

Prediction of Pharmacokinetics Prior to *In Vivo* Studies.

1. Mechanism-Based Prediction of Volume of Distribution

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ABSTRACT: In drug discovery and nonclinical development the volume of distribution at steady state (V_{ss}) of each novel drug candidate is commonly determined under *in vivo* conditions. Therefore, it is of interest to predict V_{ss} without conducting *in vivo* studies. The traditional description of V_{ss} corresponds to the sum of the products of each tissue:plasma partition coefficient ($P_{t:p}$) and the respective tissue volume in addition to the plasma volume. Because data on volumes of tissues and plasma are available in the literature for mammals, the other input parameters needed to estimate V_{ss} are the $P_{t:p}$'s, which can potentially be predicted with established tissue composition-based equations. *In vitro* data on drug lipophilicity and plasma protein binding are the input parameters used in these equations. Such a mechanism-based approach would be particularly useful to provide first-cut estimates of V_{ss} prior to any *in vivo* studies and to explore potential unexpected deviations between sets of predicted and *in vivo* V_{ss} data, when the *in vivo* data become available during the drug development process. The objective of the present study was to use tissue composition-based equations to predict rat and human V_{ss} prior to *in vivo* studies for 123 structurally unrelated compounds (acids, bases, and neutrals). The predicted data were compared with *in vivo* data obtained from the literature or at Roche. Overall, the average ratio of predicted-to-experimental rat and human V_{ss} values was 1.06 (SD = 0.817, $r = 0.78$, $n = 147$). In fact, 80% of all predicted values were within a factor of two of the corresponding experimental values. The drugs can therefore be separated into two groups. The first group contains 98 drugs for which the predicted V_{ss} were within a factor of two of those experimentally determined (average ratio of 1.01, SD = 0.39, $r = 0.93$, $n = 118$), and the second group includes 25 other drugs for which the predicted and experimental V_{ss} differ by a factor larger than two (average ratio of 1.32, SD = 1.74, $r = 0.42$, $n = 29$). Thus, additional relevant distribution processes were neglected in predicting V_{ss} of drugs of the second group. This was true especially in the case of some cationic-amphiphilic bases. The present study is the first attempt to develop and validate a mechanistic distribution model for predicting rat and human V_{ss} of drugs prior to *in vivo* studies. © 2002 Wiley-Liss, Inc. and the American Pharmaceutical Association J Pharm Sci 91:129–156, 2002

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INTRODUCTION

The volume of distribution at steady state (V_{ss}) and clearance processes characterize the dispo-

sition of drugs. The estimation of V_{ss} is therefore essential. V_{ss} corresponds to the equivalent plasma volume in which a drug is distributed into the body. In fact, V_{ss} would equal the plasma volume in addition to the sum of each tissue: plasma partition coefficient ($P_{t:p}$) multiplied with its respective tissue volume.¹ The $P_{t:p}$ represents the concentration ratio between a tissue and

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plasma at steady state. The estimation of V_{ss} *in vivo* by using $P_{t,p}$ data is a time consuming process in drug discovery and nonclinical development because determination of $P_{t,p}$ involves intravenous constant infusions to animals followed by an extraction and quantification of drugs from tissue homogenates.^{1,2} At present, V_{ss} under *in vivo* conditions is commonly determined by the product of clearance and mean residence time after a single intravenous dose of a drug according to the statistical moment theory.³ For efficient screening efforts it could be of interest to predict V_{ss} without conducting *in vivo* studies. In other words, it could be useful to predict V_{ss} *in vivo* from *in vitro* parameters routinely determined in drug discovery.

The tools available to date for predicting V_{ss} or V at non-steady state conditions are semi-empirical equations solved after their fit to *in vivo* data that have been previously collected.^{1,4,5} General or specific rules for the mechanisms underlying V_{ss} *in vivo* cannot easily be developed from such regression analyses. The semiempirical equations may provide predictions of V_{ss} , but only for the species and kinds of drugs for which the fitting exercises have previously been elaborated. The resulting regression analyses, however, show that drug lipophilicity and plasma protein binding are probably two main determinants of V_{ss} .^{1,4} Therefore, the prediction of V_{ss} prior to *in vivo* studies may potentially be performed from mechanistic equations that take into account the compound- and species-specific determinants of V_{ss} , and relate the compound-specific determinants to drug lipophilicity and plasma protein binding determined *in vitro*.

It has recently been demonstrated that *in vivo* $P_{t,p}$ of nonadipose and adipose tissues, the main compound-specific determinants of V_{ss} , can reasonably be predicted by using *in vitro* data on drug lipophilicity and plasma protein binding as sole input parameters in mechanistic tissue composition-based equations.^{6,7} The tissue composition-based equations have been validated with literature data on more than 310 $P_{t,p}$'s of mammals for about 80 structurally unrelated drugs. The mechanistic basis of these equations is that each tissue and plasma is a mixture of lipids, water, and plasma proteins in which the drug can be homogeneously distributed. In fact, the first term of these equations is based upon the drug lipophilicity in accordance with the lipophilicity-hydrophilicity balance of tissues

and plasma due to their lipid and water contents. The second term of the equations considers the binding to common proteins present in plasma and tissue interstitial space. Thus, the main assumption with these equations is that a passive diffusion process mainly governs tissue distribution of a drug to permit its partitioning into tissue lipids and water or into the extracellular water space only, and its reversible binding to common proteins present in plasma and tissue interstitial space. The drug-specific input parameters required in tissue composition-based equations (i.e., lipophilicity and plasma protein binding data) are generally available in drug discovery from *in vitro* studies. Therefore, the hypothesis of the present study was that V_{ss} *in vivo* could potentially be predicted on a mechanistic basis from data on $P_{t,p}$'s calculated with tissue composition-based equations.^{6,7} The other determinants of V_{ss} , the data on tissue volumes, can be obtained from the literature.

The objective of the present study was to use established mechanistic tissue composition-based equations to predict rat and human V_{ss} of 123 structurally unrelated drugs (acids, bases, and neutrals) without using *in vivo* data as input information.

METHODS

The overall method was based on a conventional equation that describes V_{ss} . In this equation the input parameters relative to tissue distribution (i.e., $P_{t,p}$'s) were predicted from established tissue composition-based equations, whereas the input parameters relative to the species (tissue volumes) were obtained from the literature. Two kinds of drugs were considered: those for which a homogeneous distribution into a tissue by passive diffusion can be assumed, and those for which a distribution mainly into the extracellular space can be assumed. In fact, the method was applied to 123 drugs covering a broad range of physicochemical (lipophilicity and ionization) and biochemical (plasma protein binding) properties. These drugs were chosen based on the availability of literature or Roche data on V_{ss} determined *in vivo*. The validation of the method consisted of comparing the predicted rat and human V_{ss} with corresponding experimental *in vivo* data obtained from the literature or at Roche.

