



Bayer Technology Services

Quantifying polymorphic metabolic clearance of diclofenac in CYP2C9 genotyped individuals using coupled physiology-based pharmacokinetic models

Population Approach Group of Europe July 14-16, 2006

A. Edginton¹, M. Sevestre¹, J. Kirchheiner², J. Brockmöller³, S. Willmann¹

¹ Systems Biology & Computational Solutions, Bayer Technology Services GmbH, Leverkusen, Germany

² Pharmacology of Natural Products & Clinical Pharmacology, University of Ulm, Ulm, Germany

³ Clinical Pharmacology, University of Göttingen, Göttingen, Germany

Objective

- Determine the correlation between CYP2C9 genotype and intrinsic CYP2C9 metabolism for different individuals
- Determine the importance of CYP2C9-mediated clearance to total diclofenac clearance
- Understand the advantages and limitations of using the coupled PBPK model approach



Diclofenac – What is known?

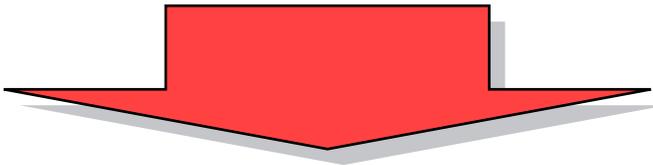
- Exclusively cleared in the liver
- Metabolism to 4'OH-diclofenac mediated by CYP2C9
- % cleared via CYP2C9 is unknown. The most recent in vitro study estimated about 25%.
- CYP2C9 is polymorphic with three alleles (*1, *2, *3)
- In vitro CYP2C9 metabolism studies inconclusive on the correlation between genotype and 4'OH-diclofenac production
- In vivo oral clearance shows no CYP2C9 genotype dependence



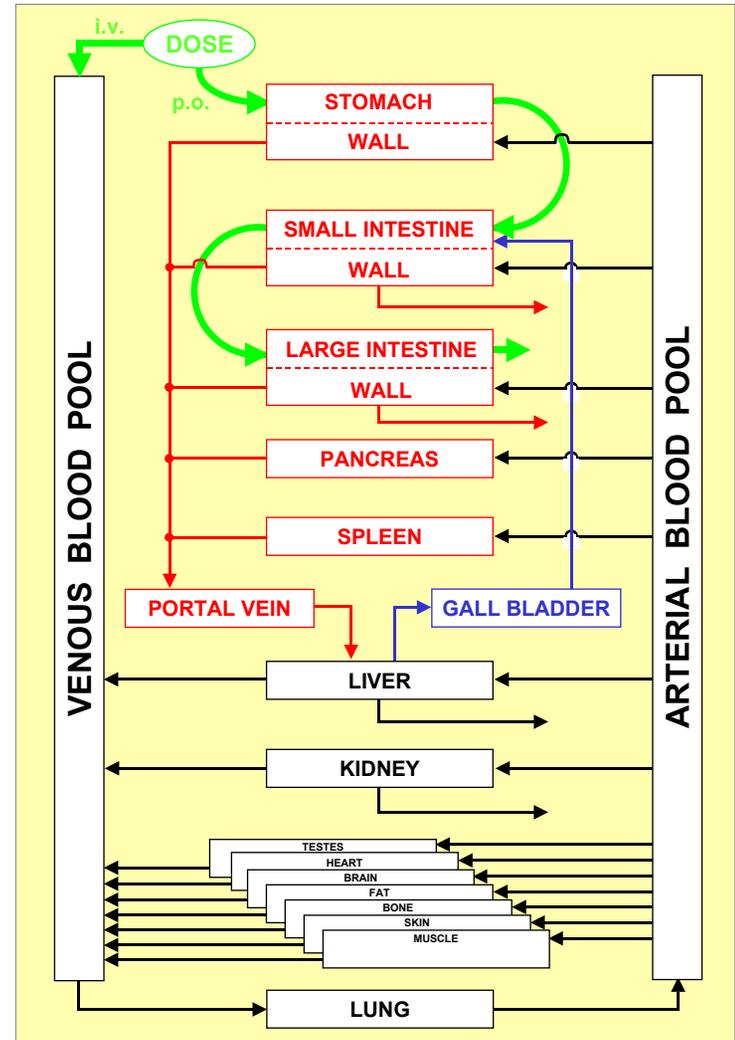
PK-Sim[®] Model-Structure

Integrated Whole Body Model:

- ⇒ Mathematical description of most important organs and their relationship to each other and to the blood pools
- ⇒ Organs separated into vascular (plasma + rbc), interstitial and cellular space
- ⇒ Active processes (metabolism, transporters) can be included in every organ



Capability for treating even very sophisticated problems.



