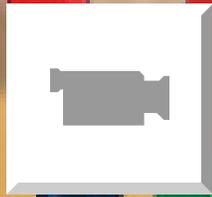




Pharmacogenomics from the Ground Up

Session 1: Concepts and Tools in Pharmacogenomics



The Human Genome

- The human genome contains 3164.7 million chemical nucleotide bases (A, C, T, and G).
- The average gene consists of 3000 bases, but sizes vary greatly, with the largest known human gene being dystrophin at 2.4 million bases.
- The total number of genes is estimated at 30,000.
- Almost all (99.9%) nucleotide bases are exactly the same in all people.
- The functions are unknown for over 50% of discovered genes.



The Human Genome

- **Less than 2% of the genome codes for proteins.**
- **Repeated sequences that do not code for proteins ("junk DNA") make up at least 50% of the human genome.**
- **Repetitive sequences are thought to have no direct functions. Over time, these repeats reshape the genome by rearranging it, creating entirely new genes, and modifying and reshuffling existing genes.**
- **During the past 50 million years, a dramatic decrease seems to have occurred in the rate of accumulation of repeats in the human genome.**



The Human Genome

- The human genome's gene-dense "urban centers" are predominantly composed of the DNA building blocks G and C.
- In contrast, the gene-poor "deserts" are rich in the DNA building blocks A and T. GC- and AT-rich regions usually are the light and dark bands we see on chromosomes.



The Human Genome

- Genes appear to be concentrated in random areas along the genome, with vast expanses of noncoding DNA between.
- Stretches of up to 30,000 C and G bases repeating over and over often occur adjacent to gene-rich areas, forming a barrier between the genes and the "junk DNA." These CpG islands are believed to help regulate gene activity.
- Chromosome 1 has the most genes (2968), and the Y chromosome has the fewest (231).



Session Goals

- **Concepts in Pharmacogenomics**
- **Metabolic Enzymes**
- **Pharmacogenomics of CYPs**
- **Clinical Examples**
- *Pharmacogenomics Jeopardy!*



Concepts in Pharmacogenomics



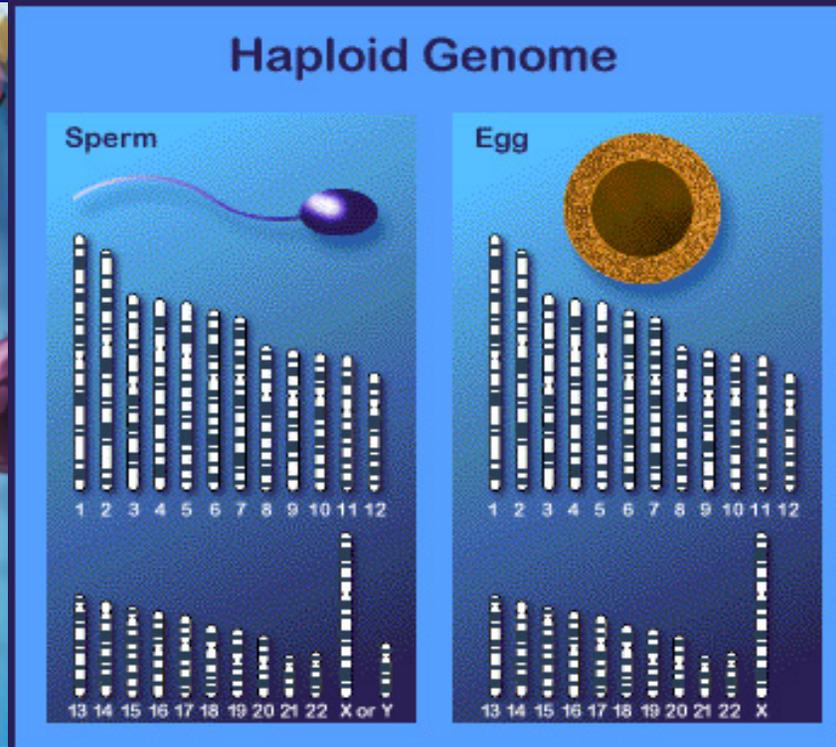
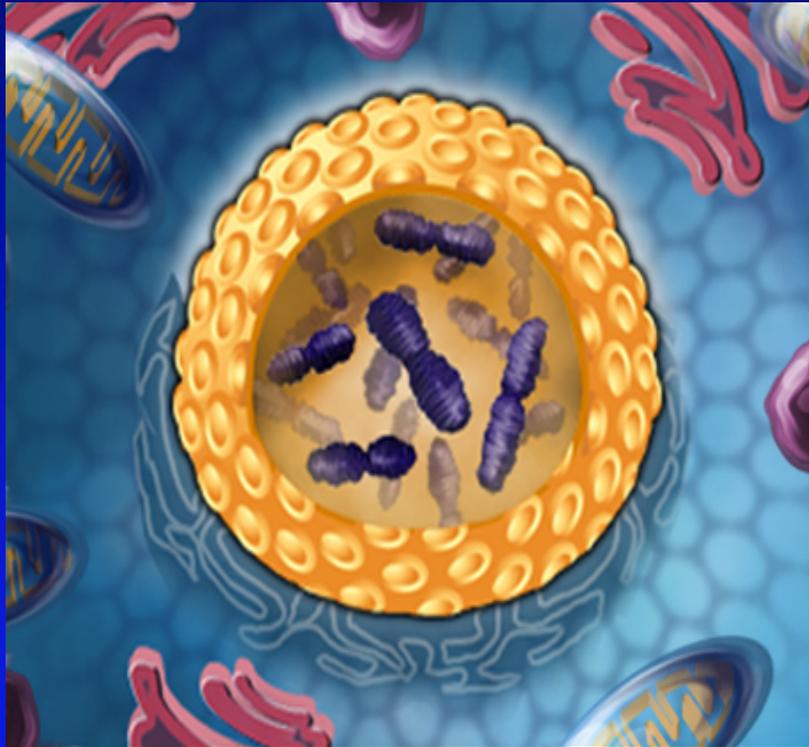
Pharmacogenetics and Pharmacogenomics

- *Pharmacogenetics*: the effect of genetic variation on drug response, including disposition, safety and tolerability, and efficacy.
- *Pharmacogenomics*: the application of genome science (genomics) to the study of human variability in drug response.
- We will study in the second session of this course *pharmacogenomics* applied to Epidermal Growth Factor Receptor (EGFR) drugs such as Tarceva, Iressa, and Erbitux.



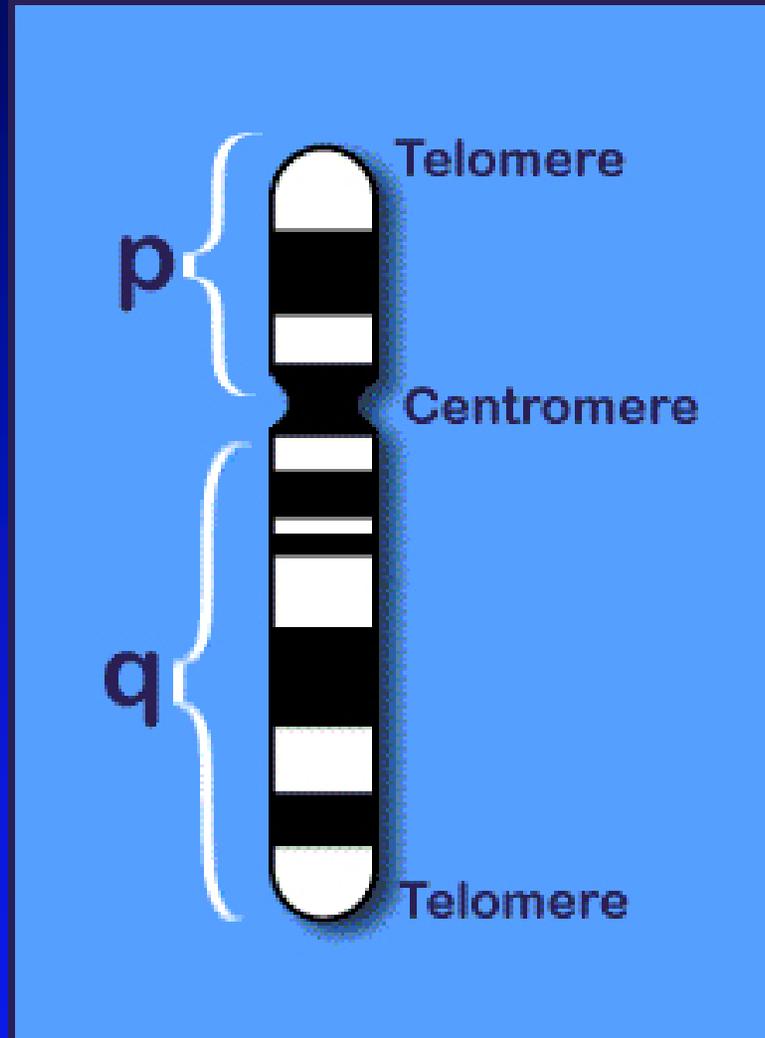
Genomic Organization: *Chromosomes*

- Every human cell with the exception of the gametes contains 23 pairs of chromosomes.