Prediction of *In Vivo* V_{ss} in Rats and Humans for 123 Structurally Unrelated Drugs

V_{ss} referring to plasma pharmacokinetics was predicted with this conventional equation:¹

$$V_{ss} = (\Sigma V_t^* P_{t,p}) + (V_e^* E:P) + V_p \quad (1)$$

where V is the fractional body volume (L/kg) of a tissue (t), erythrocyte (e), and plasma (p), $E:P$ is the erythrocyte:plasma ratio, and $P_{t,p}$ is the nonadipose and adipose tissue:plasma partition coefficients (PCs). In eq. 1, the species-specific input parameters (V_t , V_e , V_p) can be obtained from the literature, whereas the drug-specific input parameters ($P_{t,p}$, $E:P$) can be estimated without using *in vivo* data. The estimation of $P_{t,p}$ and $E:P$ is described in the following section.

Prediction of $P_{t,p}$ of Nonadipose and Adipose Tissues With Tissue Composition-Based Equations for Drugs Which Distribute Homogeneously Into Tissues Mainly by Passive Diffusion

In predicting V_{ss} , the main tissues and plasma where a drug can be distributed were chosen to cover at least 80% of the body weight. Therefore, the nonadipose tissues represent the bone + marrows, brain, gut, heart, kidney, liver, lung, skeletal muscle, skin, and spleen, whereas the adipose tissue refers to subcutaneous white fat. The *in vivo* $P_{t,p}$ values of nonadipose and adipose tissues used in eq. 1 were calculated by using *in vitro* data on drug lipophilicity and plasma protein binding as sole input parameters in mechanistic tissue composition-based equations.^{6,7} The basis and assumptions of these equations are already presented in the Introduction section. Briefly, the drug would distribute homogeneously into each tissue (and plasma) mainly by passive diffusion. Consequently, the drug would partition between lipids and water as well as bind reversibly to common proteins present in plasma and tissue interstitial space. The tissue composition-based equations used to predict the $P_{t,p}$'s at the organ level are presented as follows:^{6,7}

$$P_{t,p \text{ nonadipose}} = \frac{[P_{o:w} \times (V_{nlt} + 0.3 \times V_{pht})] + [1 \times (V_{wt} + 0.7 \times V_{pht})]}{[P_{o:w} \times (V_{nlp} + 0.3 \times V_{php})] + [1 \times (V_{wp} + 0.7 \times V_{php})]} \times \frac{fu_p}{fu_t} \quad (2)$$

$$P_{t,p \text{ adipose}} = \frac{[D_{vo:w}^* \times (V_{nlt} + 0.3 \times V_{pht})] + [1 \times (V_{wt} + 0.7 \times V_{pht})]}{[D_{vo:w}^* \times (V_{nlp} + 0.3 \times V_{php})] + [1 \times (V_{wp} + 0.7 \times V_{php})]} \times \frac{fu_p}{1} \quad (3)$$

where $P_{o:w}$ is the *n*-octanol:buffer PC of the nonionized species at pH 7.4, $D_{vo:w}^*$ is the olive oil:buffer PC of both the nonionized and ionized species at pH 7.4, V is the fractional tissue volume content of neutral lipids (nl), phospholipids (ph), and water (w), t is tissue, p is plasma, and fu is unbound fraction. Each tissue is a mixture of lipids, water, and proteins with a pH set equal to 7.4. $P_{o:w}$ and $D_{vo:w}^*$ were used to estimate hydrophobic interactions of drugs with the biotic lipids of nonadipose and adipose tissues, respectively, whereas fu_p and fu_t were used to estimate a specific macromolecular binding to common proteins present in plasma and tissue interstitial space, respectively. A detailed derivation of eqs. 2 and 3 was recently provided by Poulin et al.^{6,7} Each input parameter of eqs. 2 and 3 can be estimated prior to *in vivo* studies. In fact, the physiological data were obtained from a review of the literature, whereas data on drug lipophilicity and protein binding were determined from *in vitro* studies or were calculated (see the section concerning the estimation of these input parameters).

Prediction of $P_{t,p}$ of Nonadipose and Adipose Tissues With Tissue Composition-Based Equations for Drugs Which Distribute Mainly Into Extracellular Space

For drugs distributing mainly into the extracellular space of tissues, their $P_{t,p}$'s of both nonadipose and adipose tissues cannot be calculated with eqs. 2 and 3. Alternatively, other tissue composition-based equations obtained from the literature⁶ were used to estimate their values of $P_{t,p}$ used in the general eq. 1:

$$P_{t,p \text{ nonadipose}} = \frac{V_{eist}}{V_{eisp}} \times \frac{fu_p}{fu_t} \quad (4)$$

$$P_{t,p \text{ adipose}} = \frac{V_{eist}}{V_{eisp}} \times \frac{fu_p}{1} \quad (5)$$

where V_{eist} is the fractional volume content of extracellular (interstitial) space in tissue (t) and plasma (p). This kind of physiological data can be obtained from the literature (see the section

concerning the estimation of these input parameters).

Estimation of the Input Parameters of Eqs. 1–5

Values of the Physiological Input Parameters

Table 1 summarizes the mean rat and human data on tissue volumes used in eq. 1 (V_t , V_e , V_p), tissue composition used in eqs. 2 and 3 (V_{nl} , V_{ph} , V_w), and extracellular space used in eqs. 4 and 5 (V_{eis}). The estimation of these data from the literature is presented in the Appendix.

Values of the Biochemical Input Parameters

The biochemical parameters refer to $E:P$ for erythrocyte:plasma ratio of eq. 1 as well as fu_p and fu_t of eqs. 2–5. The $E:P$ *in vivo* was estimated from data on blood:plasma ratio ($B:P$) determined *in vitro* and the hematocrit content in blood (Ht) (assumed to be 45%) according to this conventional equation:

$$E:P = [B:P - (1 - Ht)]/Ht \quad (6)$$

For drugs investigated in the present study, the mean data on $B:P$ determined *in vitro* are presented in Tables 2–4 with their corresponding

literature sources. However, the *in vitro* data on $B:P$ were not available for all drugs of this study. Alternatively, $B:P$ used in eq. 6 was set equal to 1 for the drugs that distribute homogeneously into tissues, whereas it was set equal to 0.55 ($1 - \text{hematocrit value}$) for the drugs that distribute only in the extracellular space. This has a minor impact on the calculated V_{ss} considering that a change of $B:P$ from 0.5 to 2.0 changes V_{ss} only by a factor of 15% in the worse cases (not shown).

