

Unbound Total (Plasma) Clearance Approach in Interspecies Pharmacokinetics Correlation: Theophylline-Cimetidine Interaction

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INTRODUCTION

In a recent study (1) on the interspecies scaling of cimetidine-theophylline pharmacokinetics interactions, it was reported that the volume of distribution and half-life of theophylline in rats, rabbits, dogs and humans with or without coadministration of cimetidine are well correlated (correlation coefficients being 0.98 or 0.99) with their body weights. For theophylline hepatic clearance and unbound intrinsic clearance ($CL_{u,int}$), it was reported (1) that either in the presence or absence of cimetidine, the values for humans were smaller than expected from the extrapolation of three animal species. These results were attributed to that humans metabolize drugs at a lower rate, approximately $\frac{1}{7}$ of what would be expected on the basis of body weight (1). The lower rate of metabolism in humans was also reported to be related to the neoteny phenomenon since excellent correlation was obtained ($r = 0.999$; it was reported as 1.00) when all four species were analyzed after also considering maximum potential lifespan (MPL); correlation was made with $CL_{u,int} \times MPL$.

Unbound total (plasma) clearance (CL_u) is probably the most accurate kinetic parameter to directly measure the eliminating (excretion and biotransformation) capacity of a subject for a given compounds. It is probably also the most useful kinetic parameter since it is directly (for simplicity only linear kinetics is assumed throughout this study) related to unbound plasma concentration or its area under the curve that is in turn more directly related to the efficacy and/or toxicity of compounds (2). The unbound total (plasma) clearance approach has been employed (3,4) to successfully correlate the eliminating capacity for six β -lactam antibiotics (such as cefazolin and cefmetazole) in six species (mouse, rat, rabbit, dog, monkey and human). It has also been used to correlate the elimination of 15 extensively metabolized drugs (2), hydrochlorothiazide (5) and verapamil (6) between humans and rats. The rationale behind the interspecies correlation using unbound total clearance has been postulated (2). It was discussed (2) that mean CL_u values for all "nat-

ural" substances involved in the basal metabolism in different species can be scaled allometrically. Therefore, drugs may also resemble these "natural" substances in their clearance from the body according to the allometric relationship. The purpose of this communication is to report that such a simple approach can also be used to successfully correlate the eliminating capacity of theophylline in the absence or presence of cimetidine in the four species studied earlier (1).

METHODS

Mean total clearances (CL) of theophylline with or without co-administration of cimetidine in rats, rabbits, dogs, and humans were obtained from the same sources as reported earlier (1). The CL_u was calculated by (2)

$$CL_u = CL/f_u \quad (1)$$

where f_u is the fraction of drug unbound in plasma, being 0.58 for rabbits, dogs, and humans and 0.40 for rats (1).

Allometric analysis and statistical analysis of CL_u data from the four species were similarly made as described before (1).

RESULTS AND DISCUSSION

Mean CL_u and body weight (BW) of the four species employed in this study are summarized in Table I. The results of allometric analysis are shown in Fig. 1. It is of interest to note that the r is 0.989 for both correlations that is practically identical to those reported earlier (1) for correlation with the volume of distribution, half-life and $CL_{u,int} \times MPL$ indicating that humans are no different from the other three animal species in their ability to eliminate (in this case mainly biotransformation) theophylline. The observed CL_u for humans is lower than that predicted based on the three animal species. However, the difference is not statistically significant. The present and previous (1) results, however, do not necessarily indicate that drug interactions in general are scalable.

The unbound total clearance approach seems especially useful when there is a marked interspecies difference in plasma protein binding (2-6). Under such conditions half-life and plasma clearance may not be good parameters for correlation; this may often lead one to falsely (see examples in ref. 2) conclude that there is no general trend or predictability in eliminating capacity among different species. Thus, it appears useful first to conduct plasma protein binding study in humans and laboratory animals before performing any interspecies scaling in kinetics or dosing regimens.

Table I. Summary of Relevant Kinetic Parameters of Theophylline

Species	BW (kg)	CL_u^a (ml/min)	CL_u^b (ml/min)
Human	63	93.8	60.0
Dog	10.6	29.8	22.9
Rabbit	2.7	16.0	10.7
Rat	0.26	1.75	1.25

^a In the absence of cimetidine.

^b In the presence of cimetidine.

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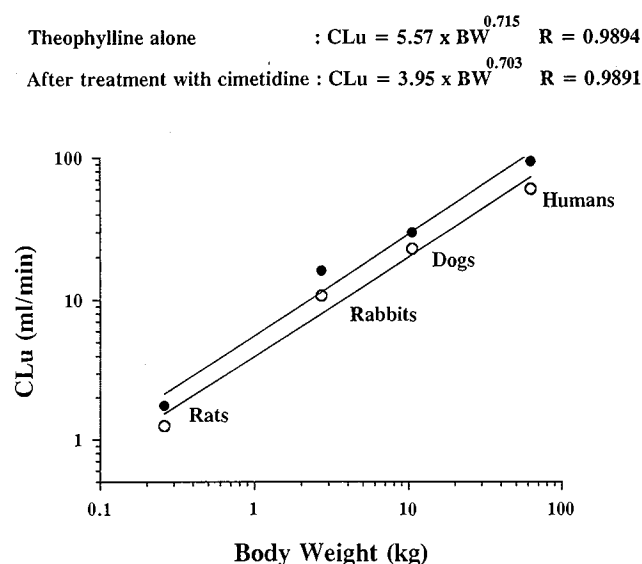


Fig. 1. Allometric relationship between unbound plasma clearance (CL_u) of theophylline alone (●) or after treatment with cimetidine (○) and the body weight (BW) of four species.

The $CL_{u_{int}}$ has been commonly used as a parameter to measure the intrinsic activity of the liver to metabolize a drug (1,7,8). This parameter is usually calculated based on a well-defined hepatic model that assumes well-stirring and lack of existence of any concentration gradients in the entire liver (8). The above basic assumptions are contradictory to the anatomy and physiology of the liver as well as the observed distribution profiles of drugs and metabolites in the liver (8,9 and references therein). Since many reported kinetic data can be adequately explained by the model-derived equations, this is then considered as supporting the validity of the model (8). Such an argument is no longer valid because the same model-derived equation can also be used to accurately or satisfactorily explain the absorption kinetics of drugs in a single-pass perfused intestinal loop (10) or the dialysis kinetics of compounds in a single-pass, artificial, rigid dialyzer (11,12). Both the perfused intestine and dialyzer are virtually unstirred preparations and their concentration gradients along the path of perfusion are well known. As expected, their absorption or dialysis kinetics can also be satisfactorily described by the classical tube model (10–12). If one accepts the notion (a reasonable one) that the liver more closely resembles the tube model than the well-stirred model, the $CL_{u_{int}}$ determined based on the well-stirred model may be much higher than the true $CL_{u_{int}}$, or that based on the tube model (11). In view of the above discussions it appears that one should be cautious in the use of $CL_{u_{int}}$ in pharmacokinetics studies. The difficulty of estimating accurate $CL_{u_{int}}$ has been extensively discussed (2). Furthermore, the role of neoteny in human evolution (13)

and drug metabolism (14) has been seriously challenged. Thus, the apparent success of using $CL_{u_{int}} \times MPL$ approach in interspecies correlation of drug metabolism may need to be re-examined. This is interesting in view of some success of using the simple CL_u approach for scale-up or correlation. This approach may be particularly valuable for comparison or correlation between humans and rats (2–5 and unpublished data).

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