

Pharmacogenetics of cytochrome P450 and its application and value in drug therapy – the past, present and future

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The human genome

- 3,120,000,000 nucleotides
- 23,000 genes
- >100 000 transcripts (!)
- up to 100,000 aa differences between two proteomes
- 10,000,000 SNPs in databases today

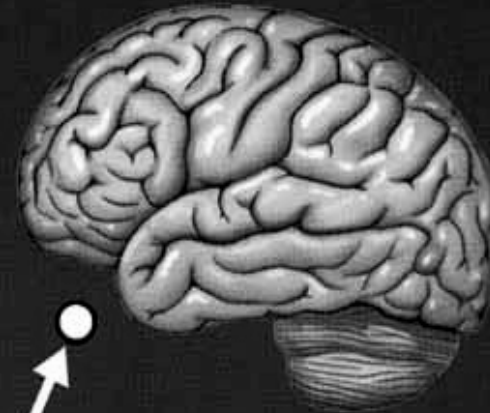
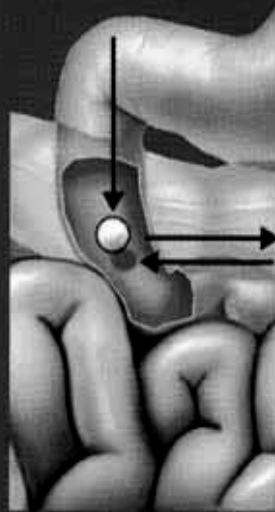
The majority of the human genome is transcribed and has an unknown function