Values of fu_p were obtained from experimental *in vitro* studies published in the literature (Tables 2–4). The mean experimental value of fu_p of each drug is presented in Tables 2–4 with the corresponding literature sources. For racemic drugs, the average for the two enantiomers is presented. For some drugs, fu_p is concentration dependent. Where a significant concentration dependency of fu_p was reported in the literature cited, fu_p values were chosen according to similar plasma concentrations in both the *in vitro* binding studies and in the *in vivo* distribution studies. The values of fu_t used for tissues were estimated from data on fu_p of plasma and tissue interstitial fluid-to-plasma concentration ratios of albumin, globulins, and lipoproteins; these ratios for mammals were estimated as equal to an intermediate value of approximately 0.5 based on literature

Table 1. Physiological Parameters for Volumes and Composition of Tissues of Adult Male Rats and Humans Used for Predicting V_{ss} ^a

Tissues	Tissue Volume		Tissue Composition (Volume Fraction of Wet Tissue Weight)						
	(Fraction of Body Weight; L/kg)		Water (V_w)		Neutral Lipids (V_{nl})		Phospholipids (V_{ph})		Extracellular (Interstitial) Space (V_{eis})
	Rat	Human	Rat	Human	Rat	Human	Rat	Human	Rat
Adipose	0.0761	0.11957	0.12	0.18	0.853	0.79	0.002	0.002	0.175
Bone	0.041476	0.085629	0.446	0.439	0.0273	0.074	0.0027	0.0011	0.42
Brain	0.0057	0.02	0.788	0.77	0.0392	0.051	0.0533	0.0565	0.162
Gut	0.027	0.0171	0.749	0.718	0.0292	0.0487	0.0138	0.0163	0.39
Heart	0.0033	0.0047	0.779	0.758	0.014	0.0115	0.0118	0.0166	0.156
Kidney	0.0073	0.0044	0.771	0.783	0.0123	0.0207	0.0284	0.0162	0.346
Liver	0.0366	0.026	0.705	0.751	0.0138	0.0348	0.0303	0.0252	0.159
Lung	0.005	0.0076	0.79	0.811	0.0219	0.003	0.014	0.009	0.484
Muscle	0.404	0.40	0.756	0.76	0.01	0.0238	0.009	0.0072	0.115
Skin	0.190	0.0371	0.651	0.718	0.0239	0.0284	0.018	0.0111	0.462
Spleen	0.002	0.0026	0.771	0.788	0.0077	0.0201	0.0136	0.0198	0.264
Plasma	0.0449	0.0424	0.96	0.945	0.00147	0.0035	0.00083	0.00225	1.00
Erythrocytes	0.0367	0.0347	—	—	—	—	—	—	—

^aMean values obtained from a review of the literature. See Appendix for the estimation and references concerning V_t , V_p , V_e , V_w , V_{nl} , V_{ph} , and V_{eis} . Data represent a rat of 250 g and a human of 70 kg.

—, Not used in the present study.

Table 2. Values of $\log P_{o,w}$, $\log D_{vo,w}$, pK_a^c , $f u_p^d$, and $B:P$ as well as Predicted and Experimental in vivo Values of V_{ss} in Adult Male Rats and Humans

Drugs	Volume of Distribution at Steady State (V_{ss}) (L/kg) ^f											
	$\log P_{o:w}^a$	$\log D_{vo:w}^b$	pK_a^c	$f u_p^d$		$B:P^e$		Rat		Human		References
				Rat	Human	Rat	Human	Predicted	Experimental	Predicted	Experimental	
Acids												
Acetaminophen	0.46	(-0.842)	9.38	0.88	1.0	(1.0)	(1.0)	0.64	0.54-0.797	0.63	0.85-1.42	[8,20-23]
Acetazolamide	-0.26	(-1.941)	7.4	—	0.044	—	(1.0)	—	—	0.32	0.38±0.05	[8,24,25]
Acitretin	6.40	(3.484)	(5.10)	0.01	—	(1.0)	(1.0)	4.60	3.57	—	—	[8,26,27]
Ascorbic acid	-1.85	-2.34	4.70	—	0.76	—	—	—	—	0.50	0.72	[4,8,28,29]
Methyl-ethyl barbiturate	0.08	(-1.338)	8.11	1.0	—	1.0	—	0.65	0.78	—	—	[8,30]
Butyl-ethylbar biturate	1.73	0.41	7.81	0.60	—	1.42	—	1.20	1.20	—	—	[8,10,30]
Pentyl-ethyl barbiturate	2.24	0.46	8.00	0.486	—	1.52	—	1.99	1.64	—	—	[8,10,30]
Hexyl-ethyl-bar biturate	2.46	(1.229)	7.74	0.18	—	1.0	—	2.16	1.80	—	—	[8,30]
Heptyl-ethyl barbiturate	2.91	(1.743)	7.78	0.07	—	1.0	—	3.07	1.67	—	—	[8,30]
Nonyl-ethylbar biturate	4.00	(2.970)	7.82	0.01	—	1.0	—	4.31	3.03	—	—	[8,30]
Captopril	0.34	(-4.671)	3.70	—	0.73	—	(1.0)	—	—	0.53	0.70-0.84	[8,31]
Dideoxyinosine	-1.24	(-2.743)	9.0	0.965	0.965	(1.0)	(1.0)	0.62	0.48-0.53	0.55	0.73-0.77	[8,32]
Isoniazid	-0.70	(-7.711)	1.82	—	1	—	(1.0)	—	—	0.56	0.59	[4,8]
Mephenytoin	1.69	(0.502)	8.51	0.63	—	(1.0)	(1.0)	1.20	1.29	—	—	[8,33-35]
Metronidazole	-0.02	(-6.152)	2.62	—	0.80	—	(1.0)	—	—	0.53	0.51-1.1	[8,36,37]
Pentobarbital	2.10	0.65	8.11	0.574	0.53	(1.0)	(1.0)	1.81	1.30±0.10	1.96	0.99	[4,8,38,39]
Phenobarbital	1.47	-0.244	7.30	—	0.543	—	0.861	—	—	0.88	0.54-0.62	[8,28,40,41]
Prazosin	1.28	(-0.373)	7.14	—	0.053	—	0.70	—	—	0.49	0.46-0.77	[8,42,43]
Tetrahydrocan nabinol	6.97	(6.421)	10.6	—	0.115	—	0.55	—	—	5.53	8.94±4.23	[8,44-46]
Thiopental	2.85	1.80	7.45	—	0.112	—	1.0	—	—	2.87	1.81-2.20	[8,38,47,48]
Bases												
Acebutolol	1.71	(-1.716)	9.67	—	0.743	—	(1.0)	—	—	1.25	1.17±0.26	[8,49,50]
Acyclovir	-1.56	(-3.084)	2.4; 3.5	—	0.85	—	(1.0)	—	—	0.52	0.69±0.19	[8,51]
Alfentanil	2.16	(1.007)	6.5	0.158	—	(1.0)	—	1.43	0.41-1.0	—	—	[8,52,53]
Alprenolol	3.34	(0.072)	9.70	—	0.24	—	(1.0)	—	—	3.04	3.37±1.24	[8,49,54]
Amsacrine	2.85	(1.511)	7.43	—	0.0298	—	(1.0)	—	—	2.24	1.56	[8,55,56]
Atropine	1.83	(-1.482)	9.57	—	0.89	—	(1.0)	—	—	1.56	2.90	[8,57,58]
Caffeine	-0.07	-1.08	10.4	—	0.80	—	1.0	—	—	0.53	0.73±0.02	[8,28,59]

