

FDA/Johns Hopkins University/PhRMA Educational Workshop
Drug Metabolizing Enzymes and Pharmacogenomic Testing-
September 13-14, 2003
Rockville, MD

**Regulatory Issues in Genotyping
Metabolizing Enzymes-
CDER Perspective**

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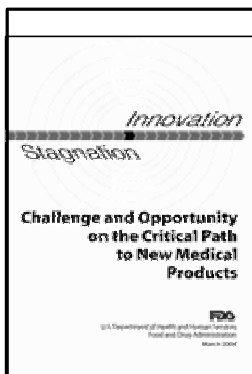
**Traditional Inefficiency in Drug
Development**

- Extremely high pre-IND failure rate
- Less than 1 in 5 IND's for NME's = NDA's
- Estimated cost per NME about \$800 million
- Time from IND to market around 8-10 years
- Multiple review cycles for most NME NDA's

Is this a systemic problem?
Is this the best that can be done?
What is needed?

Lesko, L. Pharmaceutical Sciences World Conference, Japan, May 18, 2004<

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The product development
problems we are seeing
today can be addressed,
..... generation of
predictive tools.

The new tools such as
bioinformatics, *genomics*,
imaging technologies, and
materials science.

<<http://www.fda.gov/oc/initiatives/criticalpath/whitepaper.html>>

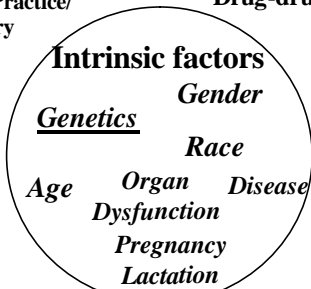
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Clinical Pharmacology and Biopharmaceutics Review

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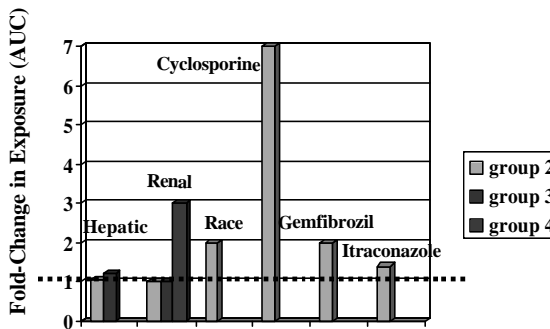
Extrinsic factors

Environmental Smoking/Diet
 Medical Practice/
 Regulatory Drug-drug interaction



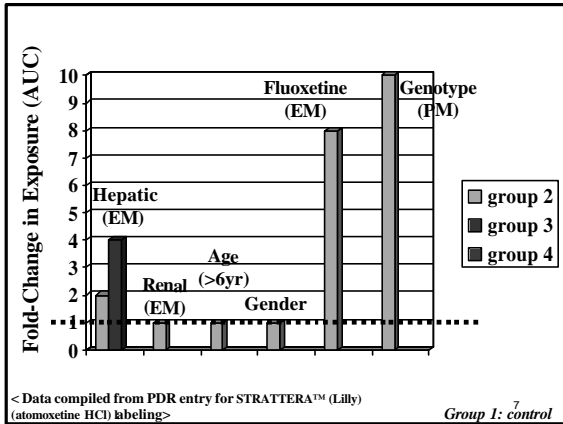
Adapted from ICH E5, 1998: <http://www.fda.gov/cder/guidance/2293fnl.pdf>

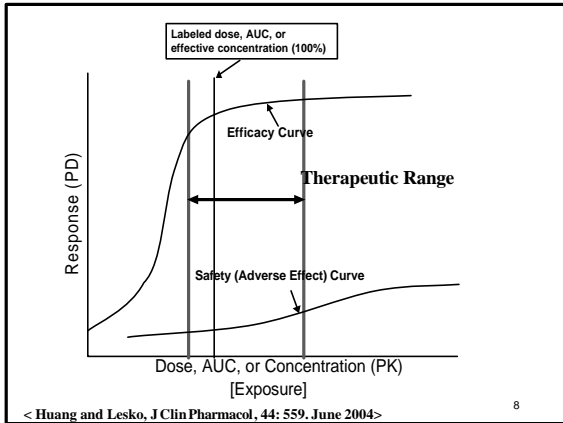
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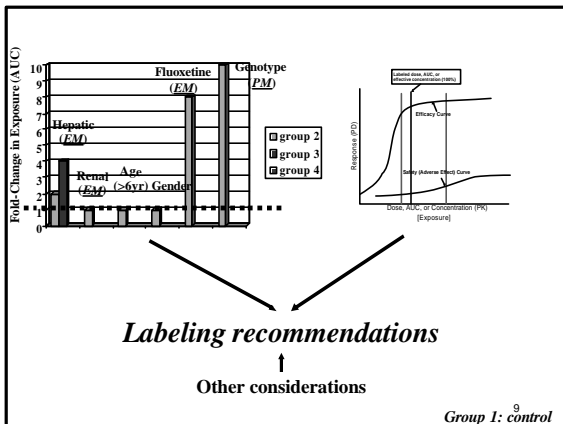