PK-Sim® Model-Inputs

- Height, weight, gender – generates organ weights, blood flows (see PK-Pop poster Friday afternoon)
 - Physico-chemistry of parent and metabolite
 - Lipophilicity
 - Molecular weight
 - Solubility
 - Acid/Base Properties
 - Unbound fraction in plasma
- } Tissue: plasma partition ratios

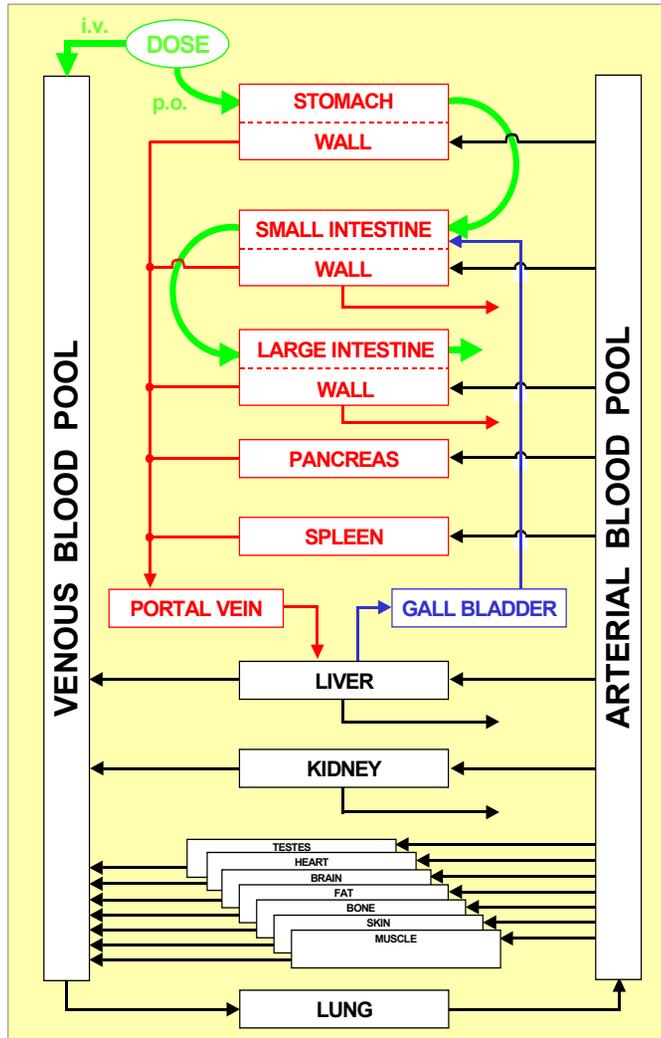
Clinical Data

- 19 CYP2C9 genotyped adult males
 - *1*1; $n = 2$
 - *1*2; $n = 4$
 - *1*3; $n = 4$
 - *2*2; $n = 3$
 - *2*3; $n = 3$
 - *3*3; $n = 3$
- Diclofenac and 4'OH-diclofenac plasma concentration time profiles over 10 hours following oral administration
- Anthropometric information of age, weight and height

Kirchheiner et al. *Br J Clin Pharmacol* 55:51-61 (2003).