Table 2. (Continued)

Drugs	Volume of Distribution at Steady State (V_{ss}) (L/kg) ^f											
	Log $P_{0:w}^a$	Log $D_{0:w}^{*b}$	pK _a ^c	$f u_p^d$		$B:P^e$		Rat		Human		
				Rat	Human	Rat	Human	Predicted	Experimental	Predicted	Experimental	
Carvedilol	4.19	(2.28)	8.40	—	0.0054	—	0.71	—	—	2.68	1.54 ± 0.11	[8,60,61]
Chlorpheniramine	3.38	(0.689)	9.13	—	0.27	—	1.34	—	—	3.26	3.17	[1,8]
Cimetidine	0.40	(−1.001)	6.80	—	0.80	—	(1.0)	—	—	0.56	1.0	[8,62]
Clonazepam	2.41	(1.337)	2.59	—	0.146	—	(1.0)	—	—	2.07	2.88	[4,8]
Cocaine	2.30	(−0.021)	8.61	0.63	—	1.0	—	2.23	2.8–3.3	—	—	[8,63,64]
Diazepam	2.99	2.07	3.5	0.137	0.022	0.38	0.58	4.10	4.50 ± 0.6	2.40	0.90–1.20	[65–68]
Diltiazem	2.84	(1.34)	7.7	0.25	0.22	(1.0)	1.0	3.46	3.60	3.09	2.91	[69–71]
Diphenhydramine	3.27	(0.685)	(9.0)	—	0.016	—	(1.0)	—	—	2.45	2.98	[4,8]
Epiroprim	2.89	1.70	6.96	0.097	0.11	1.0	1.0	3.16	4.63 ± 0.50	2.81	2.62 ± 0.36	[8]
Etomidate	3.05	(2.050)	4.5	0.191	0.235	(1.0)	(1.0)	4.70	4.03 ± 0.24	4.55	2.5 ± 1.1	[8,72–75]
Fentanyl	4.05	(1.565)	8.99	—	0.156	—	(1.0)	—	—	3.51	2.27–4.72	[8,76–79]
Gentamicin	−1.88	(−4.310)	8.20	—	0.73	—	(1.0)	—	—	0.49	0.45 ± 0.13	[8,80–82]
Ketanserin	3.29	(1.986)	7.46	—	0.05	—	(1.0)	—	—	2.88	3.7 ± 0.9	[8,83,84]
Lidocaine	2.44	(0.665)	8.01	0.38	0.296	1.27	0.90	2.37	2.13–3.11	2.18	1.08–1.51	[8,85–87]
Lorazepam	2.39	(−2.285)	11.0	—	0.107	—	(1.0)	—	—	1.69	1.60 ± 0.55	[8,88,89]
Lorainide	4.85	(1.954)	(9.5)	—	0.15	—	0.77	—	—	4.0	6.4 ± 2.4	[8,90]
Mefloquine (S ⁺)	4.49	2.56	9.1	0.024	—	(1.0)	—	4.62	7.50	—	—	[8]
Meperidine	2.72	(0.428)	8.63	—	0.357	—	(1.0)	—	—	2.73	4.17	[4,8,91]
Mepivacaine	1.95	(0.307)	7.76	—	0.225	—	0.9	—	—	1.22	1.10	[8,92,93]
Meptazinol	2.70	(−1.310)	(10.37)	—	0.729	—	(1.0)	—	—	3.30	2.36 ± 0.81	[94,95]
Methadone	3.93	1.15	8.94	0.254	0.127	(1.0)	(1.0)	5.23	6.55	3.10	4.20 ± 0.78	[8,28,96–98]
Mibefradil	3.86	(2.953)	4.80	—	0.005	—	(1.0)	—	—	2.70	3.04	[99,100]
Midazolam	2.95	(1.928)	5.82	0.027	0.019	0.55	0.55	2.91	2.54 ± 1.54	2.31	1.41 ± 0.39	[101]
Morphine	0.89	(−1.230)	8.21	—	0.76	—	(1.0)	—	—	0.64	0.92–2.12	[4,8,102]
Nicotine	1.17	−0.41	3.04; 7.8	0.84	—	0.80	—	0.80	0.98	—	—	[4,28,103–105]
Nitrazepam	2.23	1.43	3.4	—	0.123	—	1.0	—	—	1.77	1.90–2.14	[65,106]
Oxazepam	2.24	(−2.453)	2.35	0.15	0.035	(1.0)	(1.0)	1.59	1.32–4.59	1.42	0.59–1.31	[8,107–110]
Pentazocine	3.31	0.828	8.5	0.54	0.389	1.55	1.06	5.50	7.66	3.61	3.48–5.56	[1,65]
Phencyclidine	4.96	(2.176)	9.4	0.47	—	1.0	—	9.91	9.6–15.5	—	—	[8,111–114]
Prilocaine	2.11	(0.383)	7.9	—	0.715	—	(1.0)	—	—	2.12	3.70 ± 1.21	[8,115,116]
Propafenone	(3.37)	(−0.054)	(9.86)	0.01	—	(1.0)	—	3.52	4.90	—	—	[8,117,118]
Propranolol	3.20	(0.164)	9.45	—	0.099	—	(1.0)	—	—	2.61	2.79 ± 0.10	[8,49,119]
Quinidine	3.44	(1.297)	8.56	0.30	0.228	1.85	(1.0)	5.09	4.58 ± 0.80	3.45	3.50	[8,120,121]