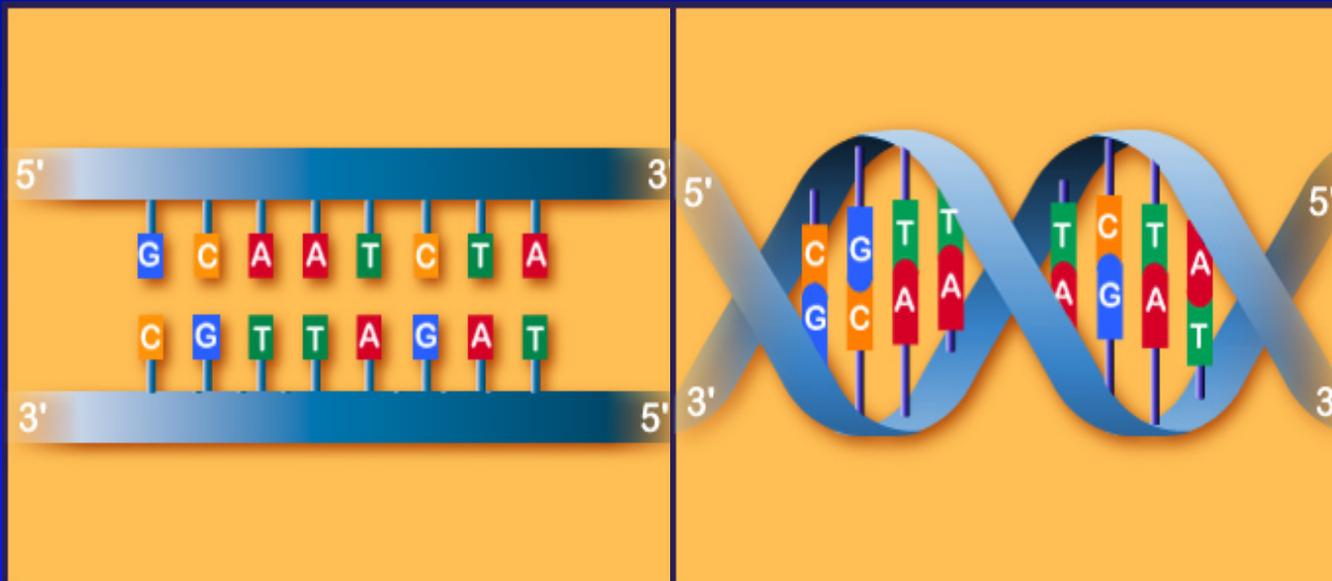
Genomic Organization: *Chromosome Structure*

- **Chromosomes carry all of the genetic material coding for all the proteins in every cell.**
- **Chromosomes consist of DNA tightly wound around special protein structures called histones.**



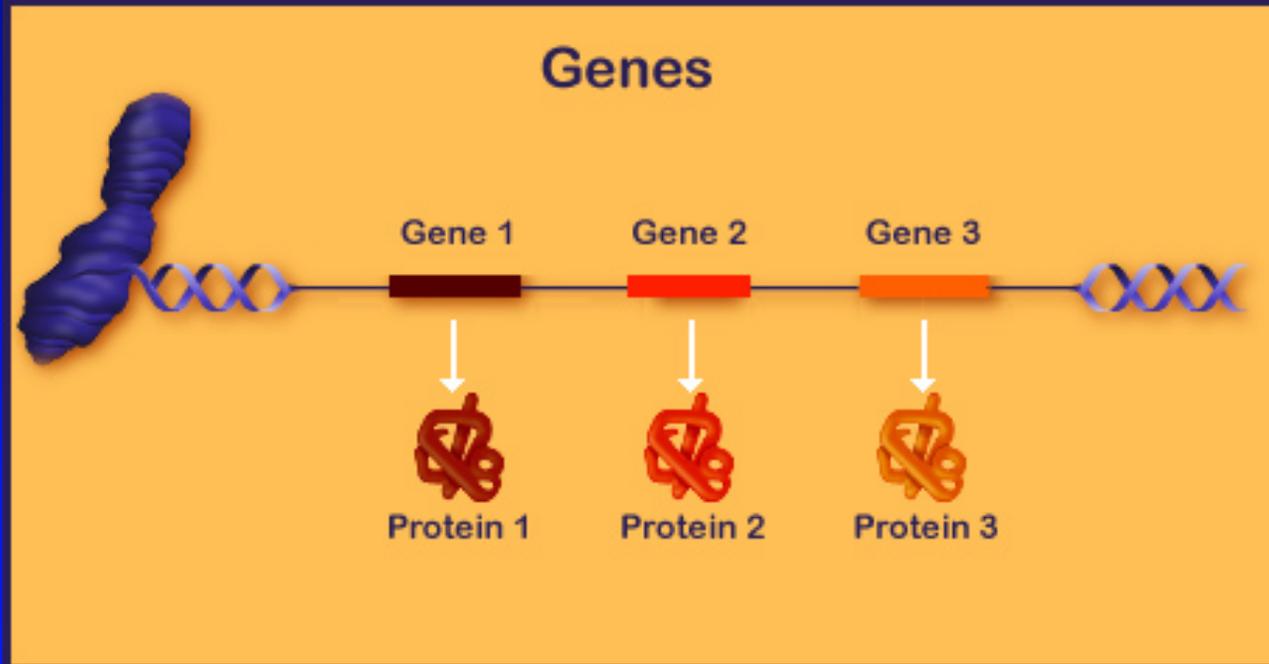
Structure of DNA

- DNA is comprised of a string of 4 *nucleotide bases*, called A, G, T and C, that are linked together in a structure called the double helix. Bases on opposite strands are always matched A-T and C-G.



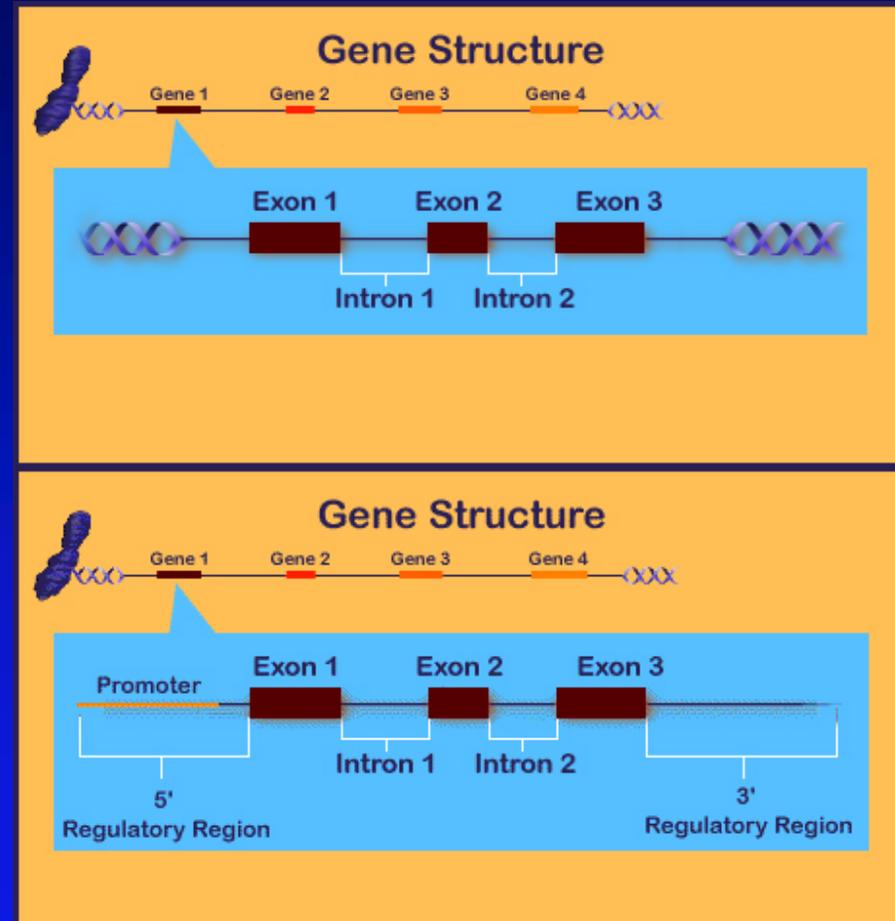
Structure of Genes

- A segment of DNA containing all of the information needed to encode for one protein is called a *gene*. The order and sequence of the base pairs in a gene determine which protein is made.



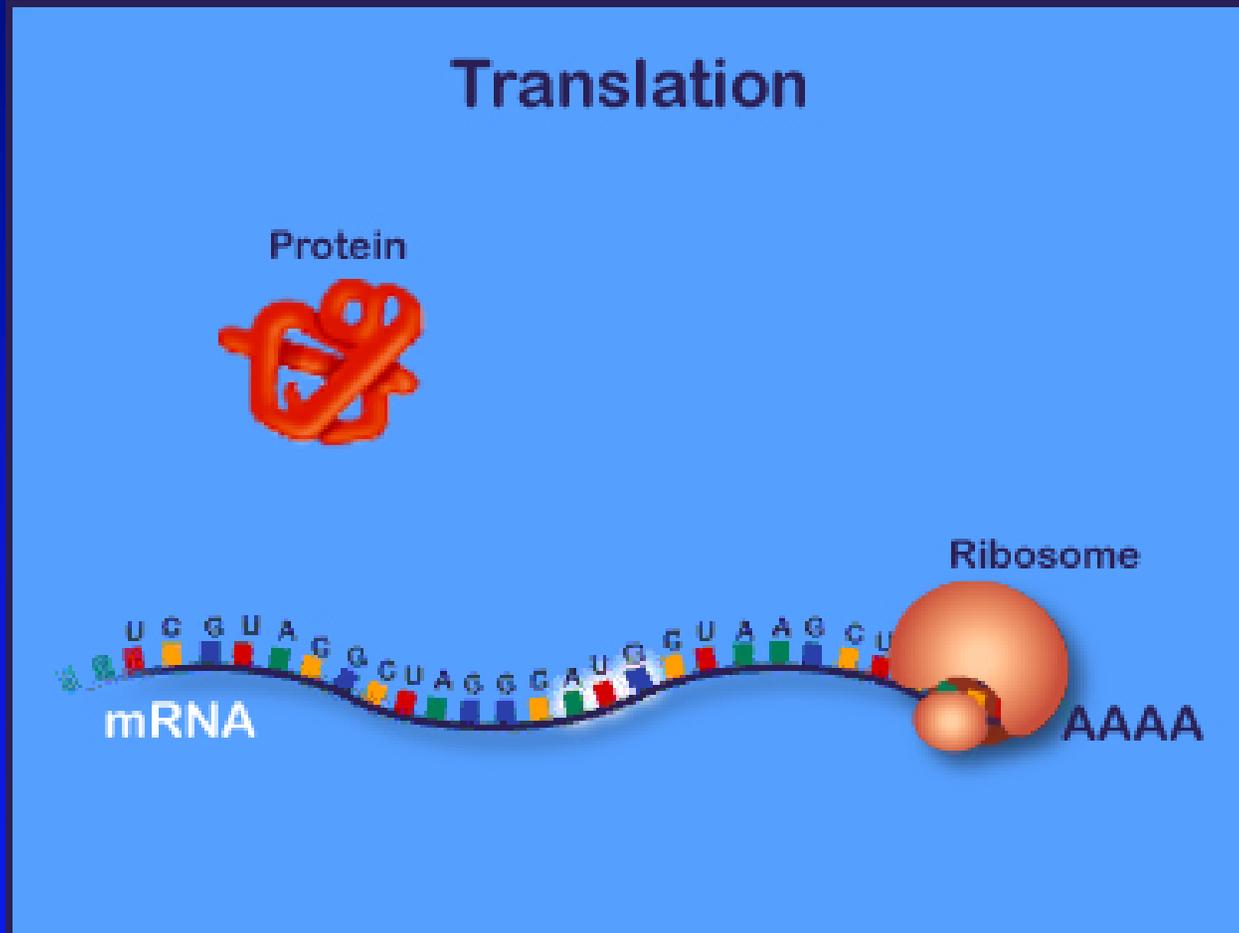
Transcription of Genes

- The transcription of DNA into messenger RNA is often assembled from discontinuous sequences in the genome called *exons*, which are separated by sequences called *introns*.
- This process is referred to as *splicing*. A genomic sequence can yield more than one splicing product.
- Transcription can be controlled through regulatory sequences known as the *promoter sequences*.