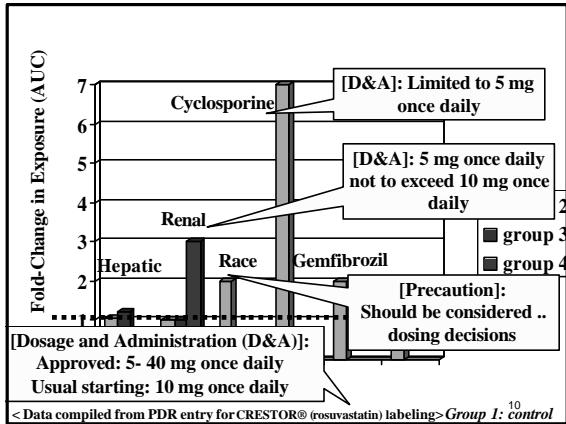
< Data compiled from PDR entry for CRESTOR® (AstraZeneca) (rosuvastatin calcium) labeling >

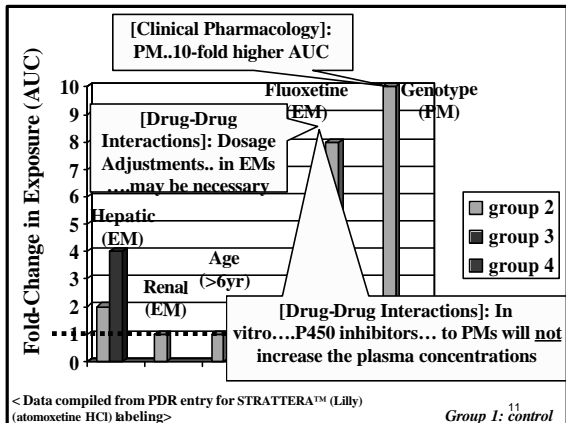
Group 1: control







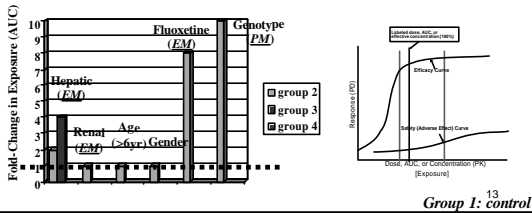




Atomoxetine Labeling

- **Laboratory Tests** : Laboratory tests are available to identify CYP2D6 PMshigher blood levels in PMs lead to higher rate of some adverse effects of STRATTERA.

What Pharmacogenomic Testing data are needed when generating these initial PK data (or clinical data) that eventually support the labeling (or future labeling) for drugs metabolized by polymorphically distributed enzymes?



Issues

- Are designations of EM and PM inadequate?
- do we need UM and IM?

CYP2D6

EM with 1 wild type *1 *1*1, *1*2,...*1*5, *1*10, *1*17...

PM with two null alleles *3*3, *3*5, *4*6,...
*3-8, *11-16,

IM? with two reduced alleles *10*10, *17*17, *4*17,...
or one reduced, one null

UM? with *2xN *1*2xN

Issues (2)

- Are designations of EM and PM inadequate?
- do we need UM and IM?
- What data to support genotype-phenotype correlation?
- How could these designations be improved?
- What are the definitions of EM and PM for all enzymes?
- “PM” applicable to all enzymes? CYP2C9?

Issues (3)

- What alleles need to be assessed before declaring a specific genotypes?
 - Can we extrapolate data from one ethnic/race group to the other?

CYP2C9

	Caucasian	Asian	African American	Hispanic
*2	8-19	0	3.2	12
*3	3.3-16	1.1-3.3	1.3	3.4
*5	--	--	0.2-1.7	0-0.5
*6	--	--	0.6-1.5	----

<Data from Xie HG, et al. Advanced Drug Delivery Reviews 54 (2002) and references therein>¹⁶

Issues (4)

- What information (PM, EM/others and specific alleles) is to be included in the drug label?
- To what extent should the test be commercially available?
- Why should the test be approved by the FDA to be in the labeling?

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Guidance for Industry Pharmacogenomic Data Submissions

DRAFT GUIDANCE

This guidance document is being disseminated for comment purposes only. Comments and suggestions regarding this draft document should be submitted within 90 days of publication to the contact person(s) in the table accompanying the availability of the draft document. Submit comments to the Division of Dockets Management (HFA-305), Food and Drug Administration, 1015 Lincoln Avenue, Silver Spring, MD 20910. All comments should be identified with the document number listed in the table of availability that publishes in the Federal Register.