Coupling Drug & Metabolite PBPK Models

Diclofenac PK-Sim Simulation

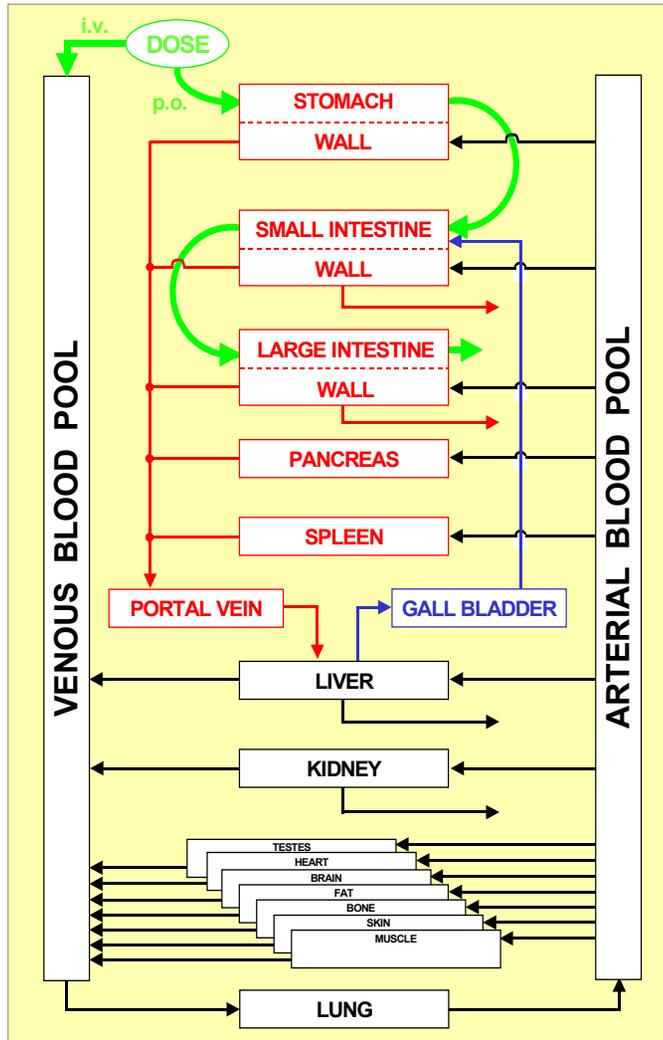


Check distribution using IV data

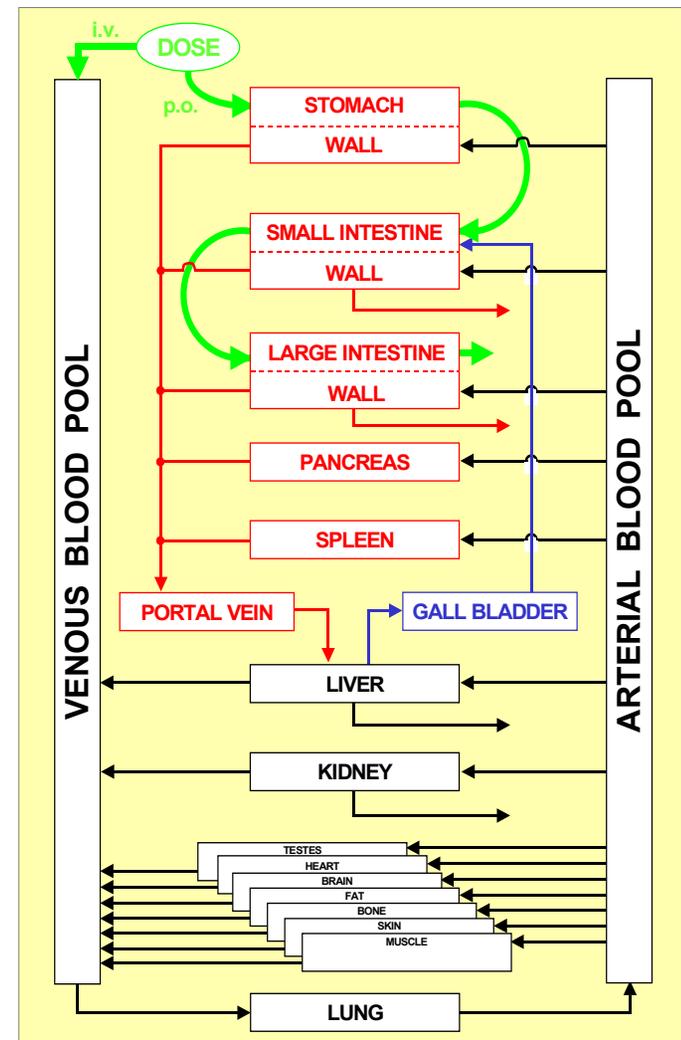


Coupling Drug & Metabolite PBPK Models

