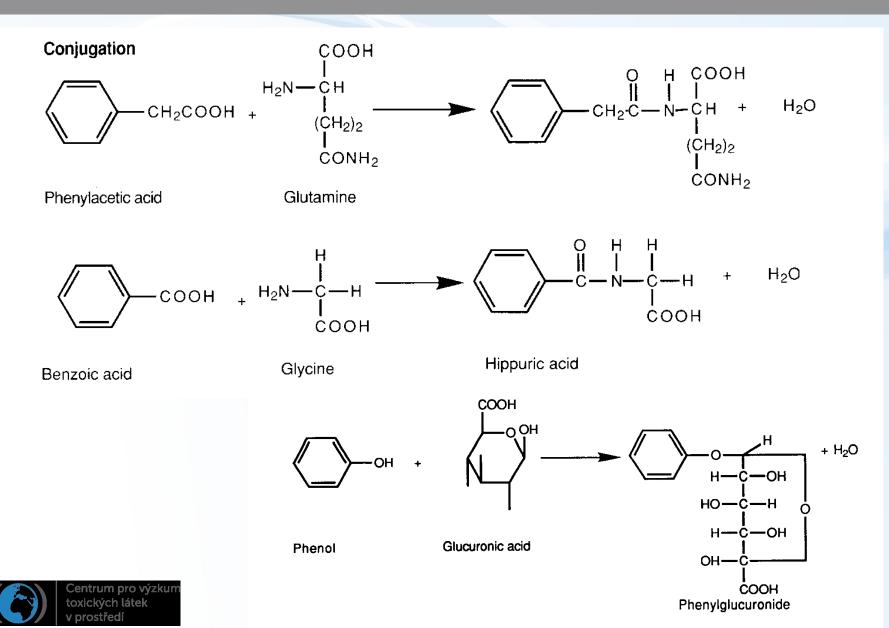
R	eaction	Enzyme	Localizationa	Substrates
H	₂ O	Epoxide hydrolase	Microsomes Cytosol	Epoxides
G	lutathione	Glutathione transferases	Microsomes	Electrophiles
	lucuronic acid JDPGA) ^b	Glucuronyl transferases	Microsomes	Phenols, thiols, amines, Carboxylic acids
	ulfuric acid APS) ^b	Sulfotransferase	Cytosol	Phenols, thiols, amines
	ethyl Group AM) ^b	N- and O- methyl transferases	Cytosol Microsomes	Phenols, amines
	cetic acid cetyl-CoA) ^b	N-acetyl transferases	Cytosol	Amines
(A ta	mino acids cetyl-CoA, urine, glycine)	Amino acid transferases	Microsomes	Carboxylic acids

Glutathione

- major donor of SH (thiol) groups in cells (MW ~ 300 g/mol)
- concentrations in tissues and blood up to 5 mM (1.5 g/L)



Examples of conjugation reactions



Xenobiotic conjugations with GSH

Glutathione



Glutathione-S-Conjugate

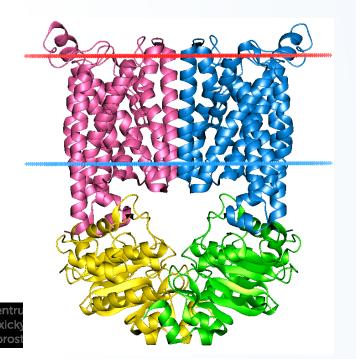
placement of aromatic halogens by glutathione

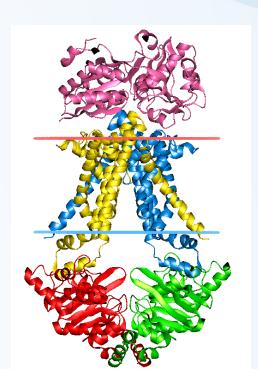




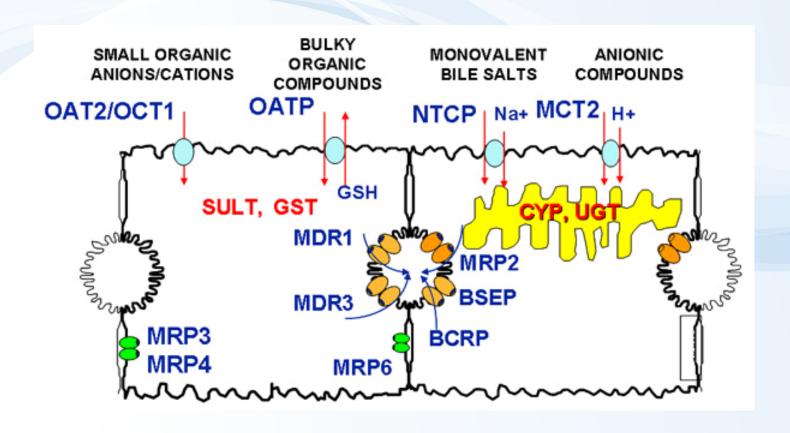
Phase III – elimination / membrane transport

- Phase III transporters
 - ATP-binding cassette transporters (ABC transporters)
 - protein superfamily (one of the largest, and most ancient in all extant phyla from prokaryotes to humans)
 - transmembrane proteins transport across extra- and intracellular membranes (metabolic products, lipids, sterols, drugs)