For questions regarding this draft document contact CDER, Liaison(s) Link(s): 301-794-5070, CDER, Regulatory Affairs, CDER, or CDER's Office of Communications.

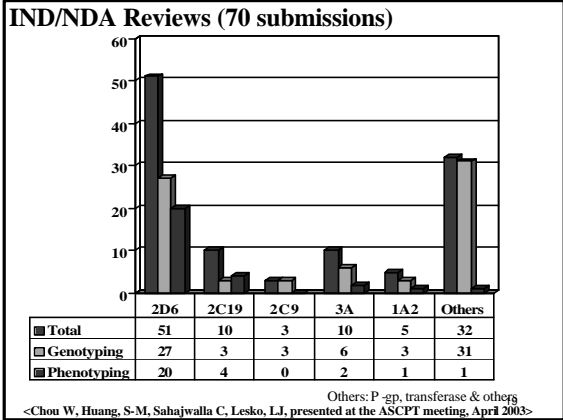
U.S. Department of Health and Human Services
Food and Drug Administration
Center for Biologics Evaluation and Research, CDER
Center for Drug Evaluation and Research, CDER
Center for Safety and Health Data Research, CDER

November 2003
Procedural

Known valid biomarker

**.....CYP2D6 and
TPMT (thiopurine
methyltransferase)**

<Guidance for Industry: Pharmacogenomic Data Submission;>
<[http://www.fda.gov/cder/guidance/index.htm_procedural\(draft\),November2003;](http://www.fda.gov/cder/guidance/index.htm_procedural(draft),November2003;)>¹⁸



Case 1
Drug A (NDA)

- *metabolized by CYP3A/CYP2D6*
- *PM 2-3 fold higher AUC than EM (high variability)*
- *PM/EM determination*
 - *earlier studies: *3, *4*
 - *later studies: *3, *4, *5, *6, *10, *16, (*17)*
- *possible reduced CYP2D6 contribution at higher doses*

Case 1 (2)
Drug A

- *Based on PK and clinical data: the sponsor suggested no special dosing requirement for PM*
- *How critical is the accuracy in the PM and EM designation in evaluating these data?*
 - *Alleles analyzed are limited and differ among studies*
 - *Data analysis grouped based on EM and PM only*
 - *Labeling mentions and makes recommendations based only EM and PM*

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**Case 2
Drug B (IND)**

- *Metabolized by CYP2C9*
- **the sponsor is conducting a drug interaction study with a CYP2C9 inhibitor, fluconazole**
- **Planning on retrospective CYP2C9 genotyping to identify impact on exposure of Drug B with fluconazole**
- *What alleles should be assessed in this study? Would the analysis of *2,*3,*5 (reduced activity), or *6 (null; but rare) appropriate?*

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**Case 3
Drug C (IND)**

- *Inhibits CYP2C19 and CYP3A*
- **the sponsor is conducting a drug interaction study with a CYP2C19/CYP3A substrate, nelfinavir**
- **Planning on retrospective CYP2C19 genotyping to identify impact on exposure of nelfinavir with drug C**
- *What alleles should be assessed in this study? Would the analysis of *2 and *3 appropriate?*

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**Today's discussion on
metabolic biomarkers**

- **Known valid biomarkers**
 - CYP2D6, CYP2C9, CYP2C19
- **Probable valid biomarkers**
 - UGT1A1
- **Exploratory biomarkers**
 - ABCB1; CYP3A4/5
 - Others

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**Special Considerations for Individual
Metabolic Biomarkers
Chair: S.-M. Huang, FDA**

CYP2D6	D. Flockhart	Indiana Univ.
CYP2C9	P. Milos	Pfizer
CYP2C19	T. Andersson	AstraZeneca
UGT1A1	Mark Ratain	Univ. Chicago
CYP3A4/5	K. Thummel	Univ. Washington
P-gp and other transporters	D. Kroetz	UCSF
Haplotype Mapping of ADME Genes	D. Goldstein	UC, London

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Discussion

- **In vitro - in vivo correlation**
- **Genotype- phenotype correlation**
- **What alleles to measure?**
- **Haplotypes?**

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Acknowledgement

OCPB Pharmacogenomics Working Group and
OCPB reviewers (case examples)

A Bhattaram F Frueh S-M Huang
M-J Kim I Lee L Lesko
A Rahman K Roy A Rudman
H Sun J Wei S Yasuda
L Zhang S Nallani J Collins
P Hinderling

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