Table 2. (Continued)

Drugs	Volume of Distribution at Steady State (V_{ss}) (L/kg) ^f											
	$f u_p^d$				$B:P^e$				Rat			
	Log $P_{0:w}^a$	Log $D_{vo:w}^b$	pK_a^c		Rat	Human	Predicted	Experimental	Predicted	Experimental	Human	References
Sematilide	(1.36)	(-2.127)	9.5; 7.6	—	—	0.96	—	—	0.98	0.74–0.77	—	[8,122,123]
Sildenafil	2.75	(1.665)	6.5	0.05	(1.0)	0.04	2.56	1.1–2.0	2.22	1.20	—	[124]
Sotalol	0.24	(-4.378)	8.3; 9.8	—	—	1.0	—	—	0.60	0.85	—	[4,8,49]
Sulpiride	0.57	(-2.443)	9.12	—	—	1.0	—	—	0.64	0.849±0.116	—	[8,125]
Theophylline	-0.02	(-2.799)	8.81	0.40	(1.0)	0.58	0.47	0.38	0.47	0.54	—	[8,126]
Verapamil	3.79	(1.752)	(8.49)	0.05	0.85	—	4.27	2.99–5.81	—	—	—	[8,127]
Neutrals												
2-Butanol	0.61	(-0.67)	—	1.0	(1.0)	—	0.70	0.73	—	—	—	[8,128,129]
<i>t</i> -Butyl alcohol	0.35	0.27	—	1.0	(1.0)	—	0.79	0.71–1.29	—	—	—	[8,128,130]
Cloprednol	1.68	(0.523)	—	—	—	0.16	—	—	0.88	0.81±0.14	—	[8,131]
Coumarin	1.39	(0.20)	—	—	—	0.853	—	—	1.08	1.29	—	[4,8]
Cyclophosphamide	0.63	(-0.648)	—	—	—	0.90	—	—	0.64	0.78	—	[4,8]
Cyclosporin	2.92	(1.906)	—	0.0602	1.283	0.0727	3.14	3.43±0.56	2.77	1.70–2.08	—	[8,132–135]
Cytosine arabinoside	-2.51	-4.0	—	—	—	0.98	—	—	0.55	0.44–0.86	—	[8,28,136,137]
Digitoxin	1.85	(0.713)	—	—	—	0.041	—	—	0.92	0.78±0.02	—	[8,138,139]
Digoxin	1.26	(0.055)	—	0.613	(1.0)	—	0.77	1.27	—	—	—	[8,140]
Enprofylline	0.33	(-0.922)	—	0.226	(1.0)	0.489	0.43	0.34	0.47	0.51	—	[8,141]
Ethanol	-0.31	-1.34	—	1.0	(1.0)	1.0	0.64	0.78–0.785	0.57	0.54±0.05	—	[8,28,128,142]
Ethoxybenzamide	0.77	(-0.491)	—	0.50	(1.0)	—	0.56	0.56	—	—	—	[8,143]
Ethylene glycol	-1.36	(-2.866)	—	1.0	(1.0)	—	0.63	0.3–0.4	—	—	—	[8,128,144]
Etretinate	(7.97)	(7.537)	—	0.01	(1.0)	—	4.65	3.07	—	—	—	[8,26,27]
Griseofulvin	2.18	(1.081)	—	—	—	0.20	—	—	1.72	1.40	—	[4,8]
Hydrocortisone	1.61	(0.445)	—	—	—	0.075	—	—	0.73	0.91	—	[4,8]
Methylprednisolone	1.82	(0.679)	—	—	—	0.232	—	—	1.13	1.38±0.25	—	[8,145]
Prednisolone	1.62	(0.456)	—	0.60	(1.0)	0.444	1.07	1.37±0.21	1.07	0.84–0.95	—	[8,145,146]
Propofol	3.79	(2.876)	—	—	—	0.022	—	—	3.04	5.82±2.94	—	[8,147]
Quercetin	(1.48)	(0.30)	—	—	—	0.02	—	—	0.60	1.13	—	[4,8]
Remoxipride	2.10	(0.992)	—	0.74	1.20	0.20	2.30	4.5±0.60	1.53	0.7±0.1	—	[8,148]
Thiopeta	0.53	(-0.759)	—	—	—	0.90	—	—	0.62	0.70±0.11	—	[8,149,150]
Toluene	2.65	2.78	—	—	—	1.0	—	—	20.1	15.4±4.4	—	[8,128,151,152]

Table 2. (Continued)

Values of $\log P_{o:w}$ (octanol:buffer PCs of nonionized species at pH 7.4), $\log D_{vo:w}^*$ (olive oil:buffer PCs of both nonionized and ionized species at pH 7.4 or 7.56), pK_a (dissociation constant), fu_p (unbound fraction in plasma), and $B:P$ (blood:plasma ratio). This table contains 89 drugs for which the predicted V_{ss} 's are within a factor of two of the corresponding experimental V_{ss} 's. A homogeneous distribution was assumed in predicting V_{ss} .

^aMean experimental data on $\log P_{o:w}$ obtained from the cited literature except for diltiazem, mibefradil, midazolam, and mefloquine (Hoffmann-La Roche). However, for etretinate, propafenone, quercetin, and sematilide, data on $\log P_{o:w}$ were calculated using the fragment constant method of Meylan and Howard⁸ in the absence of experimental data (values in brackets).

^bMean experimental data on $\log D_{vo:w}^*$ obtained from the cited literature except for diazepam, epiroprim, mefloquine, nitrazepam, and pentazocine (Hoffmann-La Roche). For toluene, its $\log D_{vo:w}$ was obtained by dividing the oil:air PC with saline:air PC. However, for drugs for which experimental data on $\log D_{vo:w}$ were not available in the literature or at Roche, they were calculated from data on $\log P_{o:w}$ and pK_a according to eqs. 7–12 as described in the Methods section (values in brackets).

^cMean experimental data on pK_a obtained from the cited literature except for diltiazem, epiroprim, lorazepam, mefloquine, mibefradil, midazolam, and oxazepam (Hoffmann-La Roche). However, for acitretin, diphenhydramine, lorcinide, neptazolin, propafenone, and verapamil the data were calculated using the method of ISIS BaseTM, version 2.2 in the absence of experimental data (values in brackets).