Diclofenac PK-Sim Simulation



4'OH-Diclofenac PK-Sim Simulation

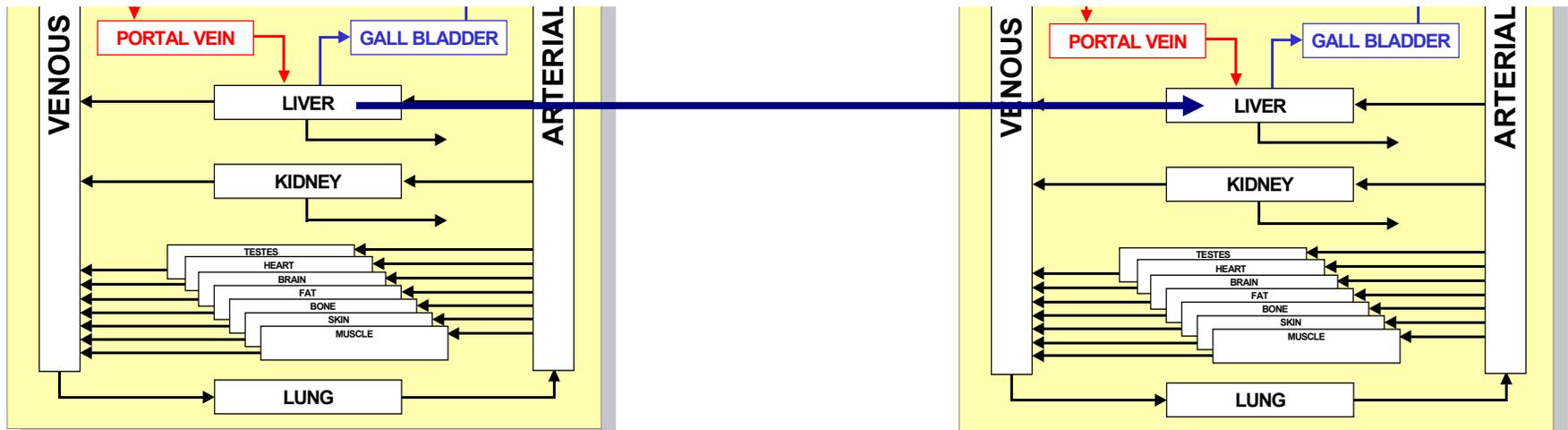


Coupling Drug & Metabolite PBPK Models

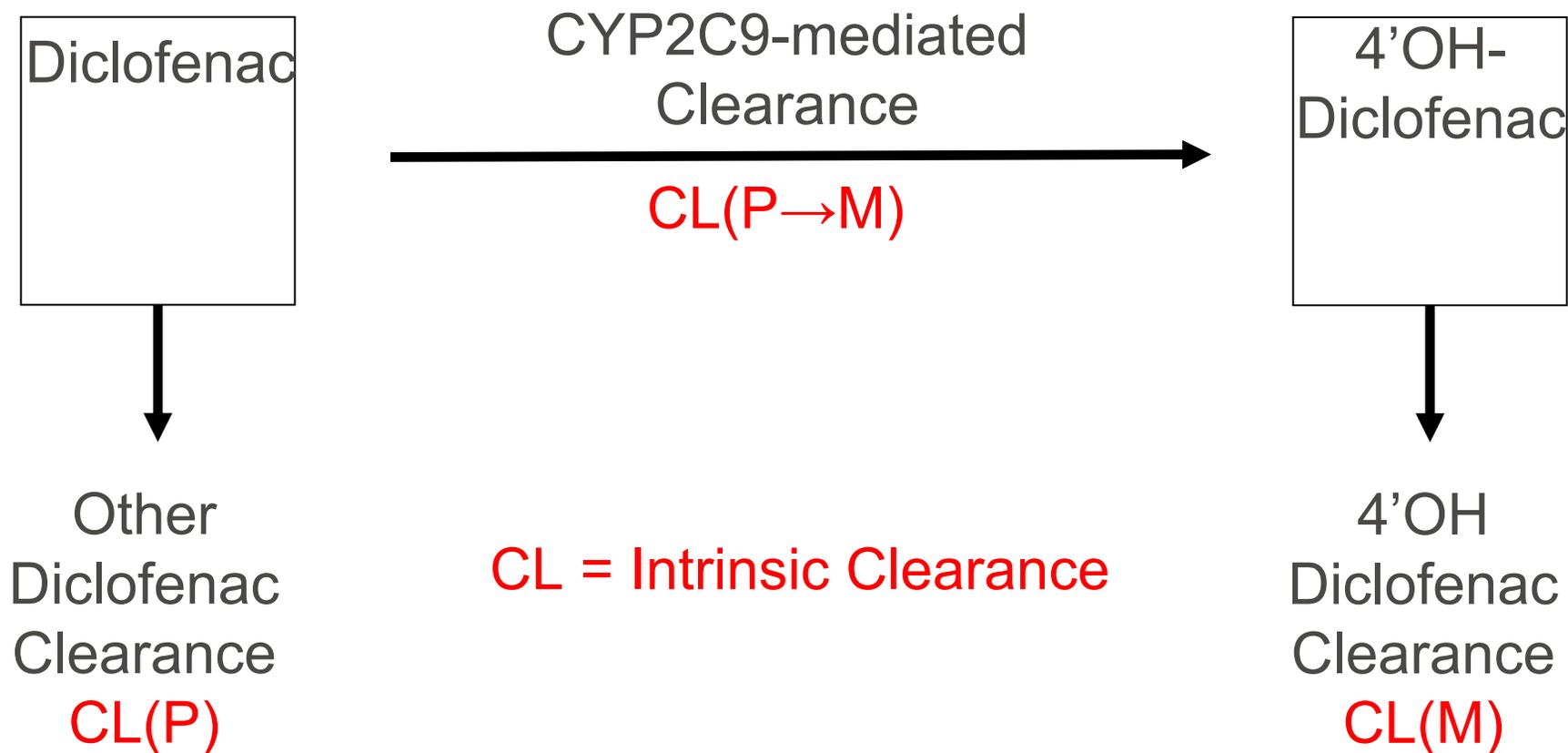