Interindividual variability in drug action

Absorption / Excretion
Slow Rapid Slow Rapid

Receptor interactions
Poor Efficient



Metabolism
Poor Efficient Ultrarapid

Drug-drug interactions

Kidney function

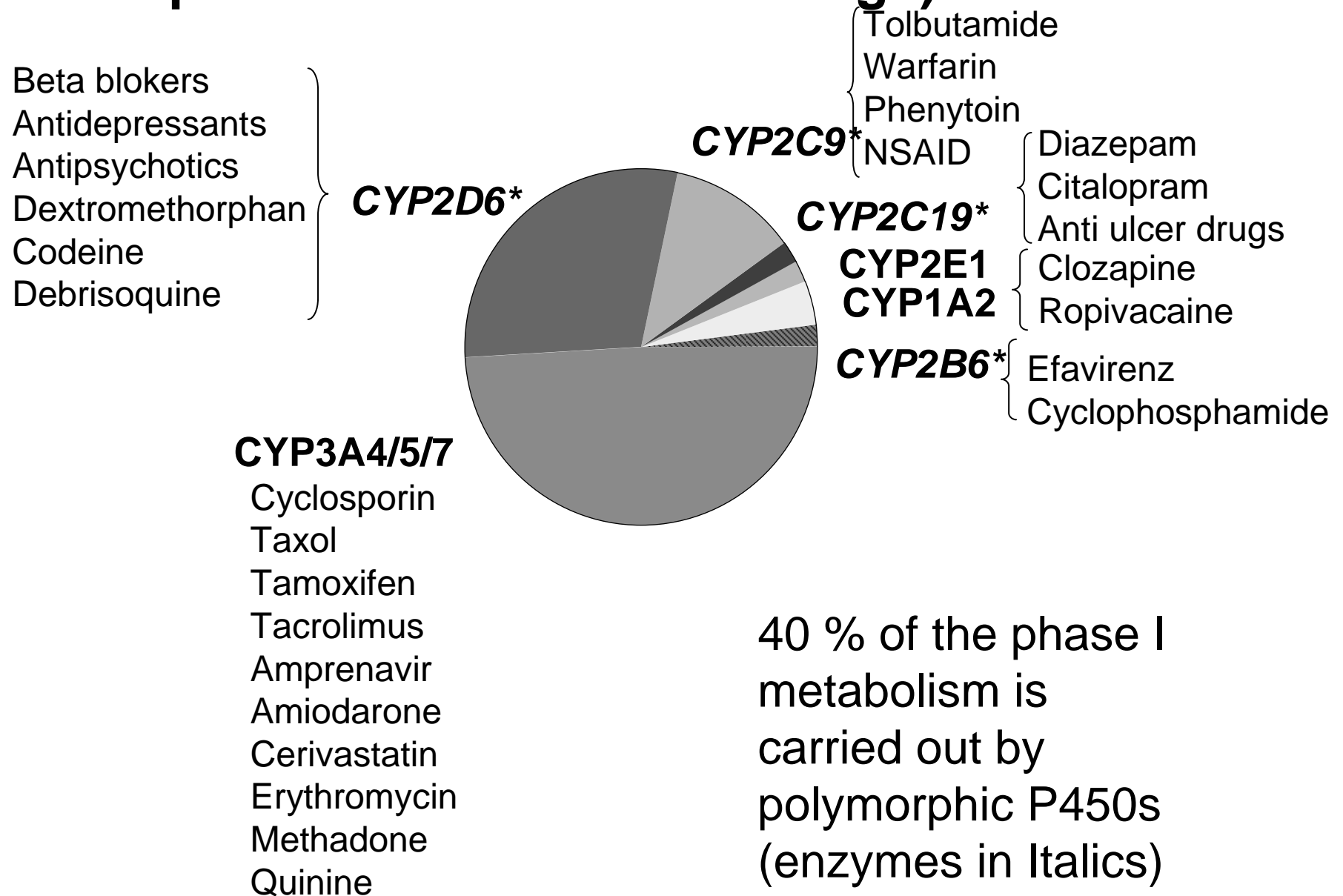
Drug-drug
drug-food
interactions



Drug-drug
drug-food
interactions



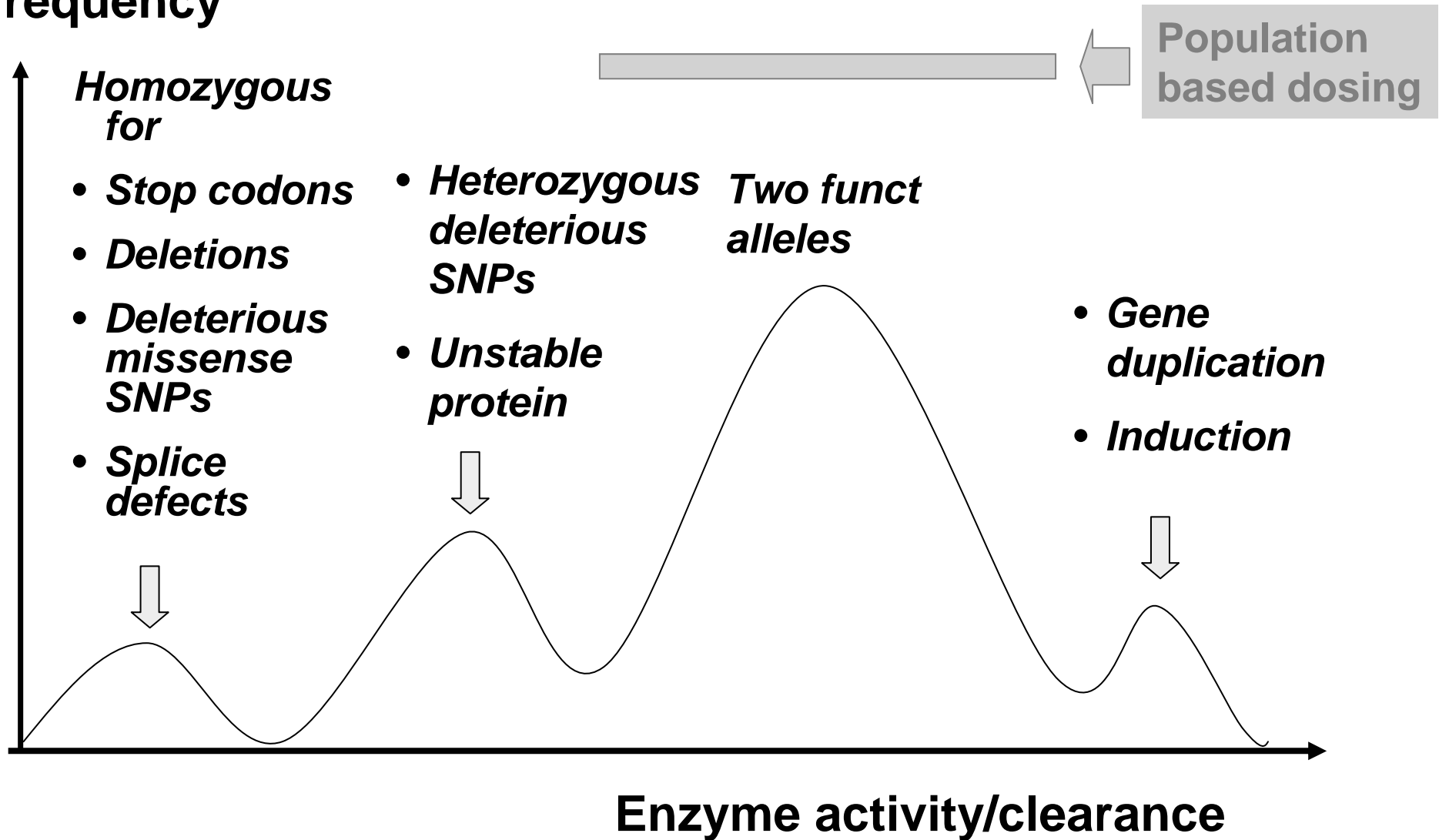
CYP dependent metabolism of drugs (80 % of all phase I metabolism of drugs)



Phenotypes and mutations

PM, poor metabolizers; IM, intermediate met; EM, efficient met; UM, ultrarapid met

Frequency





The Home Page of the Human Cytochrome P450 (CYP) Allele Nomenclature Committee

<http://www.imm.ki.se/CYPalleles/>

Webmaster: Sarah C Sim




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Advisory Board: Jürgen Brockmöller, Michel Eichelbaum, Seymour Garte, Joyce A. Goldstein, Frank J. Gonzalez, Fred F. Kadlubar, Tetsuya Kamataki, Urs A. Meyer, David R. Nelson, Michael R. Waterman, Ulrich M. Zanger.

Nomenclature files for human cytochrome P450 alleles:

CYP1A1, CYP1A2, CYP1B1, CYP2A6, CYP2A13, CYP2C9, CYP2C19, CYP2D6,
CYP2E1, CYP2J2, CYP2R1, CYP2S1, CYP3A4, CYP3A5, CYP3A7, CYP5A1,
CYP8A1, CYP21.