^dMean experimental data on fu_p in *vitro* obtained from the cited literature except for epiroprim (rat and human data), mefloquine (rat data), and midazolam (rat data) (Hoffmann-La Roche).

^eMean experimental data on $B:P$ in *vitro* obtained from the cited literature except for epiroprim (rat and human data) and midazolam (rat and human data) (Hoffmann-La Roche). However, for drugs for which experimental data on $B:P$ were not available in the literature or at Roche, a default value of 1 was presented in the table as explained in the Methods section (values in brackets).

^fData predicted from eqs. 1–3. Experimental data on V_{ss} in *vitro* obtained from the cited literature except for epiroprim (rat and human data), mefloquine (rat data), and midazolam (rat data) (Hoffmann-La Roche). The experimental V_{ss} data represent either the means alone, or the means \pm SD (single study), or the reported ranges (several studies). Experimental *in vivo* V_{ss} data were obtained for rat and human of approximately 0.250 kg and 70 kg, respectively.

—, Not used or calculated.

information.⁶ In fact, fu_t was set equal to $1/(1 + \{(1 - fu_p)/fu_p\} \times 0.5)$. The rationale of the mechanism-based estimation of fu_t from fu_p was provided in Poulin and Theil.⁶

Values of the Physicochemical Input Parameters

$P_{o:w}$ (*n*-octanol:buffer PC of nonionized species) used in eq. 2 was experimentally determined *in vitro* with conventional shake-flask methods or was calculated with a fragment constant method. The experimental or calculated $P_{o:w}$ data of several drugs are now available in a database named KowWin.⁸ This database was used in the present study to get data on $P_{o:w}$ of most drugs. For some drugs, however, data on $P_{o:w}$ are not available in the database. Therefore, $P_{o:w}$ was obtained from other literature sources including Roche data. The values of $\log P_{o:w}$ of all drugs and their references are presented in Tables 2–4.

$P_{o:w}$ is a parameter commonly determined for drugs. The availability of experimental data on $P_{o:w}$ explains why it was used in eq. 2 to estimate drug partitioning between neutral lipids and water of nonadipose tissues. Furthermore, the neutral nonpolar lipids of nonadipose tissues are mainly mixtures of triglycerides and cholesterol. The *n*-octanol may mimic the lipophilicity of such mixtures. The adipose tissue lipids, however, are mostly composed of triglycerides. In this case, it was demonstrated that *n*-octanol is not a good surrogate of triglycerides.⁷ Alternatively, olive oil was suggested as an adequate surrogate of triglycerides.⁷ Therefore, $D_{vo:w}^*$ (olive oil:buffer PCs of nonionized and ionized species) was used in eq. 3 to estimate drug partitioning between lipids and water of adipose tissue. The values of $\log D_{vo:w}^*$ determined *in vitro* with conventional shake-flask methods were obtained from the literature for some drugs (Tables 2–3). For several drugs, however, an experimental value of $\log D_{vo:w}^*$ is not available in the literature. Alternatively, the value of $\log D_{vo:w}^*$ was estimated from data on $\log P_{o:w}$ as follows. It is conventionally considered that the ionized species is soluble only in the buffer phase, but the nonionized species is soluble in both the aqueous and the lipophilic phases.⁹ As reported previously, $D_{vo:w}^*$ takes into account the nonionized and ionized species in the aqueous phase compared to $P_{o:w}$ that considers only the nonionized species. In reporting $D_{vo:w}^*$ on a basis of nonionized species only in the aqueous phase to get $D_{vo:w}$, a linear regression analysis can be obtained between

Table 3. Values of Log $P_{o:w}$, Log $D_{vo:w}^*$, pK_a , $f u_p$, and $B:P$, As Well As Predicted and Experimental $In Vivo$ Values of V_{ss} in Adult Male rats and Humans

Drugs	Log $P_{o:w}^a$	Log $D_{vo:w}^{*a}$	pK_a^a	Volume of Distribution at Steady State (V_{ss}) (L/kg) ^c								Reference	
				$f u_p^a$		$B:P^b$		Rat		Human			
				Rat	Human	Rat	Human	Predicted	Experimental	Predicted	Experimental		
Acids													
Barbital	0.65	-0.59	7.75	0.964	—	1.0	—	0.70	2.31	—	—	—	[8,38,30]
Propylethyl-barbiturate	0.66	-0.14	7.77	0.86	—	1.0	—	0.69	1.58	—	—	—	[8,10,30]
Octylethyl-barbiturate	3.78	(2.713)	7.78	0.03	—	1.0	—	4.57	2.20	—	—	—	[8,30]
Ketoprofen	3.12	(-0.822)	4.45	0.029	—	(1.0)	—	3.17	1.19	—	—	—	[8,153]
Thiopental	2.85	1.80	7.45	0.131	—	0.922	—	3.34	0.69-1.35	—	—	—	[8,38,154,155]
Bases													
Alfentanil	2.16	(1.007)	6.5	—	0.074	—	(1.0)	—	—	1.38	0.226 ± 0.048	—	[8,52,156]
Amlodipine	3.00	(0.768)	8.6	0.06	0.02	(1.0)	(1.0)	3.03	32.0	2.28	21.0	—	[8,157,158]
Amiodarone	6.66	(4.348)	9.12	0.05	—	1.1	—	6.29	16.35 ± 0.71	—	—	—	[159,160]
Biperiden	4.25	(1.972)	8.8	0.135	—	1.16	—	5.41	14.0 ± 1.4	—	—	—	[8,65,161]
Bupivacaine	3.41	(1.623)	8.16	—	0.044	—	0.60	—	—	2.70	0.63-0.76	—	[8,92,93]
Chlorpromazine	5.19	(2.531)	9.3	0.106	0.043	(1.0)	1.56	6.18	29.1	3.4	8.8-11.2	—	[1,65,162,163]
Doxorubicin	1.27	(-0.798)	8.2	—	0.277	—	(1.0)	—	—	0.61	15.4 ± 3.8	—	[8,164,165]
Disopyramide	2.58	(-1.474)	10.4	0.235	0.325	(1.0)	(1.0)	2.36	0.63	2.35	0.78	—	[1,8,166]
Etidocaine	3.69	(2.288)	7.7	—	0.06	—	0.6	—	—	3.31	1.15	—	[8,92,93]
Exaprolol	3.68	(0.877)	9.27	0.54	—	1.25	—	6.15	59.4	—	—	—	[167]
Haloperidol	4.30	(2.161)	8.66	—	0.125	—	(1.0)	—	—	4.06	17.8 ± 6.5	—	[8,168,169]
Imipramine	4.80	1.93	9.5	0.24	0.145	1.67	(1.0)	6.45	19.9	3.93	14.2 ± 1.4	—	[1,8,28,170]
Procainamide	0.88	(-2.176)	9.2	0.92	—	(1.0)	—	0.71	1.77	—	—	—	[8,171]
Methylphenidate	0.20	-0.26	8.77	0.77	—	(1.0)	—	0.61	1.47	—	—	—	[8,172,173]
Ropivacaine	2.90	(1.129)	8.07	—	0.06	—	0.69	—	—	2.32	0.55 ± 0.07	—	[8,92,93,174]
Trimethoprim	0.91	(-0.799)	7.12	—	0.304	—	(1.0)	—	—	0.52	1.96	—	[4,8]
Neutrals													
Digoxin	1.26	(0.055)	—	—	0.78	—	(1.0)	—	—	0.90	6.9-7.3	—	[8,175]