Diclofenac PK-Sim
Simulation

4'OH-Diclofenac PK-Sim
Simulation

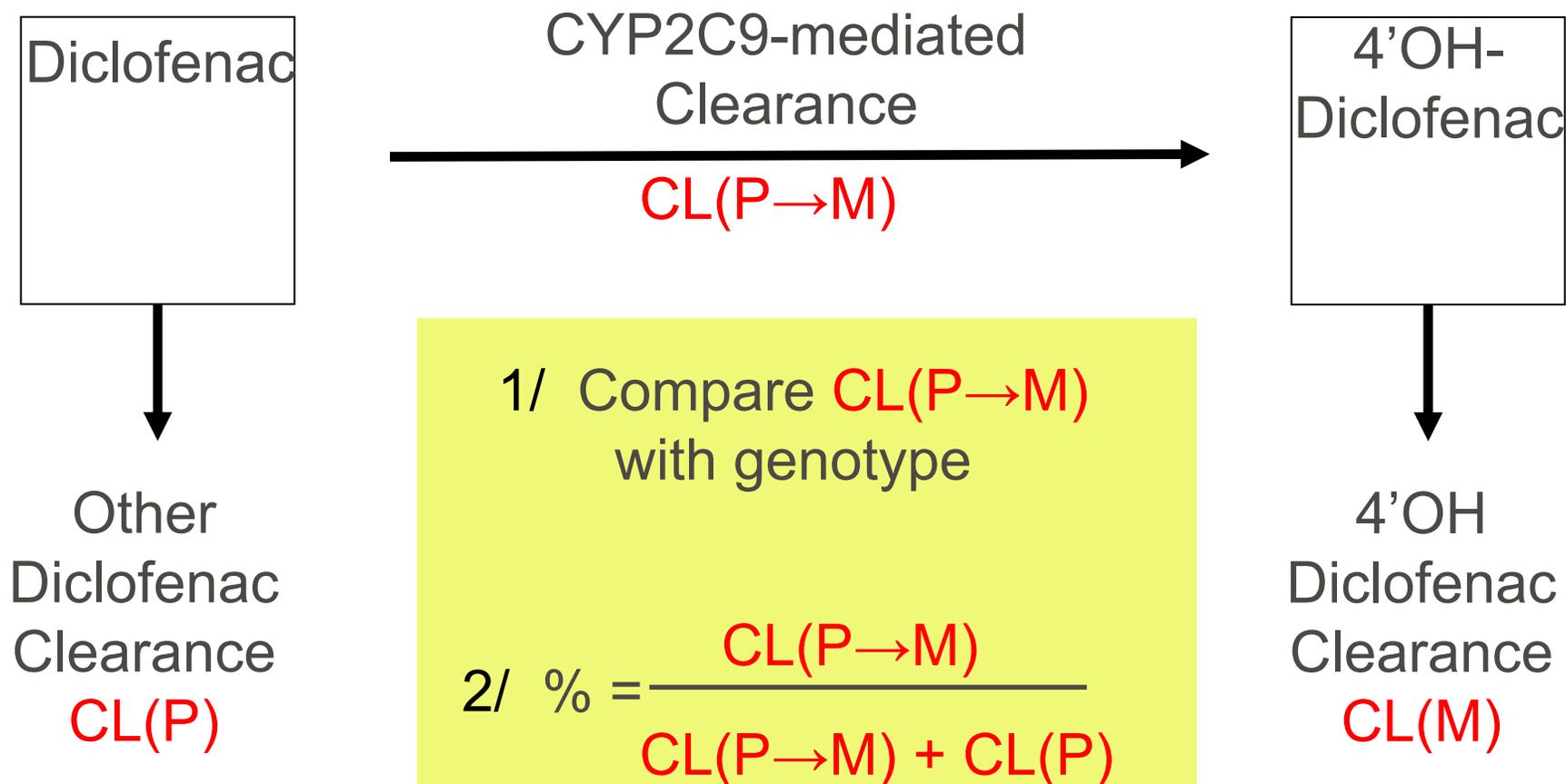
**Source of 4'OH diclofenac is diclofenac
CYP2C9-mediated metabolism in the
liver intracellular space**