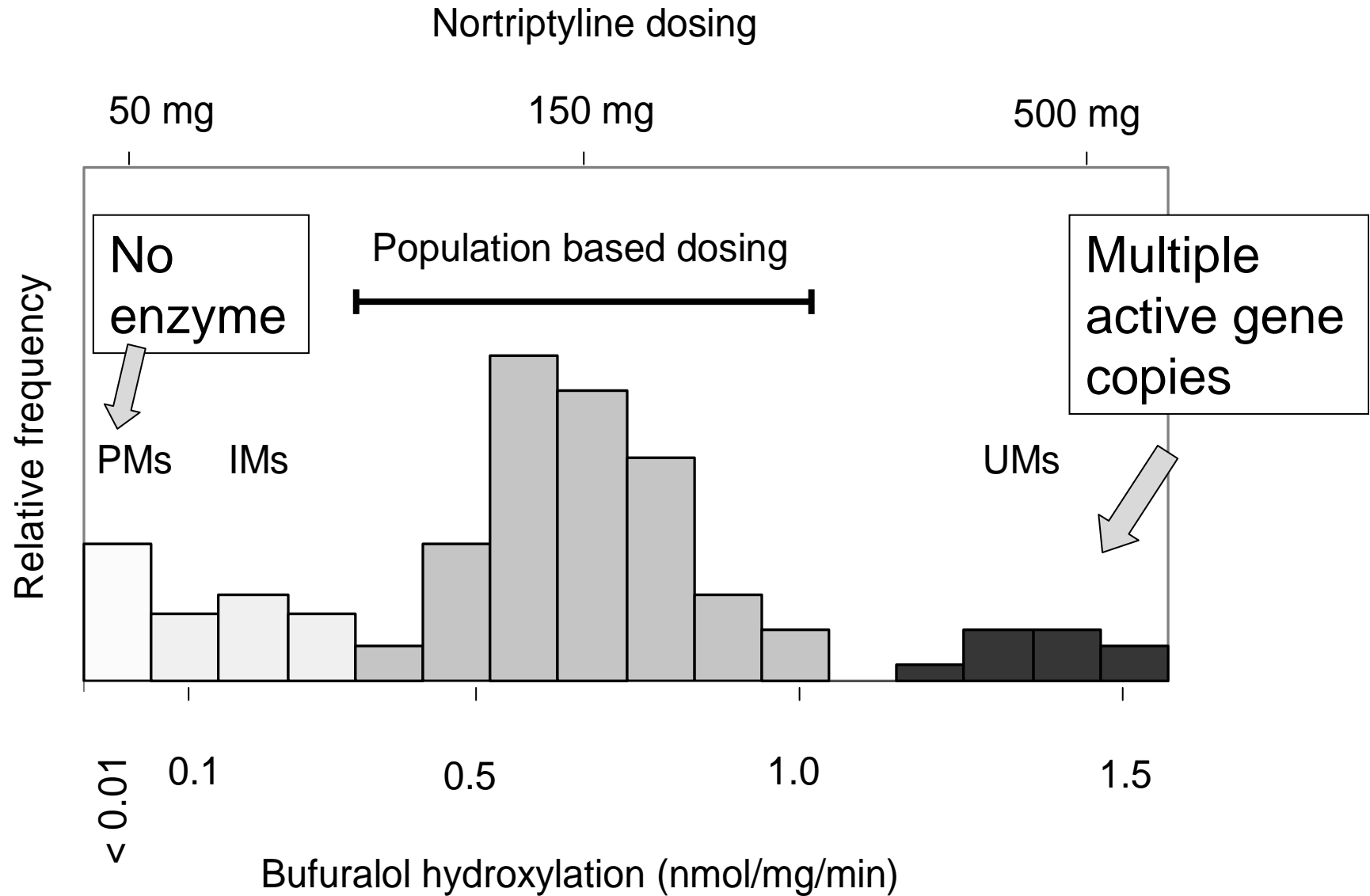


Cytochrome P450s and ADRs

- **59 % of drugs cited in ADR-studies are metabolised by polymorphic phase 1 enzymes - P450s account for 86 % of those**
- **Only 20 % of drugs which were substrates for nonpolymorphic enzymes were in the ADR reports**
- **CYP2D6 was involved in 38 % of all ADR reports**

Phillips et al
JAMA
286:2270-
2279, 2001

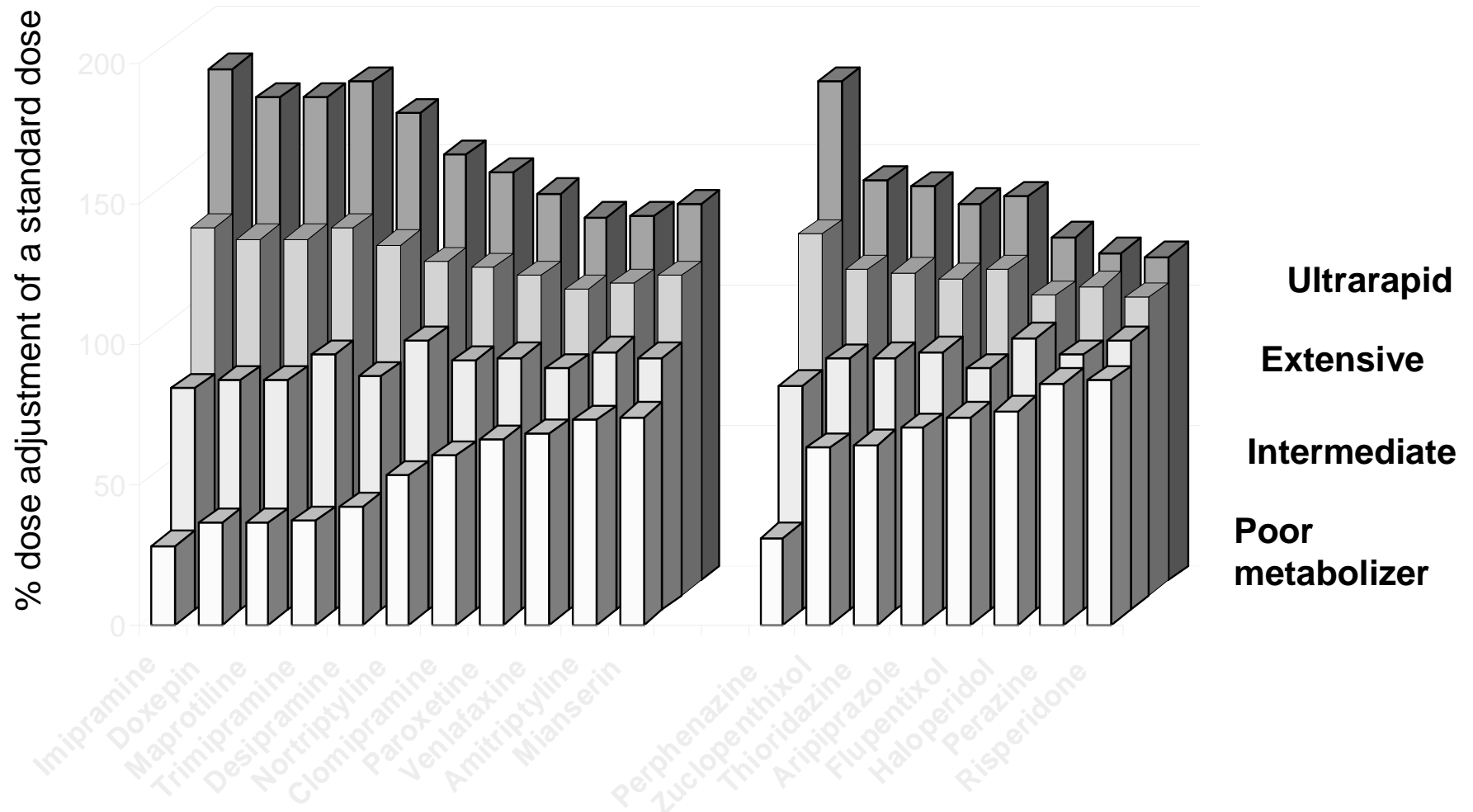
CYP2D6 and rate of metabolism in the European population