Putting It All Together: *Translation into Protein*

- Translation into protein sequences is accomplished at the ribosome.



Gene Expression in Drug Development

Gene Expression Studies

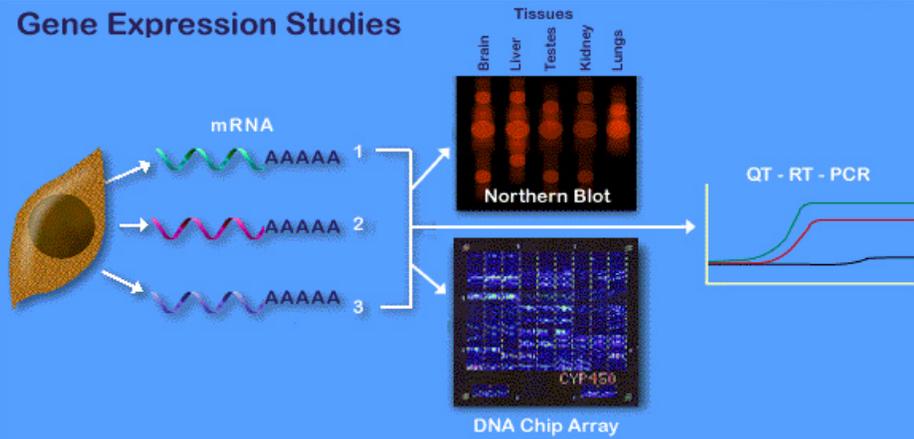
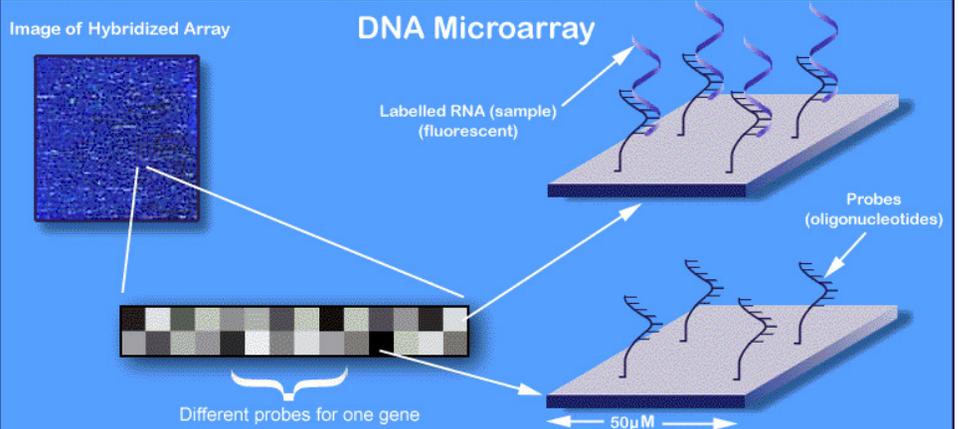
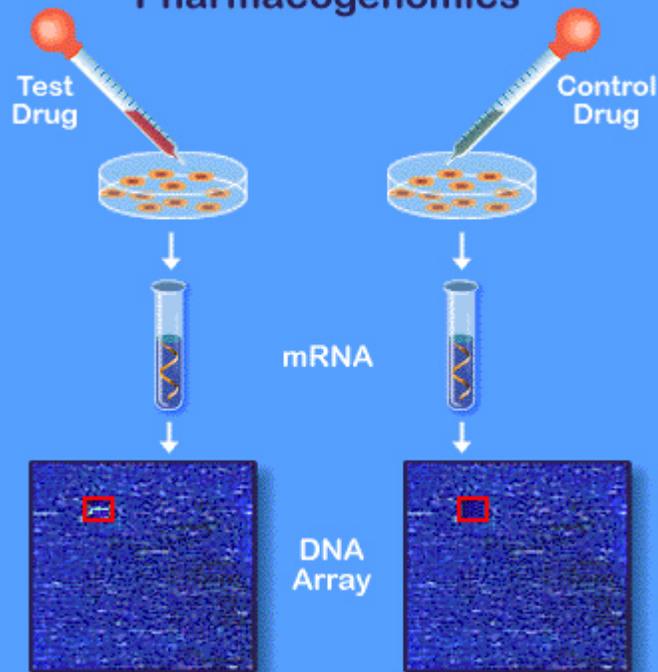


Image of Hybridized Array

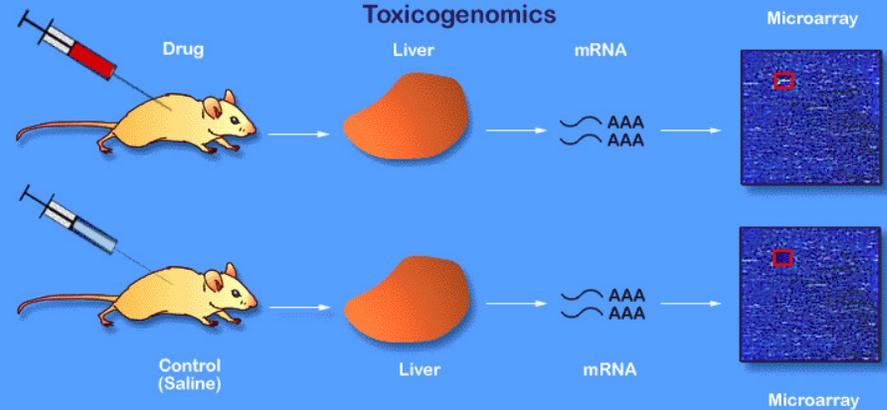
DNA Microarray



Pharmacogenomics



Toxicogenomics



The Genetic Code: *Translation into Protein Building Blocks*

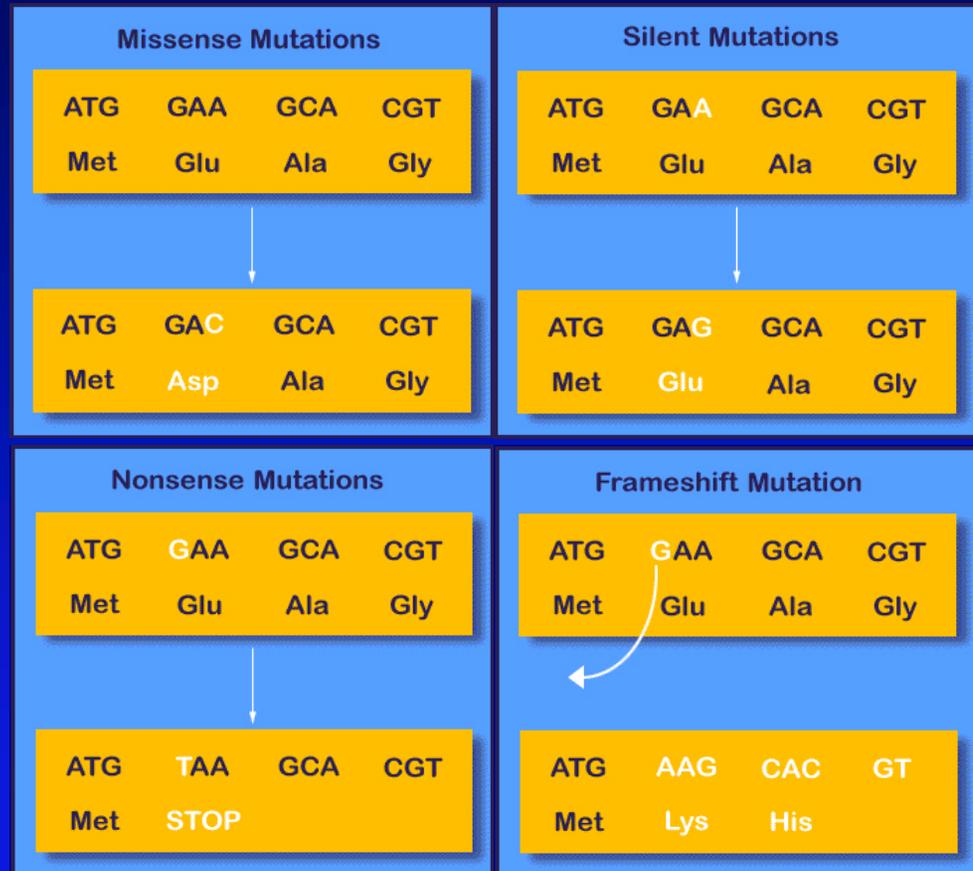
- The Genetic Code has some redundancy. But things can – and do – go wrong...