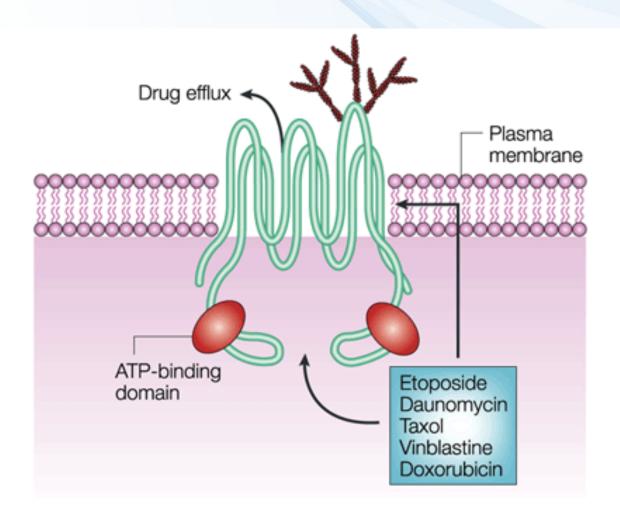
ABC transporters - examples



- MRP (MDR) multidrug resistance-associated protein family
- OATP Organic Anion Transporting Polypeptide
- P-glycoprotein

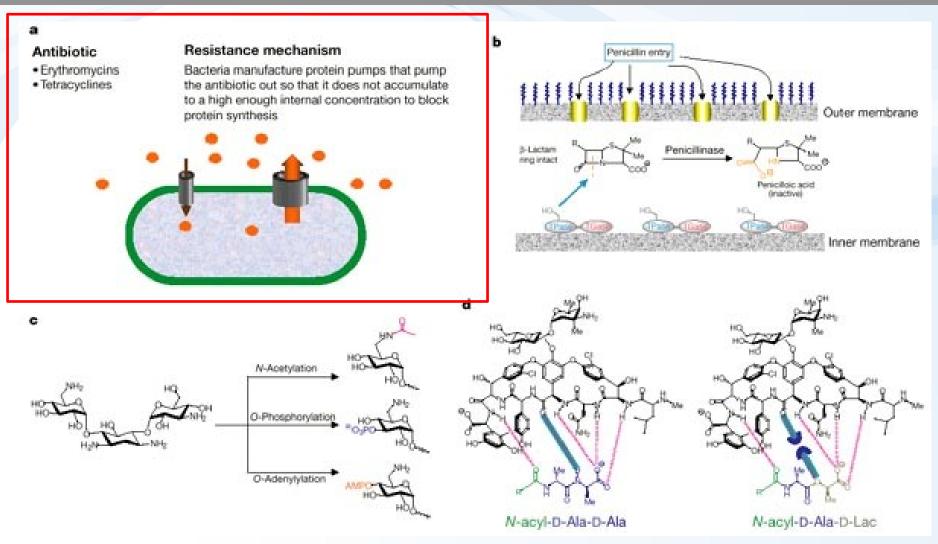


ABC one of the resistance mechanisms of tumour cells to anticancer drugs



Nature Reviews | Cancer

ABC one of the resistance mechanisms of bacteria to antibiotics





Constitutive vs Induced detoxification enzymes

- Detoxification enzymes expression
 - Constitutive low background levels (always present)
 - May be induced by substrates
 - CYP1A induction via Ah-receptor (AhR)
 - Substrate: hydrophobic organochlorine compounds (PCDDs/Fs, PAHs PCBs ...)
 [see also: lectures on nuclear receptors]
 - Other CYPs
 - Drugs → inductions of specific CYP classes
 - Phase II enzymes
 - Substrates = reactive toxicants, metabolites from Phase I
 - ABC transporters
 - Induction by respective chemicals (drugs etc)



CYP1A induction – role of AhR

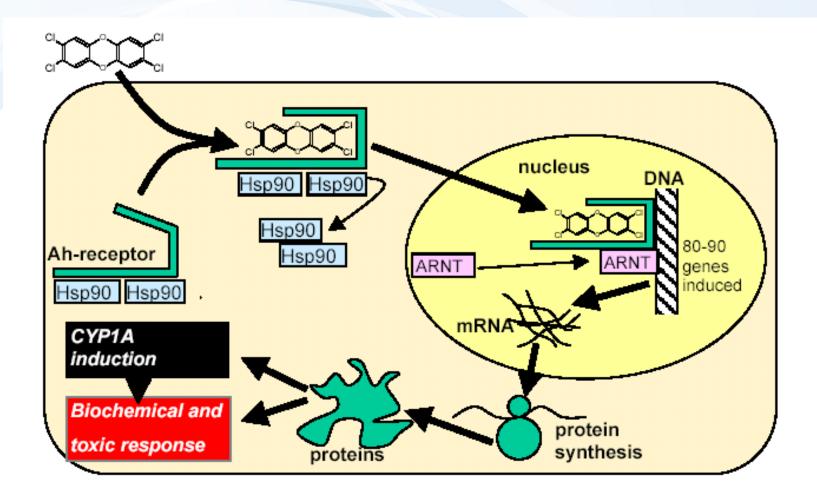


Figure 5. The mechanism of CYP1A induction mediated through the aryl hydrocarbon receptor (AhR). (Figure by M. Engwall).



Summary – "toxic consequences" of detoxification

BIOACTIVATION

- activation of pro-mutagens/pro-carcinogens etc.
- increasing side adverse effects of certain drugs

Increase in oxidative reactions – oxidative stress

- production of Reactive Oxygen Species (ROS)
 (see oxidative damage and stress lectures)
- Side toxic effects (see nuclear receptor lectures)
 - e.g. increased degradation of endogeneous compounds (retinoids – regulatory molecules degraded by CYP1A
 - Crosstalk with other mechanisms & receptors

Energy (ATP) depletion

- − chronic inductions of detox enzymes
 → permanent extra energetic demand
- Development of resistance to toxic compounds
 - Loss of efficiency of anticancer drugs, antibiotics etc.