Clearance Optimization

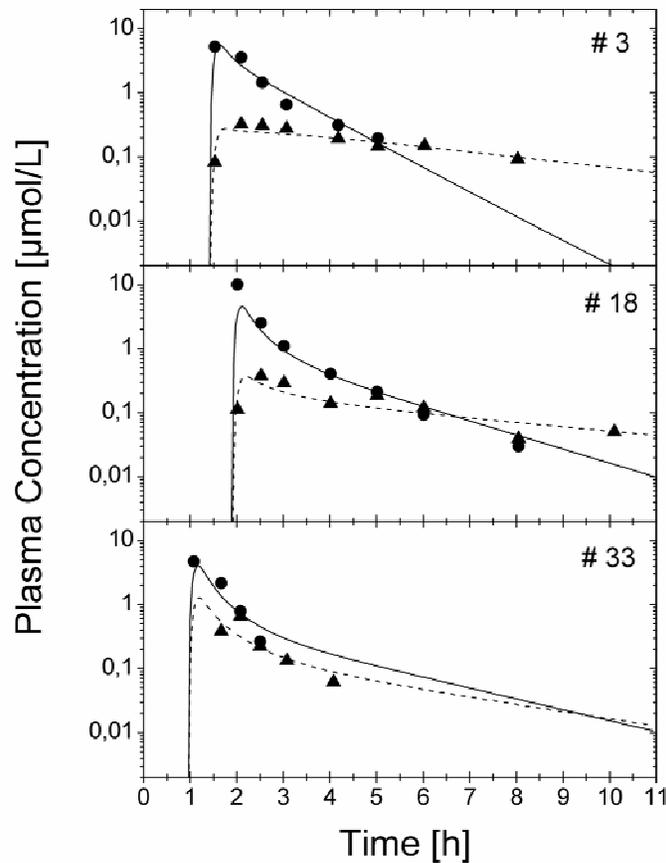


Clearance Optimization

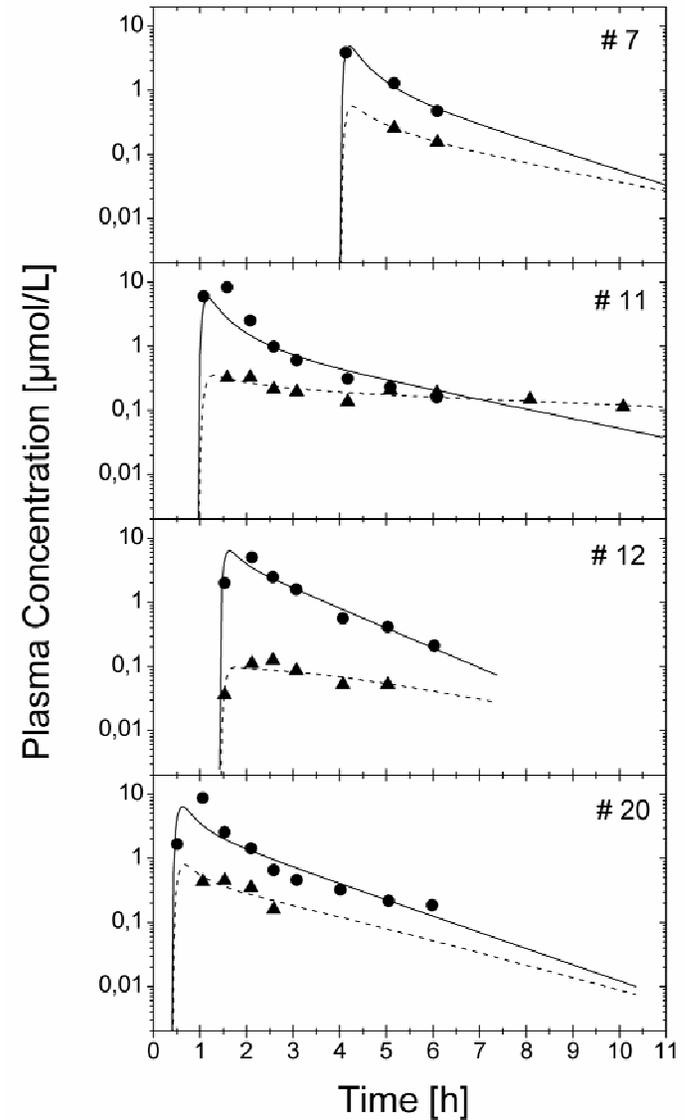