		Second Base							
		U	C	A	G				
U	UUU	Phe	UCU	Ser	UAU	Tyr	UGU	Cys	U
	UUC		UCC		UAC		UGC		C
	UUA	Leu	UCA		UAA	Stop	UGA	Stop	A
	UUG		UCG		UAG	Stop	UGG	Trp	G
C	CUU		CCU	Pro	CAU	His	CGU		U
	CUC	Leu	CCC		CAC		CGC	Arg	C
	CUA		CCA		CAA	Gln	CGA		A
	CUG		CCG		CAG		CGG		G
A	AUU		ACU	Thr	AAU	Asn	AGU	Ser	U
	AUC	Ile	ACC		AAC		AGC		C
	AUA		ACA		AAA	Lys	AGA	Arg	A
	AUG	Met / Start	ACG		AAG		AGG		G
G	GUU		GCU	Ala	CAU	Asp	GGU		U
	GUC	Val	GCC		GAC		GGC	Gly	C
	GUA		GCA		GAA		GGA		A
	GUG		GCG		GAG	Glu	GGG		G



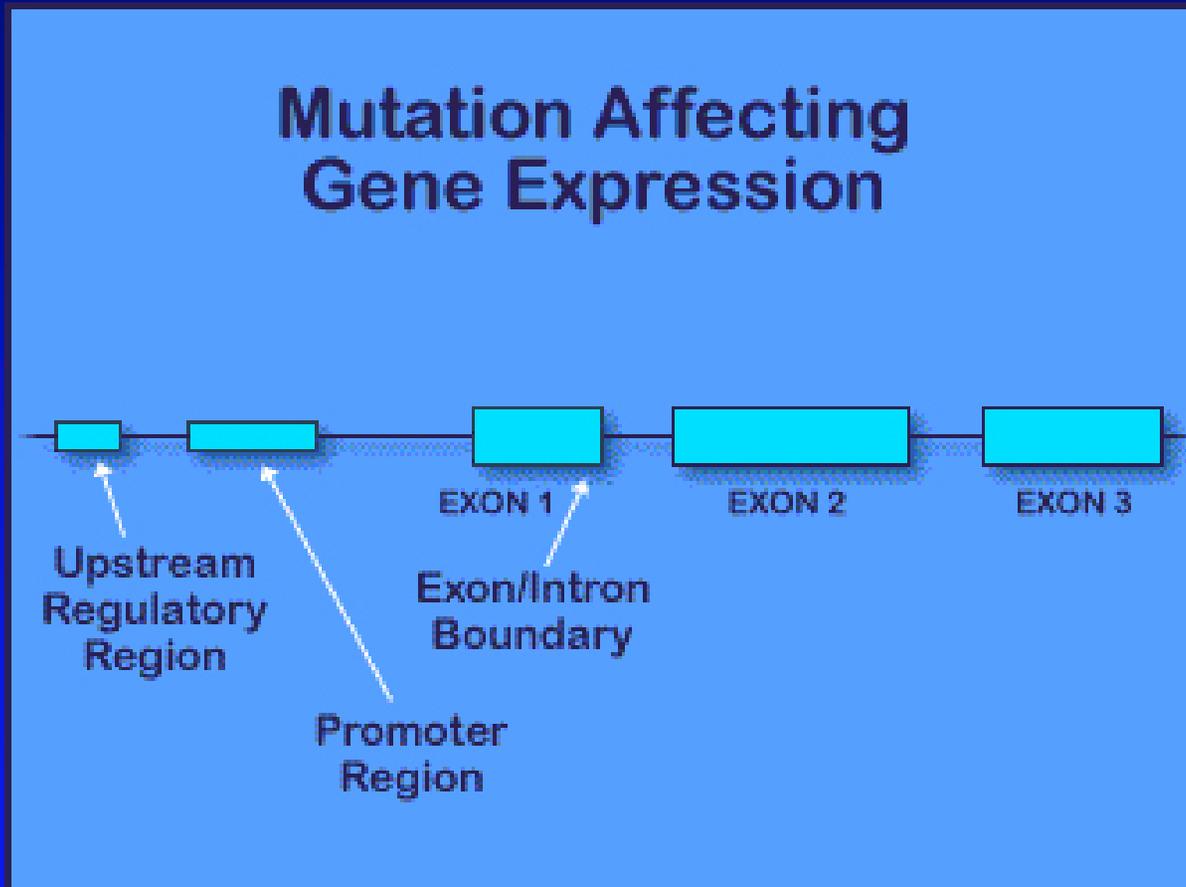
Mutations in the Genome

- One in every 1200 bases may be different in any two humans. This variation, which is called a *polymorphism*, is largely responsible for differences between how humans respond to drugs. Several types of mutations are associated with these variations:



Mutations in the Genome

- Mutations can also affect gene expression when they occur in regulatory or promoter sequences or in the exon/intron boundary.



Clinically Important Polymorphisms

- **Alter amino acid sequence of the corresponding protein (functional).**
- **Occur preferably within the exon or in the promoter region of the gene.**
- **Reasonable prevalence in the target population.**

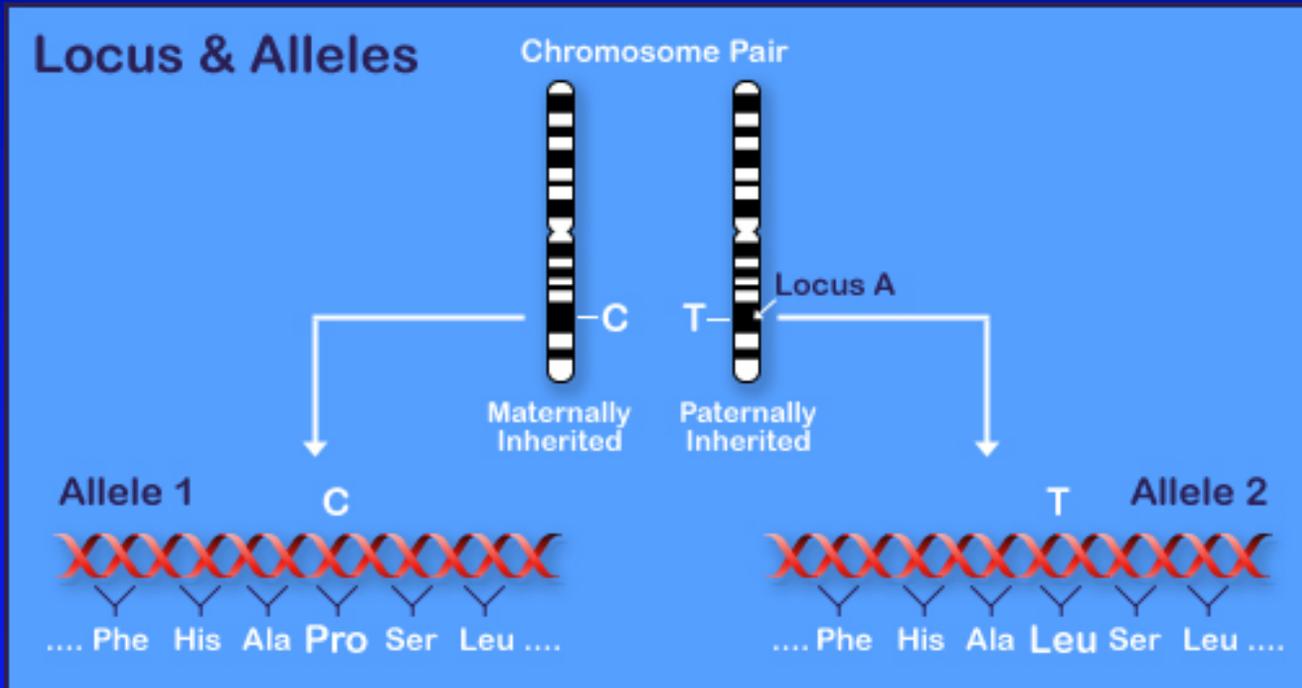


Alleles

- An allele is one of alternative forms at a genetic locus on a single chromosome. For loci in most of the genome, a human has two chromosomes, which may carry the same or two different alleles.

Genotype

- The specific genetic variants at one or more loci of an individual. Typically, the two chromosomal copies of a polymorphic site. For instance, A/A or A/G or G/G.



How to Calculate Allele and Genotype Frequencies: *The Hardy-Weinberg Equation*

$$(p^2 + 2pq + q^2) = 1, \text{ where } (p + q) = 1$$

- **Example: *cystic fibrosis***

- 1/2000 = 0.0005 live births to caucasian parents have cystic fibrosis
 - » $q^2 = 0.0005$ (0.05% of the population are *homozygotes* for the recessive *q* allele)
 - » $q = 0.02$ (the *allele frequency* is 2%)
- $p = 1 - q = 0.98$ (the *allele frequency* is 98%)
 - » $p^2 = 0.96$ (96% of the population are *homozygotes* for the dominant *p* allele)
- $2pq = 0.0392$ (3.92% of the population are *heterozygotes*, or asymptomatic carriers of the cystic fibrosis gene)



Metabolic Enzymes



Drug Metabolizing Enzymes

- **Drug Metabolism**
 - Small number of metabolic pathways
 - Microsomal enzymes
 - Liver and small intestine
- **Drugs**
 - Lipophilic
 - Low water solubility
 - Metabolizing enzymes increase water solubility
 - Water-soluble metabolites are easier to excrete



Phases of Drug Metabolism

Enzyme reaction	Metabolic reaction	Examples of enzymes
Phase I reactions		
Oxidation	Introduces hydroxyl, epoxide and ketone groups Shortens alkyl side chains Converts alcohols to aldehydes and acids	Alcohol and aldehyde dehydrogenases Amine oxidases Cytochromes P450
Reduction	Introduces hydrogen into ketones and nitro groups	Nitro- and azo-reductases
Hydrolysis	Breaks down esters to alcohols and acids	Esterases
Phase II reactions		
Acetylation	Adds acetate to polar sites	Acetyltransferases
Amino acid conjugation	Adds amino acids to polar sites	Glutathione transferases
Glucuronidation	Adds sugars to polar sites	Glucuronyl transferases
Methylation	Adds methyl groups to polar sites	Methyltransferases
Sulphation	Adds inorganic sulphate to polar sites	Sulphotransferases

