

# THE IMPACT OF STUDY DESIGN ON THE ESTIMATION OF PARAMETERS DESCRIBING MECHANISM-BASED ENZYME INHIBITION

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## Introduction

Mechanism-based inhibition (MBI) is associated with irreversible or quasi-irreversible loss of enzyme activity. This type of inhibition may be a cause of particularly profound and long-lasting drug-drug interactions, requiring synthesis of new enzyme for return of normal enzyme function. Description of the effects of MBI is based on the *in vitro* estimation of two key parameters, namely  $k_{inact}$  (the maximum rate constant for inactivation) and  $K_I$  (the concentration of inhibitor which produces half the maximal rate of inactivation). The experimental procedure involves two steps (Silverman, 1995):

- 1- pre-incubation of enzyme with inhibitor (enzyme inactivation),
- 2- incubation with a probe substrate (measurement of residual enzyme activity).

To prevent further inactivation during step 2, enzyme and inhibitor should be diluted into a solution containing a high concentration of probe substrate. In this preliminary study we have investigated the impact of dilution and 3 different probe substrate concentrations on estimates of  $k_{inact}$  and  $K_I$  for 3,4-methylenedioxymethamphetamine (MDMA), an inhibitor of CYP2D6. Although recommendations have been made for the optimal design of MBI studies (Silverman, 1995), to our knowledge, the quantitative effects of experimental variables on  $k_{inact}$  and  $K_I$  have not been investigated systematically.

## Experimental Methods

Details of the MBI experiments are given elsewhere (Heydari *et al* 2004). In brief, the pre-incubation mixture contained MDMA (2 – 40  $\mu$ M and 0.625 – 12.5  $\mu$ M for experiments with and without dilution, respectively), KCL, an NADPH-generating system, potassium phosphate buffer (pH 7.4). All reactions were started by addition of yeast microsomes expressing CYP2D6 (20 pmol), and were carried out at 37°C. In the first experimental scheme, the CYP2D6 probe, dextromethorphan (DEX) (5, 10 or 20  $\mu$ M), was added directly to the pre-incubation mixture after different pre-incubation times (1 – 5 min) (no dilution). In the second scheme, aliquots (62  $\mu$ l) were taken from the pre-incubation mixture at different times (1 – 5 min) and added to incubation tubes containing DEX solution (final DEX concentration 20  $\mu$ M ; final volume 250  $\mu$ l ; a 4 fold dilution of the MDMA in the pre-incubation mixture). The reactions were stopped after 5 min by the addition of perchloric acid.

The inhibition of CYP2D6 was determined by the conversion of DEX to dextrophan in the presence and absence of MDMA.

## References

- Heydari A *et al* (2004) Mechanism-based inactivation of CYP2D6 by methylenedioxymethamphetamine, *Drug Metab Dispos* **32**: 1213 - 1217
- Ito K *et al.* (1998) Prediction of pharmacokinetic alterations caused by drug-drug interactions: metabolic interaction in the liver. *Pharmacol Rev* **50**(3): 387-412.
- Silverman RB (1995) Mechanism-based enzyme inactivation. *Methods in Enzymology* **249**: 240 - 283

## Data Analysis

Inactivation constants ( $k_{obs}$ ) were calculated from the % inhibition of the conversion of DEX to dextrophan (Ito *et al.* 1998). Hence,  $k_{inact}$  and  $K_I$  values were estimated using the following equation (Heydari *et al.* 2004) and non-linear regression.

$$k_{obs} = \frac{k_{inact} \times [I]}{K_I + [I]}$$

Statistical inferences were made using the z-test.

## Results

Figure 1 shows the model fits to the relationship between  $k_{obs}$  and MDMA concentration obtained from experiments with no dilution and using different DEX concentrations. Increasing concentrations of DEX were not associated with statistically significant differences in estimates of  $K_I$  or  $k_{inact}$  values ( $p > 0.05$ ).

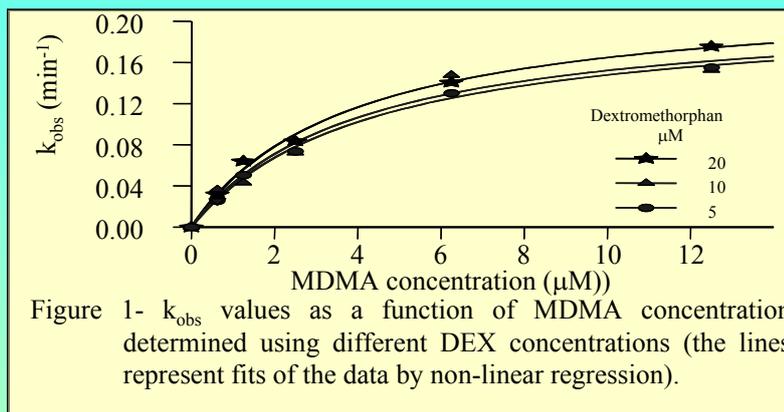


Figure 1-  $k_{obs}$  values as a function of MDMA concentration determined using different DEX concentrations (the lines represent fits of the data by non-linear regression).

A 4-fold dilution of the pre-incubation mixture resulted in higher values of  $k_{inact}$  and  $K_I$  relative to those observed without dilution ( $0.29 \pm 0.03$  SE vs  $0.21 \pm 0.02$   $\text{min}^{-1}$  ;  $p < 0.02$  and  $12.9 \pm 3.6$  vs  $3.5 \pm 0.8$   $\mu$ M ;  $p < 0.01$ ; respectively).

## Discussion

The observations indicate that probe substrate concentrations in excess of  $K_m$  ( $\approx 1.7$   $\mu$ M for DEX) will not have a major impact on  $K_I$  and  $k_{inact}$  values. Dilution after pre-incubation minimises any additional MBI that may occur during the subsequent incubation period and increases the value of  $K_I$ . However, the fold dilution that fully prevents inactivation during the incubation stage requires further evaluation as the literature is inconsistent in this regard (Ghanbari *et al.* in preparation)

These preliminary data emphasise the importance of study design in the accurate characterisation of the kinetics of MBI, with implications for *in vitro-in vivo* extrapolation of the extent of MBI-based drug-drug interactions.