Based on the European population with 7 % PMs and 5.5% UMs overall

Ingelman-Sundberg, M. *Trends Pharmacol Sci* 2004, **25**:193-200

CYP2D6-based dose adjustments for antidepressants and antipsychotics



CYP2D6 and the European population

20-30 million subjects
have no CYP2D6
enzymes (PMs)



- Too slow drug metabolism
- Too high drug levels at ordinary dosage
- High risk for ADRs
- No response from certain prodrugs (e.g. codeine)

15-20 million subjects
have *CYP2D6* gene
duplications (UMs)



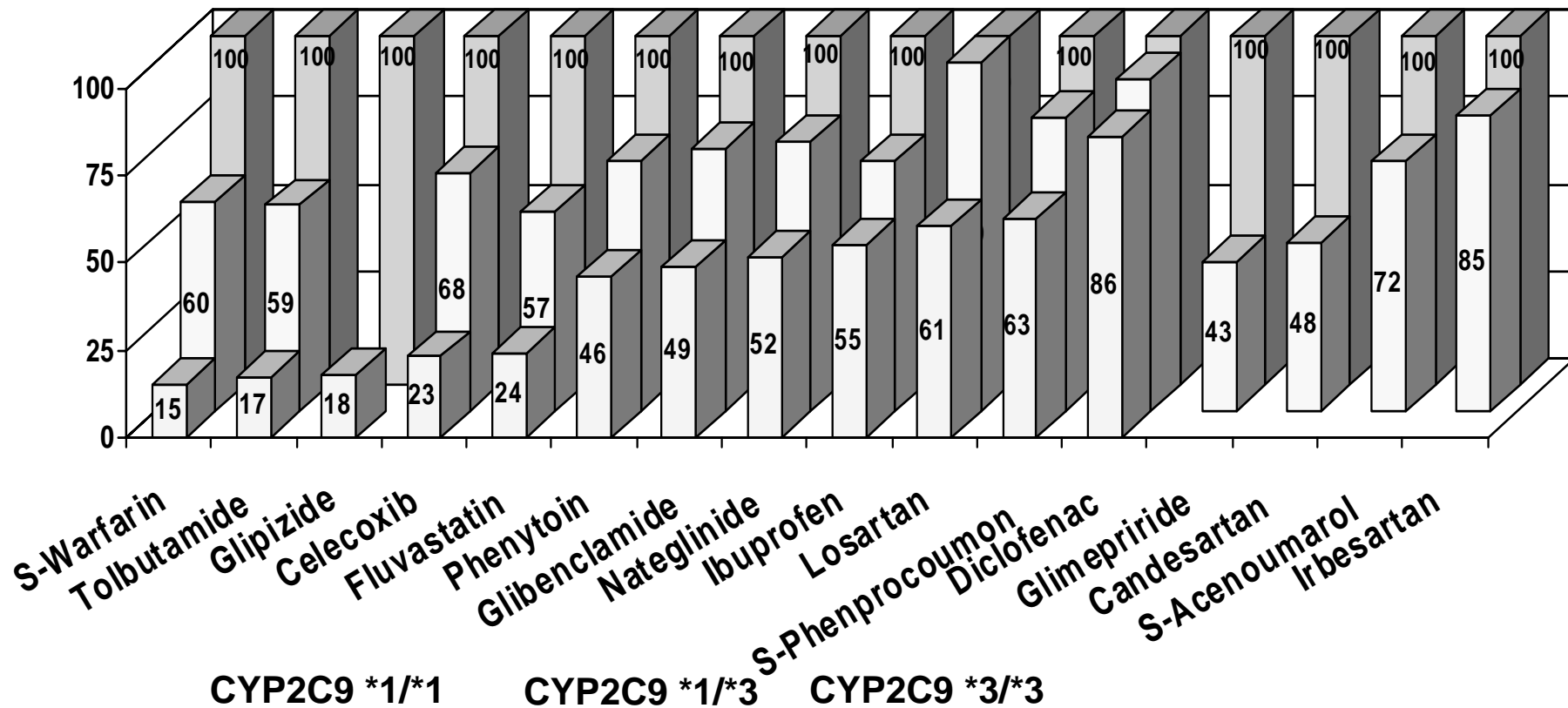
- Too rapid drug metabolism
- No drug response at ordinary dosage -
Non-responders