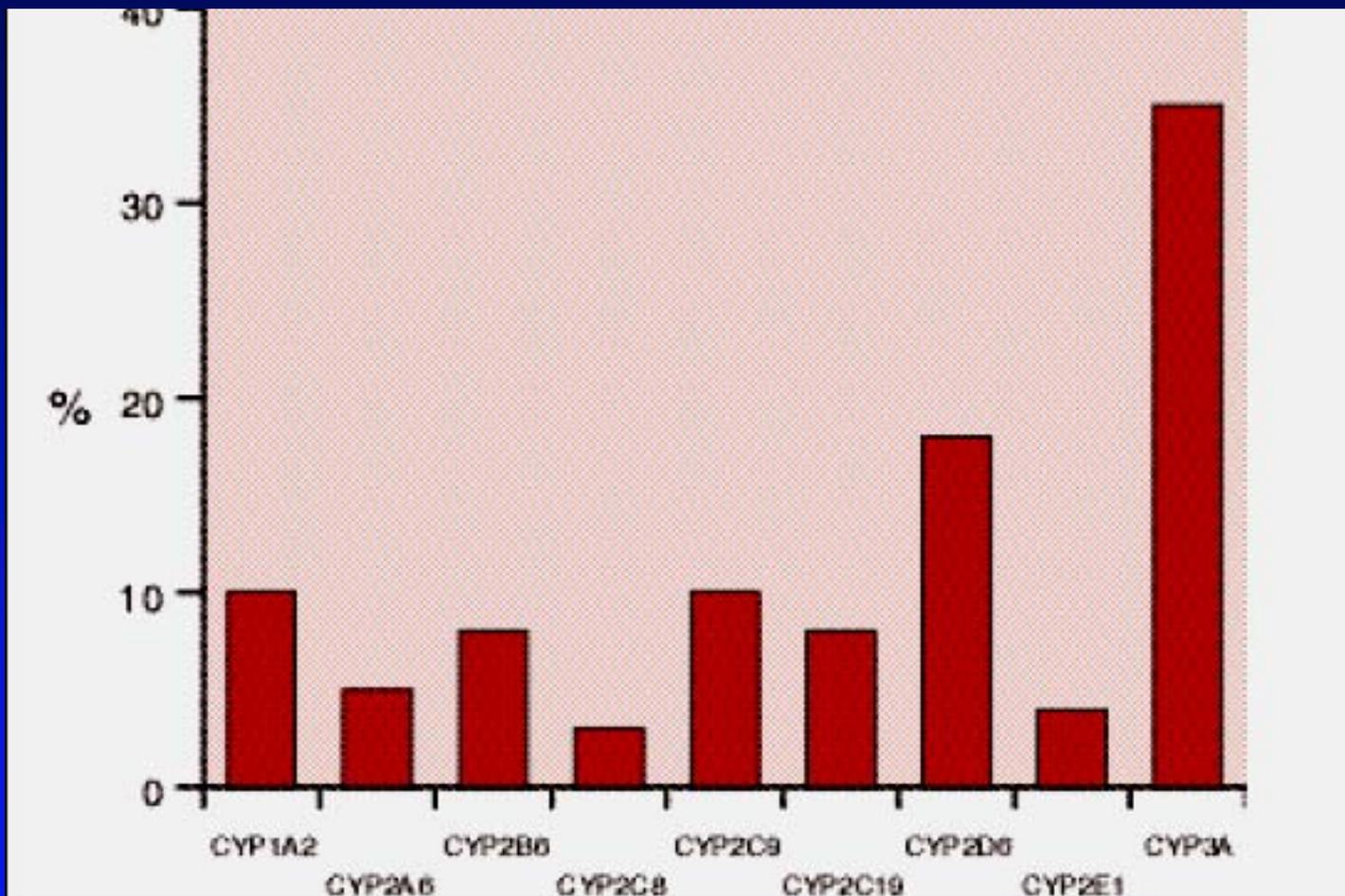


P450 Enzymes

- **57 Different Active Genes**
- **17 different families**
- **CYP1, CYP2 and CYP3 are primarily involved in drug metabolism.**
- **CYP2A6, CYP2B6, CYP2C9, CYP2C19, CYP2D6, CYP2E1 and CYP3A4 are responsible for metabolizing most clinically important drugs**
- **Reactions**
 - **Aliphatic oxidation**
 - **Aromatic hydroxylation**
 - **Sulphoxide formation**
 - **N-oxidation and N-hydroxylation**
 - **N-/O-/S-dealkylation**
 - **Oxidative or reductive dehalogenation**



P450s: *Who Work Most Often*



Major P450 Families and Their Isoforms

CYP family/subfamily	Isoform*	Organ localisation	Polymorphic substrates
1A	1A1	Primarily extrahepatic	Yes†
	1A2	Primarily hepatic	Yes†
	1B1	Primarily extrahepatic	Yes†
2A	2A6	Hepatic and extrahepatic	Coumarin, nicotine
	2A7	Hepatic and extrahepatic	
	2A13	Primarily extrahepatic	
2B	2B6	Primarily hepatic	Yes†
2C	2C8	Primarily hepatic	
	2C9	Primarily hepatic	Tolbutamide, warfarin
	2C18	Primarily hepatic	
	2C19	Primarily hepatic	Diazepam, propranolol
2D	2D6	Primarily hepatic	Codeine, nortriptyline, ecstasy
2E	2E1	Primarily hepatic	Yes†
2F	2F1	Primarily extrahepatic	
2G	2G1	Olfactory mucosa	
2J	2J2	Primarily extrahepatic	
3A	3A4	Primarily hepatic	Yes†
	3A5	Primarily hepatic	Yes†
	3A7	Foetal liver	
	3A43	Primarily hepatic	
4A	4A11	Primarily hepatic	
4B	4B1	Primarily extrahepatic	

*The role in drug metabolism of additional CYP isoforms in these families (eg, CYP4F2, CYP4F8) is not well described

† Multiple variants exist for these polymorphic isoforms but the consequences on the substrates they metabolise are not yet fully documented — see <http://www.imm.ki.se/CYPalleles/default.htm>



Nuclear Receptors in Enzyme Induction of Drug Metabolizing Enzymes

Receptor	CYPs induced	Inducers	Other enzymes induced
Aryl hydrocarbon (Ah/XRE*)	CYP1A1, CYP1A2, CYP1B1	Cigarette smoking, barbecued food and omeprazole	Glutathione transferases (GST) Glucuronyltransferases (UGT)
Constitutive androstane (CAR)	CYP2B6, CYP2A6, CYP3A†	Phenobarbital, amobarbital, secobarbital butobarbitone, heptobarbitone, glutethimide, promethazine	Epoxide hydrase, GST, UGT Cytochrome P450 reductase
Pregnane X (PXR)	CYP3A4, CYP3A5, CYP3A43†, CYPs2A†, CYPs2C†, CYP2E1†	Rifampicin, carbamazepine, dexamethasone, phenylbutazone, phenytoin, sulfadimidine, sulfinpyrazone; phenobarbital†, St John's Wort	Cytochrome P450 UGT†
Peroxisome proliferator-activated (PPAR)	CYP4A	Fibrate anti-hyperlipidaemics	Cytochrome P450 reductase
TR	CYP reductase	Thyroid hormone (T3)	
Unknown	2E1	Ethanol and chloral hydrate, isoniazid	

*XRE = xenobiotic responsive element (nuclear binding site) for the Ah receptor

†These items are suspected but not yet proven



Drug	Associated CYP	Drug	Associated CYP	Drug	Associated CYP
Pharmaceuticals					
Afenitil	2D6, 3A4	Bacamide	1A2, 2D6	Naoripityline	2D6
Alprazolam	3C19, 3A4	Racemate	1C9, 3C19, 3A4	Olansapine	1A2
Amiodarone	1A2, 2C9, 3A4	Racemate	1C9, 3C19, 2D6, 3A4	Onapazole	1A2, 3C19, 3A4
Amitriptyline	3C9, 2D6, 3A4	Rapheazine	2D6	Ondansetron	1A2, 2D6
Amprenavir	3A4	Rarbiposin	1C9	Orphenadrine	2D6
Antemide	3A4	Ratanide	1A2, 3A4	Oxibrocain	1A2
Atomoxetine	3C9, 3A4	Raxatin	1C9	Pachicol	3C8, 3A4
Azaxidine	3A4	Raxoxamine	1A2, 3C19, 2D6, 3A4	Paracetamol	3C19
Bromocriptine	3A4	Genitocid	1A4	Paracetamol	1A2, 3E1, 3A4
Budonide	3A4	Gibercumide	1A4	Paroxetine	1A2, 2D6
Bupropion	2D6	Granisetron	1A4	Perindolone	3C19
Captopril	2D6	Grisoflavin	2A6	Perphenone	2D6
Carbamazepine	1A2, 2B6, 2C8, 3C9, 3A4	Haloperidol	1A2, 2D6, 3A4	Perphenone	2D6
Carisoprodol	3C19	Hessobarbital	1A2, 3C9, 2C19	Phenobarbital	1A2, 2A6, 2B6, 2C8, 3C9, 3A4
Carisofol	2D6	Ibuprofen	2C9	Phenylbutazone	3C9
Chloramphenicol	2A6, 3A4	Ibuprofen	2A6, 2B6, 3A4	Phenylephrine	1A2, 3C9, 3C19, 3A4
Chlorazepate	1A2	Imipramine	1A2, 3C19, 2D6, 3A4	Phenylephrine	2A6
Chloroquine	2D6, 3A4	Inflamivir	1A4	Phenylephrine	3C9
Chlorpheniramine	2D6	Inflamivir	2D6	Phenylephrine	3A4
Ciclesonide	3A4	Inflamivir	2E1	Phenylephrine	3A4
Cimetidine	3C9, 3C19, 3A4	Inflamivir	1C19, 2D6, 2E1, 2E1,	Phenylephrine	3A4
Cinnarizine	2D6	Inflamivir	1A4	Phenylephrine	2B6
Ciprofloxacin	1A2, 2D6, 3A4	Inflamivir	1C8, 3A4	Phenylephrine	3A4
Cisapride	3A4	Inflamivir	1A4	Phenylephrine	3C19, 3A4
Citalopram	3C19, 2D6, 3A4	Itrazoxazole	1A4	Phenylephrine	2B6
Clarithromycin	1A2, 3A4	Ivacaftor	1A4	Phenylephrine	2D6, 3A4
Clonidine	3C19, 2D6	Ketocozazole	1A4	Phenylephrine	1A2, 2D6
Clopidogrel	2B6, 3A4	Lamoprazole	1C19, 3A4	Quetiapine	3A4
Coltrimazole	3A4, 3A4	Lercanidipine	1A4	Quinidine	2D6, 3A4
Conjunctivitis	1A2, 2D6, 3A4	Lignocaine	2B6, 3A4	Raboprazole	3C19, 3A4
Cocaine	1A6, 3A4	Lauride	1A4	Racemate	3C8
Codine	2D6, 3A4	Lopinavir	1A4	Raboprazole	2A6, 3C8, 3C9, 3C19, 3A4
Colchicine	3A4	Lorazepam	1A4	Raboprazole	2A6, 2B6, 3C8, 3C9, 3C19, 3A4
Conium	3A4	Lorazepam	1C9, 3A4	Raboprazole	2A6, 2B6, 3C8, 3C9, 3C19, 3A4
Cyclophosphamide	2A6, 2B6, 3A4	Lorazepam	1A4	Raboprazole	2D6
Dapoxetine	3A4	Meprobamate	2D6	Raboprazole	1A2, 2D6, 3A4
Debrisoquine	2D6	Mefenamic acid	1C9	Raboprazole	2B6
Debrisoquine	2D6	Meloxicam	1A4	Raboprazole	3A4
Desferrioxamine	2B6, 3A4	Methadone	2D6, 3A4	Raboprazole	3A4
Desferrioxamine	2D6, 3A4	Methocarbamol	1A2, 2A6, 2E1	Raboprazole	2A6, 3C9
Desferrioxamine	1A2	Methylprednisolone	1A4	Raboprazole	3A4
Diazepam	1A2, 2B6, 2C19, 2D6, 3A4	Metoprolol	2D6	Raboprazole	2D6, 3A4
Didanosine	3C9	Metronidazole	1A4	Raboprazole	2B6, 2E1
Digoxin	3A4	Mexiletine	1A2, 2D6	Raboprazole	3A4
Digoxin	3A4	Minoxidil	2D6	Raboprazole	3A4
Diltiazem	3A4	Misoprostol	1C9, 3A4	Raboprazole	3A4, 3A4
Divalproex	2E1	Misoprostol	2B6, 3A4	Raboprazole	3C9, 3A4
Doxazepam	3C8, 3A4	Misoprostol	1A4	Raboprazole	3A4
Efavirenz	3A4, 3A4	Moclobemide	1C19	Raboprazole	3C9, 3A4
Eralopril	3A4	Morphine	2D6	Raboprazole	3A4
Eralopril	2E1	Naproxen	1A2, 3C9	Raboprazole	1A2, 2B6, 2D6, 3A4
Ergotamine	3A4	Nefedipine	1A4	Raboprazole	3C9
Erythromycin	1A2, 3A4	Nefedipine	1C19, 3A4	Raboprazole	2D6, 3A4
Etidocaine	1A2, 3A4	Nefedipine	2B6, 3A4	Raboprazole	2B6, 3A4
Etidocaine	2B6	Nefedipine	1A4	Raboprazole	1A2, 2E1, 3A4
Ethinylestradiol	3A4	Nefedipine	1A6, 2B6	Raboprazole	2D6
Etoposide	3A4	Nefedipine	1A4	Raboprazole	3C19, 2D6
Etoposide	3A4	Nefedipine	1A4	Raboprazole	2D6
Etanercept	3A4	Nefedipine	1A4	Raboprazole	3C8, 3C9
Etanercept	3A4	Nefedipine	1A2	Raboprazole	3C9



