Co-operativity in the *in vitro* kinetics of cytochrome P450 (CYP) mediated drug metabolism will have minimal impact on *in vivo* metabolic clearance M. Jamei¹, A. Rostami-Hodjegan^{1,2}, G. T. Tucker^{1,2} <u>M.Jamei@Simcyp.com</u>

1- Simcyp Ltd, Blades Enterprise Centre, John St, Sheffield, S2 4SU, UK 2- Academic Unit of Clinical Pharmacology, University of Sheffield, Sheffield, UK

Introduction

Some CYPs exhibit atypical kinetics under *in vitro* conditions, and a two-site binding model has been proposed to accommodate negative and positive hetero- or homo-tropic co-operativity. The perfused rabbit liver with constant rate input of drug has been used to demonstrate possible *in vivo* consequences of this behaviour (Chen *et al.*, 2004), while there is no unequivocal evidence from *in vivo* studies in humans to support the importance of atypical metabolic kinetics.

In this study we have assessed the potential impact of homotropic co-operativity (substrate inhibition or activation), using the two-site binding model nested in a mechanistic physiologically-based pharmacokinetic (PBPK) model implemented in Simcyp[®].

Methods

Figure 1 shows a schematic of the two-site binding model, where [E], [S] and [P] are enzyme, substrate and product concentrations, [SE], [SES] and [ES] are enzyme-substrate complex concentrations, α and β are the co-operativity parameters and K_s and K_p are dissociation and metabolic rate constants, respectively.

$$[E] + [P] \xleftarrow{K_{p}} [SE] \xleftarrow{\alpha K_{s}} [SES] \xrightarrow{\beta K_{p}} [ES] + [P] \text{ or}$$

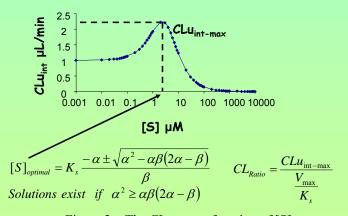
$$[E] \xleftarrow{K_{s}} f \xrightarrow{\alpha K_{s}} [SES] \xrightarrow{\alpha K_{s}} [ES] + [P]$$

Figure 1 – A schematic of the two-site binding model.

The governing equations for this model is:

$$v = \frac{\frac{\left[S\right]}{K_{s}} + \frac{\beta\left[S\right]^{2}}{\alpha K_{s}^{2}}}{1 + \frac{2\left[S\right]}{K_{s}} + \frac{\left[S\right]^{2}}{\alpha K_{s}^{2}}} V_{\text{max}} \qquad CLu_{\text{int}} = \frac{v}{\left[S\right]}$$

where v is the rate of metabolism, CLu_{int} is unbound intrinsic clearance and V_{max} is the maximum rate of metabolism. From the first-order differential the concentration at which maximum intrinsic clearance occurs can be determined. Figure 2 shows how CLu_{int} changes as a function of substrate concentration.





Simcyp[®] version 6.1 was used to generate a virtual North European Caucasian population and their individual plasma drug concentration-time and hepatic and intestinal intrinsic clearance – time profiles.

In order to compare the effects of autoactivation on AUC and C_{max} , intrinsic clearances for all metabolic routes were calculated using Michaelis-Menten (MM) and two-site binding models.

Results

The effects on *in vivo* kinetics of a range of co-operativity numbers (α and β) reported from *in vitro* studies of different drugs were investigated. As a worse case scenario, the most extreme values of α and β (0.03-23.8) that have been reported (Lin *et al.*, 2001) were used in simulation studies to examine the *in vivo* implications of autoactivation.

The time-dependences of the hepatic CL_{int} of alprazolam and tolbutamide based on MM kinetics and the two-site binding model are compared in Figures 3 and 4.

ose = 0.5 mg
$$i\nu$$
, α = 0.03, β = 25, $CL_{int-atypical} / CL_{int-MM}$ = 61.94
AUC.... = 0.0923 (mg/L h) AUC... = 0.0919 (mg/L h)

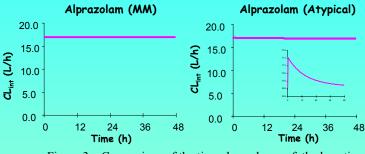
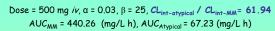


Figure 3 – Comparison of the time-dependence of the hepatic CL_{int} of alprazolam predicted by Michaelis-Menten and Two-Site Binding models,.



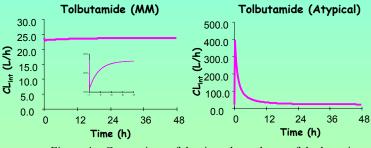


Figure 4 – Comparison of the time-dependence of the hepatic CL_{int} of tolbutamide predicted by Michaelis-Menten and Two-Site Binding models.

Discussion

The results indicate that for most drugs given in relatively low therapeutic doses (*e.g.* alprazolam) atypical enzyme kinetics are unlikely to be manifest in significant changes in *in vivo* clearance. Even with tolbutamide, given in relatively high doses, an impact is only evident when extreme values of the co-operativity numbers are applied. This is consistent with a lack of any discernable dose (concentration)-dependency in the *in vivo* kinetics of the drugs investigated in our study. PBPK modelling is a useful tool to clarify areas of *in vitro-in vivo* extrapolation where systematic *in vivo* studies on metabolic issues such co-operativity are lacking.

References

Chen, Q. *et al.*(2004), Effect of quinidine on the 10-hydroxylation of R-warfarin: species differences and clearance projection, *J Pharmacol Exp Ther* **311**, 307.

Lin, Y., *et al.* (2001), Substrate inhibition kinetics for cytochrome P450-catalyzed reactions. *Drug Metab Dispos*, **29**, 368.