Diclofenac Results

CYP2C9 *2*2

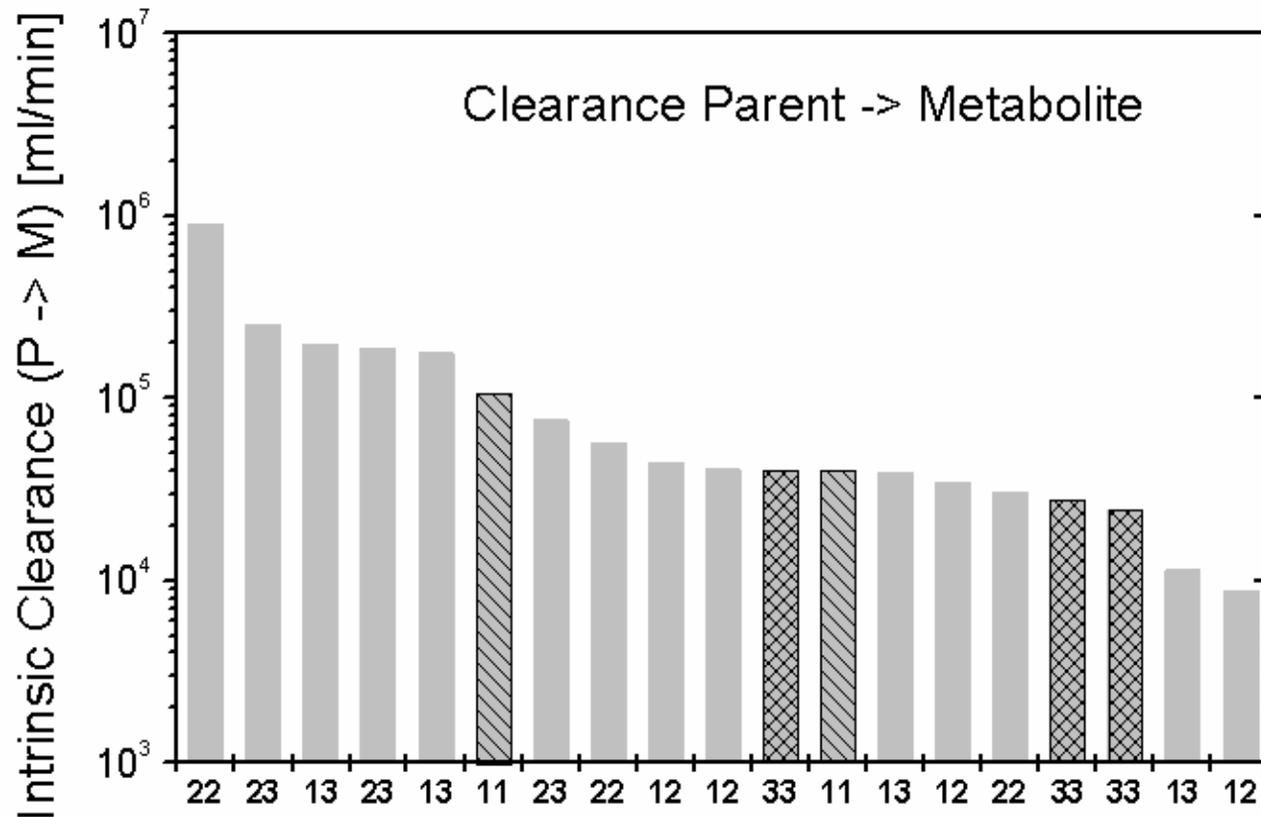


CYP2C9 *1*3



Diclofenac Results

Weak genotype-phenotype correlation!