Drug	Associated CYP	Drug	Associated CYP
Pharmaceuticals			
Toremifene	3A4	Valproic acid	2A6, 2B6, 2C9, 3A4
Tramlycypromine	2A6, 2C19	Venlafaxine	2D6
Tretinoin	2C8, 3A4	Verapamil	1A2, 3A4
Trimethoprim	2C9	Vinblastine	3A4
Trimipramine	2D6	Vincristine	3A4
Tropisetron	2D6	Vinorelbine	3A4
Unlicensed products and environmental agents			
Acetofenac	2C9	Flurythromycin	3A4
Ajmeline	2D6	Fumfylline	1A2
Alpidem	3A4	Germander	3A4
Alprenol	2D6	Glutethimide	2B6
Amiflamine	2D6	Grapefruit juice (naringenin)	3A4
Aprindine	2D6	Guanoan	2D6
Arachidonic acid	2B6	Hydrocodone	2D6
Benzphetamine	2C8, 3A4	Isoaflrole	1A2
Bufurakol	2D6	Josamycin	3A4
Caffeine	1A2, 2E1	Ketones	2E1
Cannabis (THC)	2C9, 3A4	Lauric acid	2B6
Chlorproguanil	2C19	Lobeline	2D6
Chlorzoxazone	2E1	Lornoxicam	2C9
Coumarin	2A6	Mephentoin	2B6, 2C19
Cyclobenzaprine	3A4	Metamphetamine	2D6
Delavirdine	2D6, 3A4	Methoxychlor	2B6
Deprenyl	2D6	Methoxyflurane	2B6, 2E1
Dichloralphenazone	2A6	Methoxyphenamine	2D6
Dithiocarb	2A6, 2E1	Midecamycin	3A4
Ebastine	3A4	Minaprine	2D6
Ecstasy (MDMA)	2B6, 2D6	Niludipine	3A4
Encainide	2D6	Nitrendipine	3A4
Ethanol	2C9, 2E1	Norfluoxetine	2D6
Felbamate	2C19	Oxidipine	3A4
Flunarizine	2D6		
		Phenacetin	1A2
		Phenformin	2D6
		Possinomycin	3A4
		Propylajmaline	2D6
		Quercetin	3A4
		Rauhimbine	2D6
		Remoxipride	2D6
		Rokitamycin	3A4
		Seratroclast	2B6, 2C9
		Sparteine	2D6
		St John's Wort	3A4
		Sulphamethizole	2C9
		Sulphaphenazole	2C9
		Suprofen	2C9
		Tacrine	1A2
		Teniposide	2C19, 3A4
		Terguride	3A4
		Tienilic acid	2C9, 2D6
		Tomoxetine	2D6
		Triazolam	3A4
		Trifluoperidol	2D6
		Trimethadione	1A2, 2E1, 3A4
		Troglitazone	3A4
		Troleandomycin	3A4
		Zonisamide	3A4

Readers are advised to check the extent of possible interactions shown here using additional sources eg, the British National Formulary (BNF). Drugs are shown with their associated CYPs. Those CYPs in black are known to metabolise the drug, while those in blue are also inhibited and those in red also induced by the drug. Occasionally, the drug is not a substrate of the CYP it induces/inhibits eg, quinidine is not a substrate of CYP2D6.

Non-pharmaceutical agents may not be licensed in the UK but are shown because of their practical or theoretical importance.

Abbreviations: MDMA, methylenedioxymetamphetamine; THC, tetrahydrocannabinol.

Compiled from multiple sources including 7, 8, 11, 12, 14, 16, 17 and <http://www xenotechllc.com> and <http://medicine.iupui.edu/flockhart/>



Pharmacogenomics of CYPs



Polymorphic P450s

- **Genetically Polymorphic P450s Associated with Changes in Drug Effects**
 - CYP2C9
 - CYP2C19
 - CYP2D6
 - The correlation between pheno- and genotype for CYP3A is not fully understood.



CYP Genotypes

- **Abnormal CYP alleles**
 - **Abolished enzyme activity**
 - » Deleted gene or defect in allele
 - **Reduced or altered activity**
 - » Allele is not fully functional but some functionality remains
 - CYP2C9*2 and *3
 - CYP2D6*10 and *17
 - **Increased activity**
- **CYP2D6: *genotype predicts phenotype***



CYP2D6 Potential Phenotypes

- *Poor Metabolizers*
 - lack functional enzyme.
- *Intermediate Metabolizers*
 - heterozygous for one functional and one deficient allele
 - have two partially defective alleles that cause reduced metabolism
- *Extensive Metabolizers*
 - two normal alleles
 - often majority of population
 - “normal metabolizers
- *Ultra-Rapid Metabolizers*
 - duplicated or multiduplicated functional CYP2D6 genes with extremely high metabolic capacity.



Phenotypic Effects of P450 Pharmacogenomics

Drug	Slow metabolizer phenotype	Fast metabolizer phenotype
Prodrug, needs metabolization to work (eg. codeine is metabolized by CYP 2D6 to morphine)	Poor efficacy Possible accumulation of prodrug	Good efficacy, rapid effect
Active drug, inactivated by metabolization (example is omeprazole)	Good efficacy Accumulation of active drug can produce adverse reactions May need lower dose	Poor efficacy Need greater dose or slow release formulation



Phenotyping and Genotyping

- **Phenotyping**

- Dose subjects with a compound or compounds that are metabolized to a product exclusively by the enzyme systems in question.
- Collect plasma or urine samples
 - » Single time point
 - » Over a period of time
- Analyze for model compound and metabolite
- Ratio of concentrations of compound and its metabolite is used to measure metabolic capacity for a specific P450.

- **Genotyping**

- Collect blood (> 1 ml)
- Isolate DNA from nucleated blood cells.
- Amplify number of copies of DNA by the Polymerase Chain Reaction (PCR).
- Genotype by sequencing or probing.



Pharmacogenomics in Drug Development

- **DNA samples taken for ADME genotyping in drug development**
 - Routine if one enzyme is known as the predominant route of metabolism.
- **Compounds with narrow safety margin**
 - Reduce risk of developing concentration-dependent side effects when treated with standard doses
 - Exclude poor metabolizers (if the parent drug is predominantly biologically active)
 - Exclude ultra-rapid metabolizers (if metabolite is predominantly biologically active)
- **Compounds with wide therapeutic window**
 - Dose adjustments based on pharmacogenomic tests.
 - Increase opportunity for regulatory approval on subpopulation.
 - Less important if compound and metabolite have similar activity.
- **Troubleshooting**
 - Retrospective analysis in subjects with side effects or lack of therapeutic effect.
 - Prediction of ethnic variation explaining profiles in different populations.