Table 3. (Continued)

Drugs	Volume of Distribution at Steady State (V_{ss}) (L/kg) ^c									
	f_u^a			$B:P^b$			Rat			Human
	Rat		Human	Rat		Human	Predicted		Experimental	Reference
	$P_{o:w}^a$	$\log D_{vo:w}^*$	pK_a^a	pK_a^a	$\log D_{vo:w}^*$	pK_a^a	$P_{o:w}$	$\log D_{vo:w}^*$	pK_a^a	
Felodipine	3.86	(2.954)	—	0.004	—	(1.0)	—	2.68	10.3 ± 3.0	[8,176]
Zwitterionics										
Ciprofloxacin	0.28	(0.986)	6.2; 8.6	0.52	—	(1.0)	—	0.47	1.74–1.90	[8,177–179]
Enoxacin	–0.20	(–1.654)	6.3; 8.3	0.485	—	(1.0)	—	0.44	1.96 ± 0.43	[8,177,180,181]

This table contains 25 drugs for which the predicted V_{ss} 's are greater by a factor of two than the corresponding experimental V_{ss} 's. A homogeneous distribution was assumed in predicting V_{ss} .

^aMean experimental data on $\log P_{o:w}$, $\log D_{vo:w}^*$, pK_a , and f_u *in vitro* obtained from the cited literature. However, for drugs for which experimental data on $\log D_{vo:w}^*$ were not available in the literature they were calculated from $\log P_{o:w}$ and pK_a according to eqs. 7–12 as described in the Methods section (values in brackets).

^bMean experimental data on $B:P$ *in vitro* obtained from the cited literature. However, for drugs for which experimental data on $B:P$ were not available in the literature, a default value of 1 was presented as explained in the Methods section (values in brackets).

^cData predicted from eqs. 1–3. Experimental data on V_{ss} *in vivo* obtained from the cited literature. The experimental V_{ss} data represent either the means alone, the means ± SD (single study), or the ranges (several studies). Experimental *in vivo* V_{ss} data were obtained for rat and human of approximately 0.250 kg and 70 kg, respectively. —, Not used or calculated.

experimental data on $\log D_{vo:w}$ and $\log P_{o:w}$ of several organic chemicals (acids, bases, neutrals):¹⁰

$$\log D_{vo:w} = 1.115 \times \log P_{o:w} - 1.35, \quad (7)$$

$$n = 104, r = 0.99$$

For the prediction of $P_{t:p}$ of adipose tissue with eq. 3, data on $D_{vo:w}^*$ are required instead of $D_{vo:w}$.⁷ Therefore, data on $\log D_{vo:w}^*$ were obtained as follows from data on $\log D_{vo:w}$ calculated with eq. 7. For this, it was assumed that $D_{vo:w}^*$ and $D_{vo:w}$ differ by a factor corresponding to the ionized species in the aqueous phase based on classical Henderson-Hasselbalch equations:

$$1. \text{ Monoprotic acid: } \log D_{vo:w}^* = \log D_{vo:w} - \text{Log} (1 + 10^{\text{pH} - \text{p}K_a}) \quad (8)$$

$$2. \text{ Monoprotic base: } \log D_{vo:w}^* = \log D_{vo:w} - \text{Log} (1 + 10^{\text{p}K_a - \text{pH}}) \quad (9)$$

$$3. \text{ Diprotic acid: } \log D_{vo:w}^* = \log D_{vo:w} - \text{Log} (1 + 10^{\text{pH} - \text{p}K_{a1} + \text{pH} - \text{p}K_{a2}}) \quad (10)$$

$$4. \text{ Diprotic base: } \log D_{vo:w}^* = \log D_{vo:w} - \text{Log} (1 + 10^{\text{p}K_{a1} - \text{pH} + \text{p}K_{a2} - \text{pH}}) \quad (11)$$

$$5. \text{ Zwitterionic: } \log D_{vo:w}^* = \log D_{vo:w} - \text{Log} (1 + 10^{-\text{p}K_{a2} + \text{pH} + \text{p}K_{a1} - \text{pH}}),$$

$$\text{p}K_{a1} = \text{acid}, \text{p}K_{a2} = \text{base} \quad (12)$$

6. Neutral compound (see eq. 7).

To estimate $D_{vo:w}^*$ with the above equations, additional data on $\text{p}K_a$ are required. The experimental or calculated value of $\text{p}K_a$ of each drug is presented in Tables 2–3, which includes also the corresponding literature sources. The calculated values of $\log D_{vo:w}^*$ are presented in Tables 2–3. Note that data on $P_{o:w}$ and $D_{vo:w}^*$ expressed in a logarithmic form need to be transformed in the inverse log form before their use in eqs. 2 and 3 to predict $P_{t:p}$ of nonadipose and adipose tissues, respectively.