resulting in

Relevant for 15 % of all drugs used

Pharmacogenetics based dose adjustments: *CYP2C9**3

% reduction in oral clearance



Kirchheiner & Brockmüller, 2005

Warfarin dosage variation: 0.5-8 mg/day

CYP2C9



25% prediction



+ VKORC1



52% prediction



+ age and weight



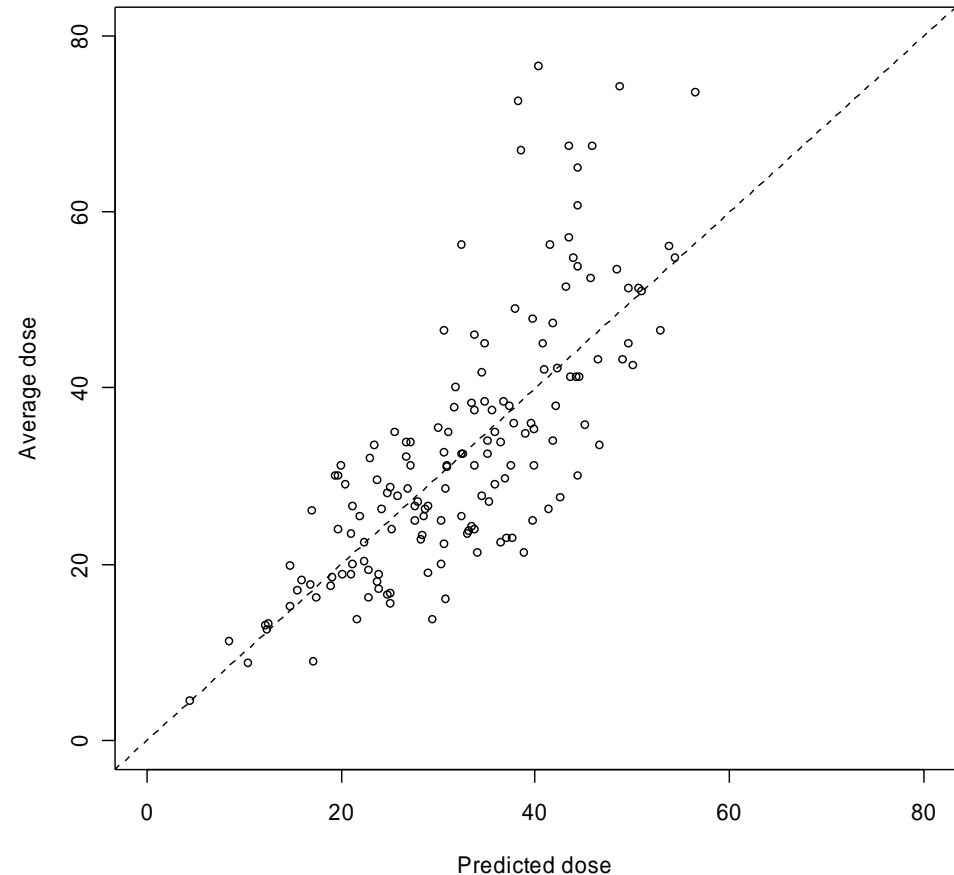
62% prediction

VKOR = vit K
epoxide reductase

Multiple regression model explains 61% of variance in warfarin dose

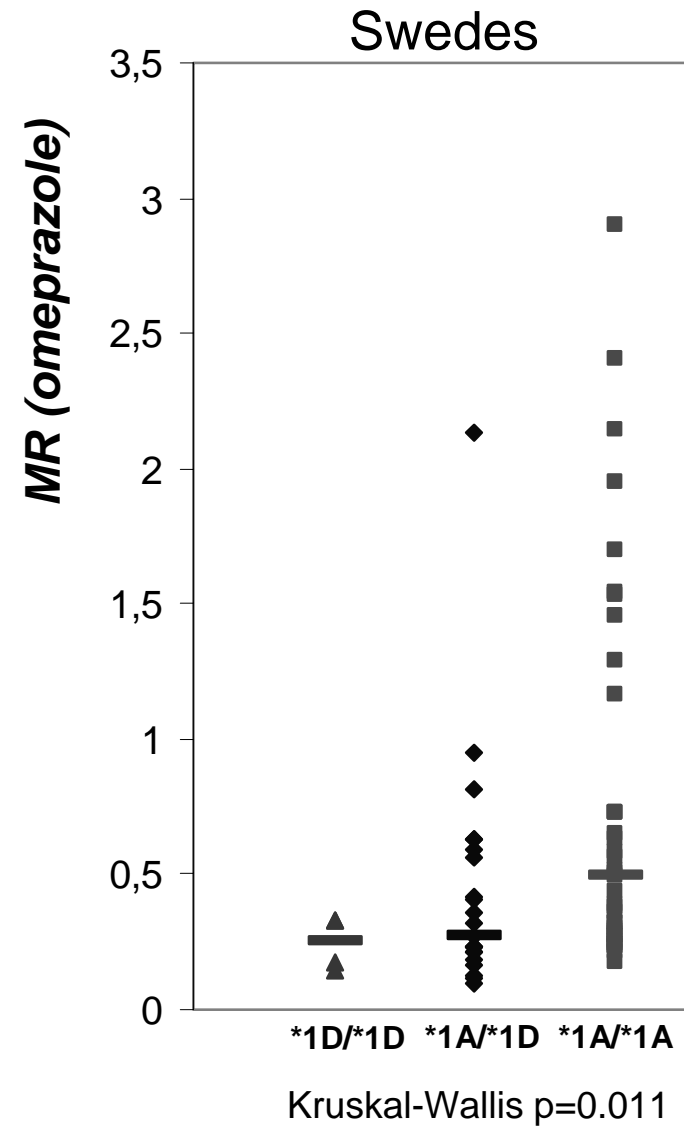
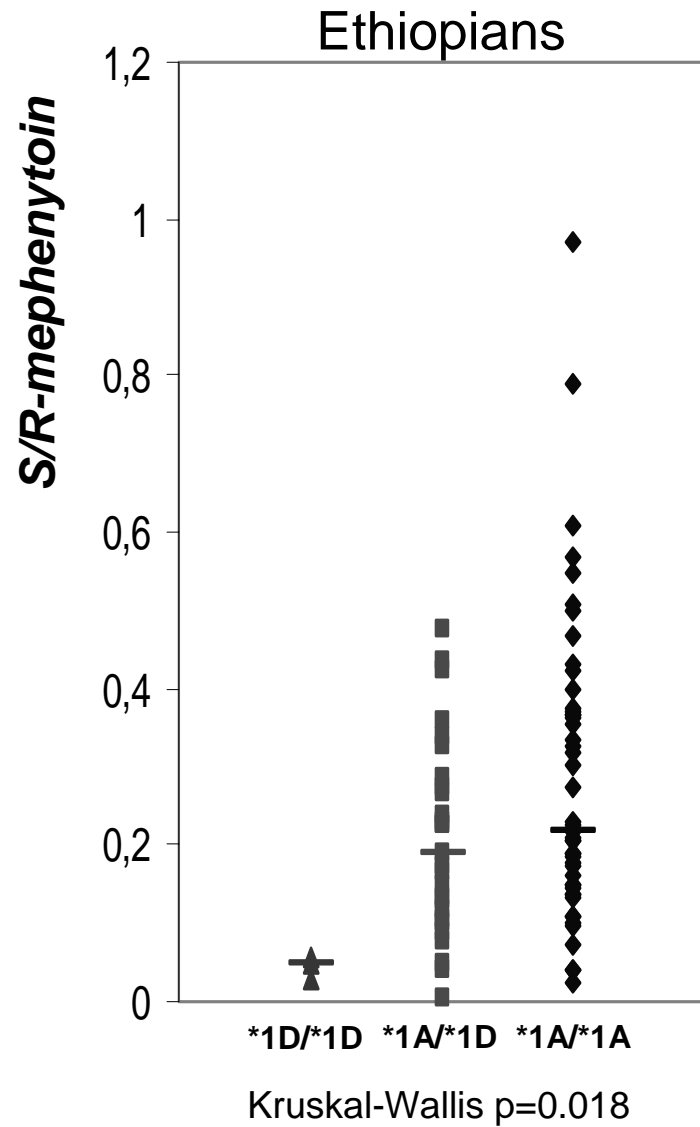
<i>Variables</i>	<i>Dose</i>	<i>p</i>
VKORC1	<.0001	
CYP2C9	<.0001	
PROC	0.0541	
Age	0.0002	
Bodyweight	0.0002	
Indication	0.0406	
Interaction	0.1018	