% of Total Clearance Due to CYP2C9

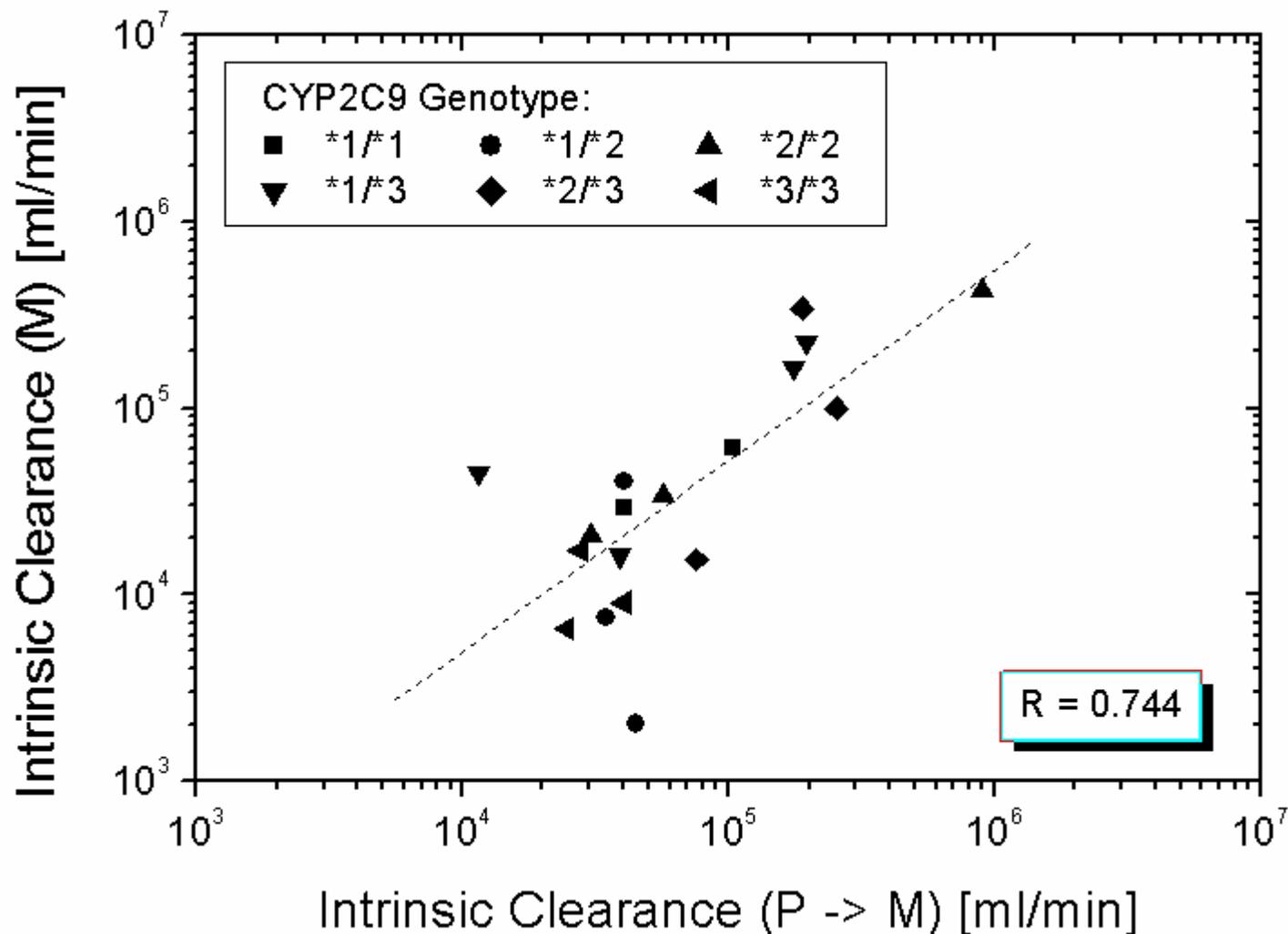
- Based on optimized clearance values, $CL(P \rightarrow M)$ as a percentage of total diclofenac intrinsic clearance was:
 - **Median 7.2%**
 - **Mean 19.8%**
 - **Range 1.9 to 93.8%**

% of Total Clearance Due to CYP2C9

- Based on optimized clearance values, $CL(P \rightarrow M)$ as a percentage of total diclofenac intrinsic clearance was:
 - **Median 7.2%**
 - **Mean 19.8%**
 - **Range 1.9 to 93.8%**

A weak pheno/genotype correlation along with a limited importance of CYP2C9 likely the reason why no in vivo correlation between oral diclofenac clearance and genotype is observed

Correlation between $CL(P \rightarrow M)$ and $CL(M)$



Potential Limitations

- Distribution is adequately estimated from physico-chemistry of the parent (can test) and metabolite (assumed)
- Requires a metabolite curve with observed data surrounding both the apparent C_{max} and terminal phase

Defines $CL(M)$

Primarily defines $CL(P \rightarrow M)$

Uses of Method

- Determine the relative importance of the metabolite to the overall clearance of the parent
- Determine if there is an *in vivo* genotypic effect for polymorphic enzymes
- Generate a measure of the inter-individual variation in intrinsic clearances
- Examine age-related differences in metabolite production



Thank you

Acknowledgement

Thorsten Hartman from Nimbus Technology for the lipophilicity (Log MA) and serum binding measurements.