Clinical Examples



Labeling Regulations

“If evidence is available to support the safety and effectiveness of the drug only in selected subgroups of the larger population with a disease, **the labeling shall describe the evidence and identify specific tests needed for selection and monitoring of patients who need the drug.**”

- 21 CFR 201.57



How Does It Read?: Examples of Pharmacogenomic Information in the Drug Label

Brand Name (generic name)	Labeling section	Labeling Statement
HERCEPTIN® (trastuzumab) August 2002	INDICATIONS AND USAGE	HERCEPTIN should be used in patients whose tumors have been evaluated with an assay validated to predict HER2 protein overexpression (see <u>PRECAUTIONS</u> : <u>HER2 Testing</u> and <u>CLINICAL STUDIES</u> : <u>HER2 Detection</u>).
Purinethol (6-Mercapto- purine) July 2004	WARNINGS DOSAGE and ADMINISTRATI ONS	Individuals who are homozygous for an inherited defect in the TPMT (thiopurint-S-methyltransferase) gene may be unusually sensitive to the myelosuppressive effects of mercaptopurine and prone to developing rapid bone marrow suppression following the initiation of treatment. (see DOSAGE AND ADMINISTRATION). Patients with inherited little or no thiopurine S-methyltransferase (TPMT) activity are at increased risk for severe PURINETHOL toxicity from conventional doses of mercaptopurine and generally require substantial dose reduction. The optimal starting dose for homozygous deficient patients has not been established (see <u>CLINICAL PHARMACOLOGY</u> , <u>WARNINGS</u> and <u>PRECAUTIONS</u> sections)
(thioridazine) July 2003	CONTRA- INDICATIONS	thioridazine is contraindicated in patients, comprising about 7% of the normal population, who are known to have a genetic defect leading to reduced levels of activity of P450 2D6 (see <u>WARNINGS</u> and <u>PRECAUTIONS</u>).
STRATTERA (atomoxetine) March 2003	Drug-Drug Interactions Laboratory Tests	In EMs, inhibitors of CYP2D6 increase atomoxetine steady-state plasma concentrations to exposures similar to those observed in PMs. Dosage adjustment of STRATTERA in EMs may be necessary when coadministered with CYP2D6 inhibitors, e.g., paroxetine, fluoxetine, and quinidine (see <u>Drug Interactions under PRECAUTIONS</u>). In vitro studies suggest that coadministration of cytochrome P450 inhibitors to PMs will not increase the plasma concentrations of atomoxetine. <u>CYP2D6 metabolism</u> --Poor metabolizers (PMs) of CYP2D6 have a 10-fold higher AUC and a 5-fold higher peak concentration to a given dose of STRATTERA compared with extensive metabolizers (EMs). Approximately 7% of a Caucasian population are PMs. Laboratory tests are available to identify CYP2D6 PMs. The blood levels in PMs are similar to those attained by taking strong inhibitors of CYP2D6. The higher blood levels in PMs lead to a higher rate of some adverse effects of STRATTERA (see <u>ADVERSE REACTIONS</u>).



Mercaptopurines

- **Leukemia indication**
 - **Converted to nucleotides for incorporation into DNA by hypoxanthine phosphoribosyl transferase (HPRT).**
 - **Mercaptopurine-derived products block DNA replication and lead to tumor cell death.**
- **Mercaptopurine metabolism**
 - **Thiopurine methyltransferase (TPMT) converts mercaptopurine into an inactive metabolite called methylmercaptopurine.**
 - » **90% homozygous for wild type allele and metabolize product normally**
 - **Toxicity is low, but relapse is high.**
 - » **Some are poor metabolizers**
 - **Toxicity is high.**
 - » **0.3% are homozygous for these variants**
 - » **High risk of myelosuppression and secondary tumors.**
- **Label: “Recommendation to use pharmacogenetic testing to guide dosing”.**

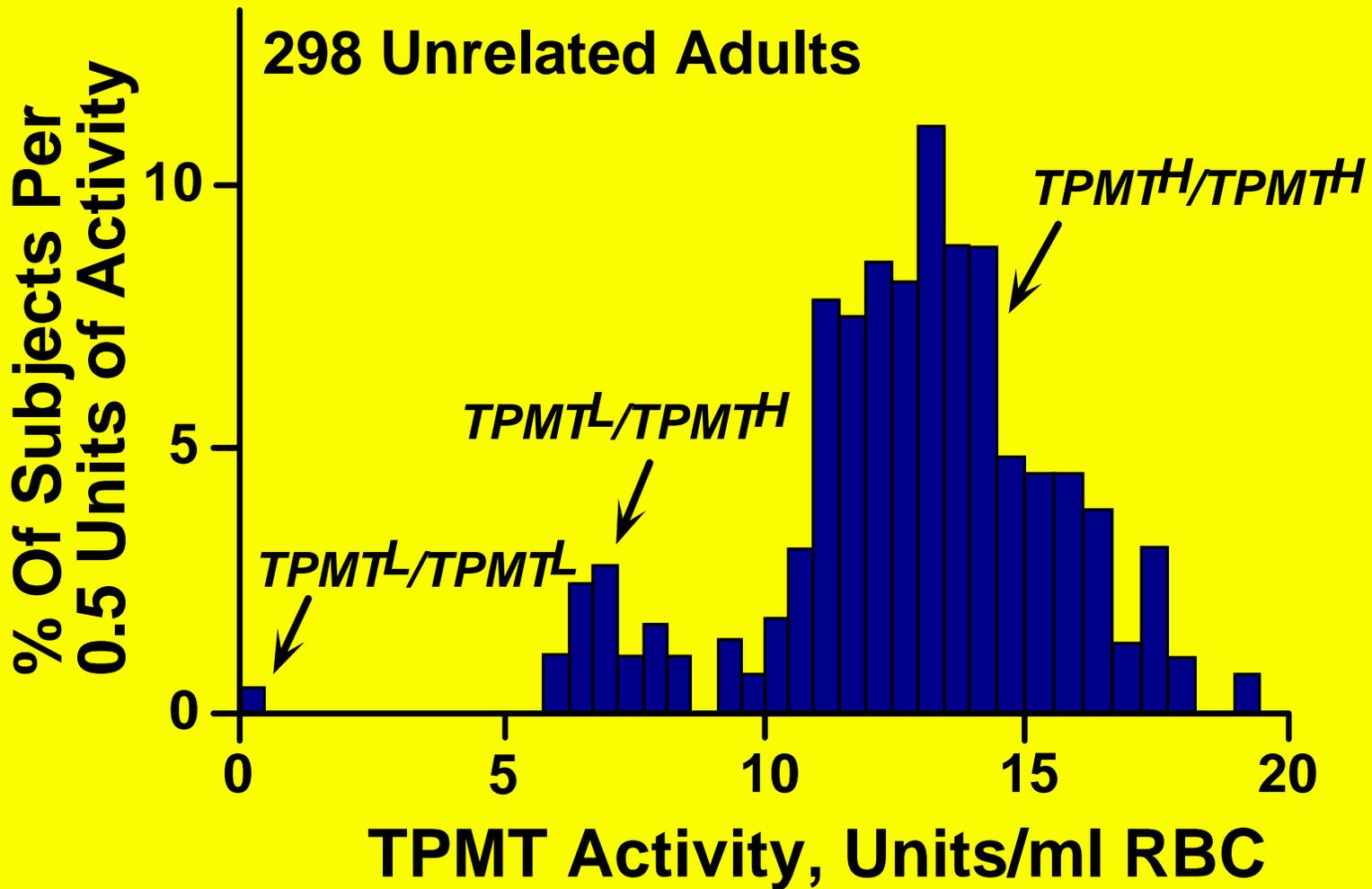


2005 FDA Pharmacogenomic Guidance *Valid Biomarkers*

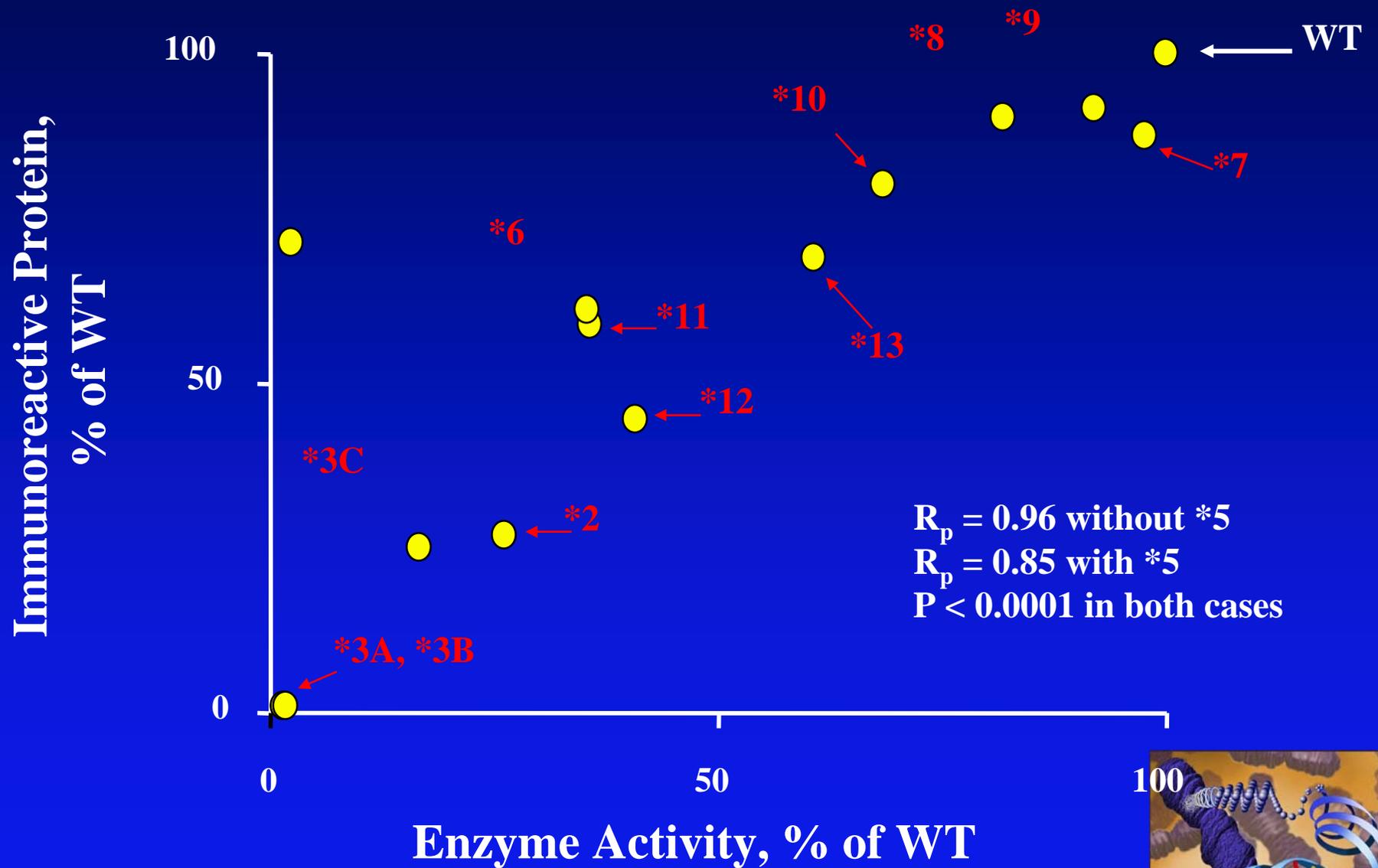
- Cytochrome P450 2D6 (CYP2D6)
- Thiopurine S-methyltransferase (TPMT)