Total r^2 for the model = 0.614



Wadelius M, Chen LY, Eriksson N, Ghori J, Wadelius C, Bentley D, McGinnis R, Deloukas P.
Uppsala University, Sweden and the Wellcome Trust Sanger Institute, UK.

A novel ultrarapid *CYP2C19* allele (*CYP2C19*17*)



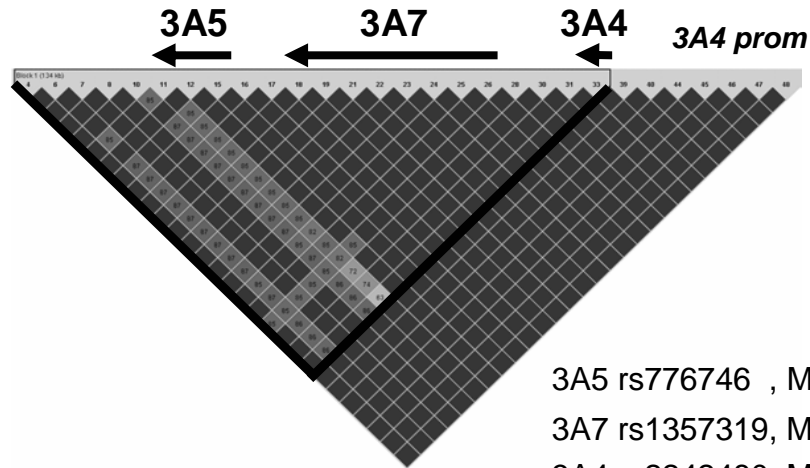
Frequency and effect of *CYP2C19**17 on clinical parameters

Population	Allele frequency	*1/*1	*1/*17	*17/*17
Swedes	20.1	155	80	9
Ethiopians	17.9	126	60	4
Tanzanians	16.3	63	23	3
Chinese*	5.0	54	6	0

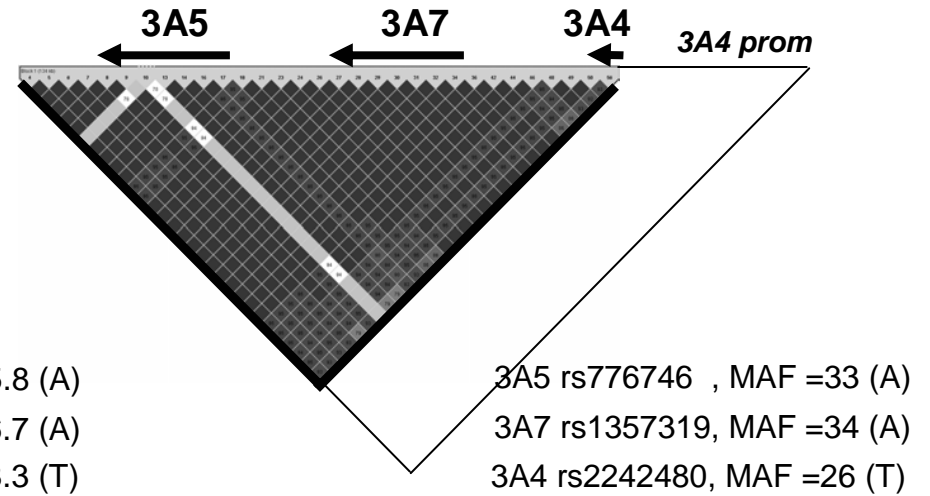
	*17/*17	*1/*17	*1/*1	*1/*2
n	4	23	38	16
Omeprazole MR	0.25±0.10	0.44±0.44	0.77±0.70	1.17±0.75
Pred omeprazole AUC	742±93	926±410	1240±658	1620±710
Predicted intragastric pH	3.50	3.75	4.20	4.50

CYP3A locus: INTERETHNIC DIFFERENCES

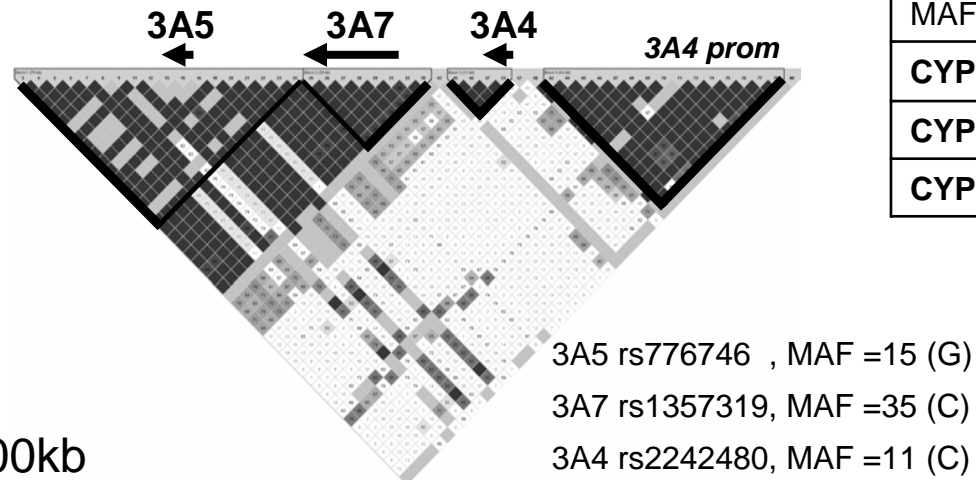
European



Chinese



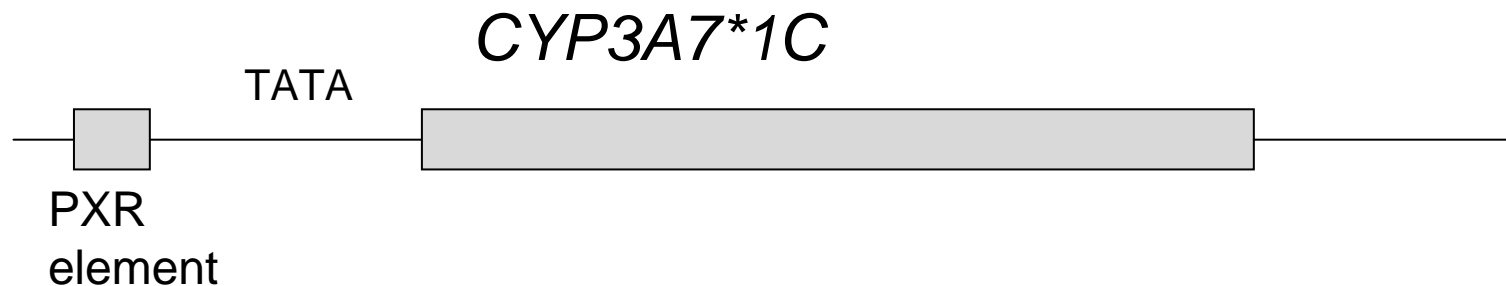
African



200kb

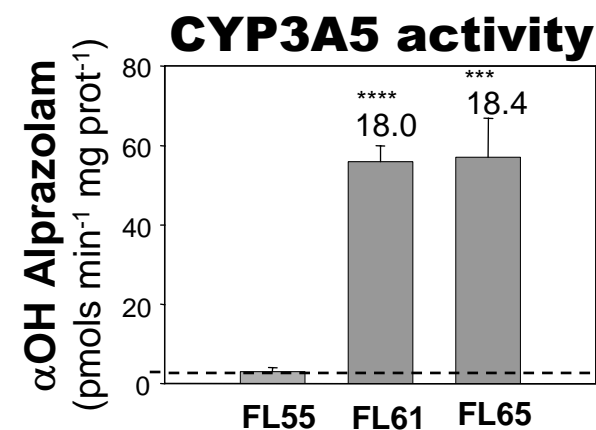
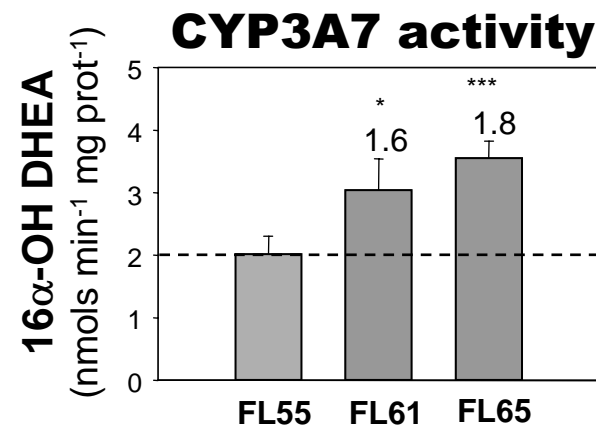
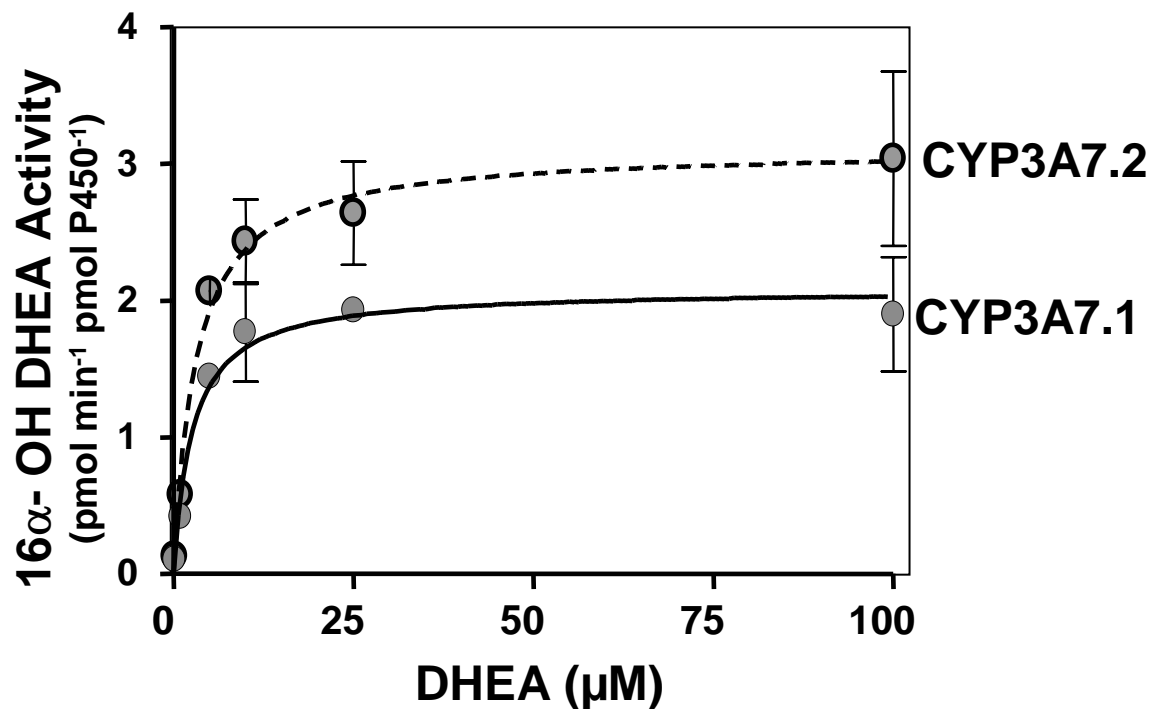
MAF (%)	European	Chinese	Africans
CYP3A5 rs776746	6 (A)	33 (A)	15 (G)
CYP3A7 rs1357319	7 (A)	34 (A)	35 (C)
CYP3A4 rs2242480	8 (T)	26 (T)	11 (C)