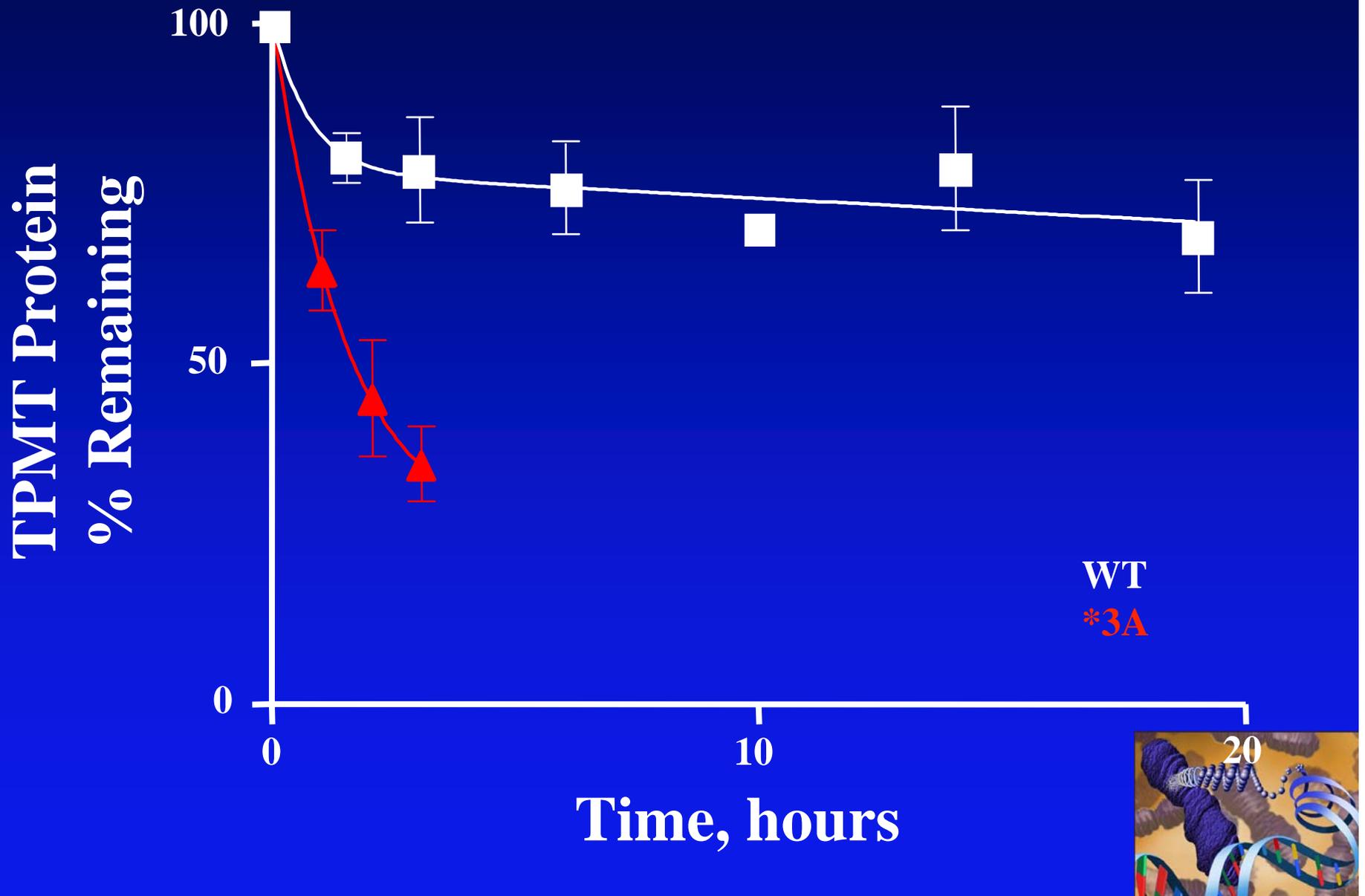
Human RBC TPMT



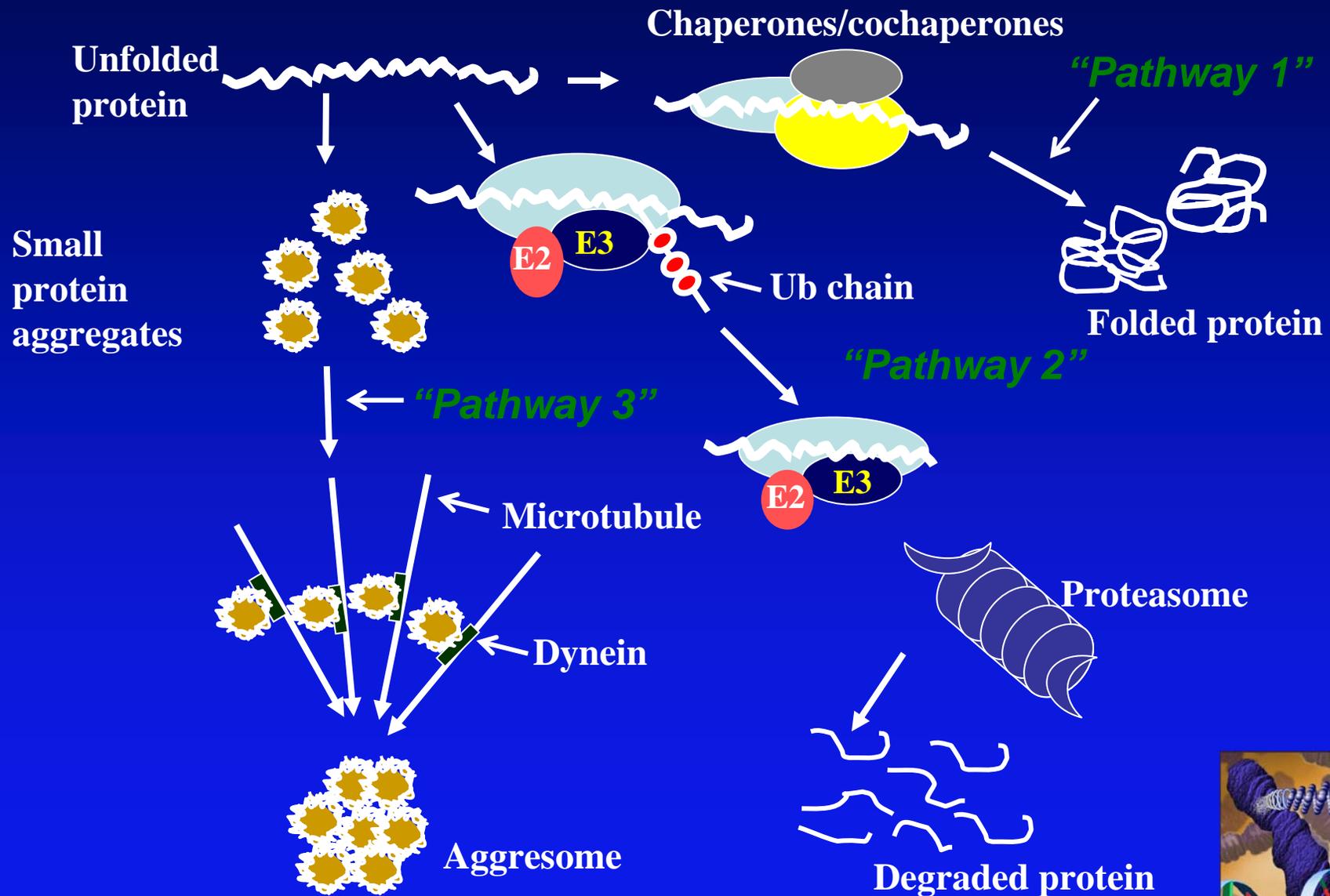
Human Recombinant TPMT Allozymes



Human TPMT Protein Degradation



Protein Folding, Degradation and Aggregation



TPMT Genetic Polymorphism

Clinical Consequences

- **Low TPMT**
 - Increased thiopurine toxicity
 - Increased risk for secondary neoplasm
- **High TPMT**
 - Decreased therapeutic effect



6-Mercaptopurine and TPMT Polymorphism

- *Labeling* – absence of PGx information in label discussed at CPSC and Pediatric Oncology Subcommittee in 4/03 and 7/03
- *New labeling* - revised by sponsors in consultation with FDA- includes data on increased risk of severe myelosuppression for TPMT activity-deficient genotypes
- Informs clinicians about option of using TPMT testing to guide treatment with 6MP



General Process for Updating Labels with PGx Information

- **Develop the appropriate questions**
- **Capture the relevant evidence**
- **Abstract and summarize the evidence**
- **Evaluate the quality of studies**
- **Assess the overall strength of evidence**
- **Consider other factors in relabeling decision**
- **Determine specific language for label**



Warfarin

- **Optimal use hampered by 10-fold interpatient variability in doses required for therapeutic response.**
- **CYP2C9 allelic variants are associated with impaired elimination of warfarin and exaggerated anticoagulatory responses to the drug.**
 - CYP2C9*1 is the wild type
 - Other variants include CYP2C9*2, CYP2C9*3, CYP2C9*4 and CYP2C9*5
 - » Allelic frequencies depend on ethnicity
- **185 patients genotyped for CYP2C9 and treated with warfarin**
 - Variant groups required more time to achieve stable dosing with a median difference of 95 days.
 - Gene-dose effect relationship suggested when comparing the *1/*1, *1/*2, and *1/*3 genotypes, with corresponding mean maintenance doses of 5.63, 4.88, and 3.32 mg.
 - *Conclusion:* CYP2C9*2 and CYP2C9*3 polymorphisms are associated with increased risk of over coagulation and bleeding events.
 - In general, subjects who are heterozygous for these two alleles require a 60 to 75% lower dose of warfarin compared to homozygous wild-type patients.



What would you do?