Polymorphically determined expression of CYP3A7 in human adult liver

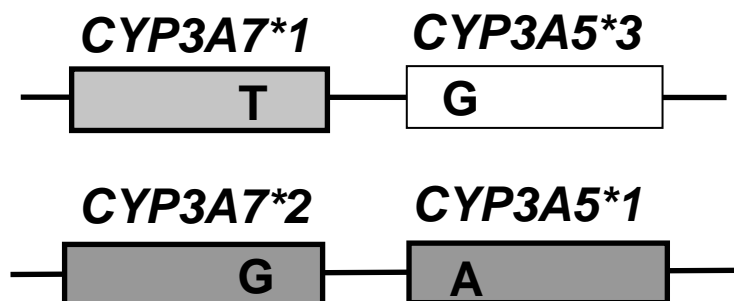


- one in 10 adult livers expressed CYP3A7 at 24-90 pmol/mg (9-36% to total CYP3A levels in these livers).
- 5/7 livers with *CYP3A7*1C* expressed CYP3A7 protein.
- In 57 livers CYP3A7 was present at 4 pmol/mg, *higher* than that of CYP3A5.

Sim SC et al. *Pharmacogenet Genomics*. 2005 Sep;15(9):625-31.



HAPLOTYPE



Estimated Frequency (%)

	Caucasians	Chinese	Africans
	90	72	20
	7	27	62

Genotyping for CYPs

Enzyme	substrates	phenotypes
CYP2B6	– cyclophosphamide, efavirenz	IM/EM
CYP2C9	– warfarin, antidiabetics, phenytoin, celecoxib	PM/IM/EM
CYP2C19	– antiulcer drugs, citalopram	PM/EM/UM
CYP2D6	– antidepressants, antipsychotics, codeine, tramadol, perhexiline, antiemetic drugs,	PM/IM/EM/UM

Examples of clinical impact of cytochrome P450 pharmacogenetics

Disease	Enzyme	Dose % of ctrl		Examples
		UMs	PMs	
Depression	CYP2C9			Bipolar disorders and valproate PMs and SSRIs Non-responders (UMs) and side effects of tricyclics (PMs)
	CYP2C19		40	
	CYP2D6	200	30	
Psychosis	CYP2D6	160	30	Haloperidol and parkinsonian side effects
Ulcer	CYP2C19		20	Dosing of PPIs pH and gastrin changes
Cancer	CYP2B6			Cyclophosphamide metabolism Non-response of antiemetic drugs (UMs)
	CYP2D6	250	60	
CV	CYP2C9		30	warfarin dosing (acenocoumarol) Irbesartan and blood pressure response; Perhexiline neuropathy and hepatotoxicity,
	CYP2D6	160	30	
Pain	CYP2D6			Codeine no response (PMs)
Epilepsia	CYP2C9			Phenytoin pharmacokinetics and side effects

Conclusions, pharmacogenetics in future drug treatment

- In 15-25 % of the cases of drug treatment genotyping will be very important for prediction of drug efficacy and drug toxicity
- For certain drugs genotyping will be relevant in 100 % of the cases
- Predictive genotyping might prevent 10-20 % of serious and fatal adverse drug reactions
- In 15-35 % drug metabolism is influenced by polygenic factors and difficult to foresee by genotyping
- In 50 % of the cases genetic factors are of limited importance

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Drug Reaction Testing

Do not alter the dosage amount or schedule of any drug you are taking without first consulting your doctor or pharmacists.

Research shows that of all the clinical factors such as age, sex, weight, general health and liver function that alter a patient's response to drugs, genetic factors are the most important. This information becomes even more crucial when you consider the fact that adverse reactions to prescription drugs are killing about 106,000 Americans each year -- roughly three times as many as are killed by automobiles. This makes prescription drugs the fourth leading killer in the U.S., after heart disease, cancer, and stroke.

We currently offer CYP2D6, CYP2C9, CYP2C19, and CYP1A2 screens that can help your physician or pharmacist predict your particular response to many prescription, OTC (over-the-counter) and herbal medicines including those used to treat depression, anxiety, seizures and psychoses; blood pressure, anticoagulation and other heart medicines; anti-diabetic agents, and many pain relievers. These include such important medications as Coumadin (Warfarin), Prozac, Zoloft, Paxil, Effexor, Hydrocodone, Amitriptyline, Claritin, Cyclobenzaprine, Haldol, Metoprolol, Rythmol, Tagamet, Tamoxifen, Valium, Carisoprodol, Diazepam, Dilantin, Premarin, and Prevacid (and the over-the-counter drugs, Allegra, Dytuss and Tusstat). [Click here to view a more complete list of drugs processed through these pathways.](#)

Approximately half of all Americans have genetic defects that affect how they process these drugs. There are four different types of metabolizers, and we all fall into one of these categories for the variable pathways in

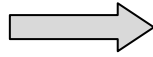
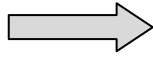
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Testing Consultant.

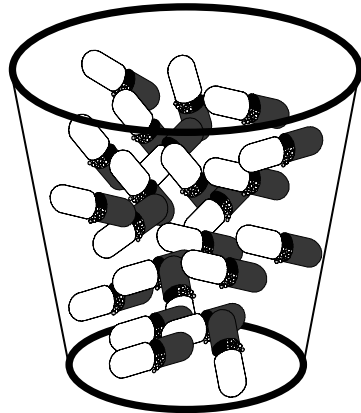


Future (cont...)

- Large prospective studies with well characterized patients on monotherapy
- Pharmacogenetics is used during drug development
- The urgent need concerns old drugs – industry will not finance such studies
- Pharmacogenetics is not used in the clinics unless required by regulations
- Development of guidelines of critical importance
- Implementation in the clinics after guidelines will occur



Specific
genotype



Specific
Drug



Combine