In the literature cited in the tables, data on $\log P_{o:w}$ and $\log D_{vo:w}^*$ were determined at a temperature of about 25 or 37°C in most cases. However, the known data on temperature dependence of $\log P_{o:w}$ and $\log D_{vo:w}^*$ as well as other solvent:buffer PCs indicate that the temperature dependence is relatively minor for several organic compounds ($\Delta \log \text{ solvent:buffer PCs} = 0.009/^\circ\text{C}$).^{10–12} A temperature dependence of $\log P_{o:w}$ and $\log D_{vo:w}^*$, however, may be a potential source of error in

Table 4. Values of $\log P_{o:w}$, pK_a , and $f_{u,p}$ As Well As Predicted and Experimental *In Vivo* Values of V_{ss} (L/kg) in Adult Male Rats and Humans

Drugs	$\log P_{o:w}^a$	pK_a^b	$f_{u,p}$	$B:P$	Rat Data ^c		References
					Predicted V_{ss}	Experimental V_{ss}	
Cefazolin	-0.58	2.33	0.272	(0.55)	0.16	0.208	[8,182]
Ceftazidime	-1.36	—	0.10	(0.55)	0.14	0.24	[183]
Ceftriaxone	-1.47	2.33	0.725	(0.55)	0.21	0.33 ± 0.19	[184]
Glycyrrhetic acid	6.90	5.26	0.17	(0.55)	0.15	0.178 ± 0.068	[8,185]
Glycyrrhizin	2.80	2.6,2.6,5.3	0.05	(0.55)	0.14	0.184 ± 0.015	[8,186]
Interferon	—	—	1.0	(0.55)	0.23	0.26	[187]
Lamifiban	—	—	0.92	(0.55)	0.23	0.24	[188]
Nicotinic acid	0.36	4.75	0.55	(0.55)	0.19	0.289	[8,189]
Tolcapone	3.13	4.93	0.001	(0.55)	0.13	0.21	—

This table contains nine drugs for which tissue distribution is assumed to be restricted to the extracellular space based on the literature and for which the predicted V_{ss} 's are within a factor of two of the corresponding experimental V_{ss} 's.

^aMean experimental data on $\log P_{o:w}$ obtained from the cited literature except for ceftriaxone and tolcapone for which $\log P_{o:w}$ was estimated using the fragment constant method of Meylan and Howard.⁸ Data on $\log P_{o:w}$ were not used for predicting V_{ss} of these types of drugs, which do not penetrate into cells (see the Methods section).

^bData on pK_a estimated with the method of ISIS BaseTM except for nicotinic acid for which the experimentally determined pK_a was obtained from the literature.⁸ Data on pK_a were not used for predicting V_{ss} of this type of drug, which does not penetrate into cells (see the Methods section).

^cAll data of tolcapone were obtained at Hoffmann-La Roche. The experimental data on $B:P$ *in vitro* were not available in the literature. Therefore, a default value of 0.55 (1 – hematocrit value) was used for each drug that does not penetrate into cells (values in brackets). Data on $f_{u,p}$ represent only the means (single study), whereas data on V_{ss} represent the means alone or the means \pm SD (single study). Data on V_{ss} were predicted from eqs. 1, 4, and 5 or were experimentally determined under *in vivo* conditions for rat and human of approximately 0.250 kg and 70 kg, respectively.

—, Data not available.

calculating $P_{t,p}$'s at 37°C. Nevertheless, PC data at 25 or 37°C were used in absence of these data at 37°C for all drugs. It is of interest to remember that the main purpose of the present study was only to verify if reasonable rough first estimates of V_{ss} *in vivo* could be obtained as required in early drug discovery. For the same reason, $\log P_{o:w}$ and $\log D_{vo:w}^*$ were obtained only at pH 7.4 or 7.56 considering that: these data are not available at other pHs based on the literature, the physiological pH of plasma and extracellular space of tissues is close to 7.4, the overall pH value of an intact tissue is currently unknown for *in vivo* conditions, and the change of *in vivo* $P_{t,p}$ with pH is also unknown.

Validation of the Mechanism-Based Method Used to Predict V_{ss}

The values of V_{ss} of 123 structurally unrelated drugs predicted based on eq. 1 were compared with corresponding *in vivo* values available in the literature (most cases) or obtained at Hoffmann-La Roche. The *in vivo* values of V_{ss} for all drugs and their corresponding literature sources are

presented in Tables 2–4. Only V_{ss} 's (i.e., not an apparent V , $V\beta$, or V_z) estimated from plasma concentration-time profiles determined after intravenous administrations to rats and humans were considered for the purpose of the present study. For a few drugs, however, the values of V_{ss} are expressed on a basis of blood concentration. In this case, V_{ss} was multiplied with the blood: plasma ratio to get V_{ss} for plasma. As much as possible, values of V_{ss} 's were obtained for healthy adult male rats ($\cong 250$ g) and humans ($\cong 70$ kg). It was observed that V_{ss} is concentration-dependent for some drugs. In this context, see the Methods section concerning the estimation of $f_{u,p}$.

The experimental or calculated values of V_{ss} , $f_{u,p}$, $B:P$, $P_{o:w}$, or $D_{vo:w}^*$ were taken from several literature sources of the last 30 years. It is important to note that in collecting these values, no effort or judgment was made to differentiate experimental and analytical errors from true variability, or to classify the values according to the techniques used for their determination/estimation. Sensitivity and variability analyses are beyond the scope of the present study considering that the main objective was to verify

if the current methodology could provide rough first-cut estimates of V_{ss} prior to any *in vivo* experiments as required in drug discovery.

RESULTS

It is of interest to note that fitting exercises were omitted to improve the predictions. In fact, the current study presents only first predictions of V_{ss} by using one set of specified data as the input in eq. 1. The experimental values of V_{ss} *in vivo* obtained from the literature for 123 drugs were compared with the corresponding predicted values. For the 123 drugs investigated in the present study the overall average ratio of predicted-to-mean experimental rat and human V_{ss} values was 1.06 (SD = 0.817, $r = 0.78$, $n = 147$; Tables 2–4, Figure 1A). In fact, 80% of all predicted values were within a factor of two of the corresponding experimental values obtained from the literature. The drugs can therefore be separated into two groups.

First Group

The first group contains 98 drugs for which the predicted V_{ss} 's were within a factor of two of those experimentally determined (average ratio of 1.01, SD = 0.39, $r = 0.93$, $n = 118$; Tables 2 and 4, Figure 1B). These 98 drugs include 89 drugs for which a homogeneous tissue distribution by passive diffusion can be assumed (Table 2), and nine other drugs for which a distribution limited to the extracellular space can be assumed (Table 4). The current assumptions on tissue distribution considered for predicting V_{ss} are probably not violated for these 98 drugs, which include acids, bases, and neutral compounds.

Second Group

The second group contains 25 drugs (acids, bases, zwitterions) for which the predicted and experimental V_{ss} 's differ by a factor larger than two (average ratio of 1.32, SD = 1.74, $r = 0.42$, $n = 29$; Table 3, Figure 1C). It was assumed that these 25 drugs distribute homogeneously into tissues by passive diffusion. Thus, additional relevant distribution processes were probably neglected in predicting V_{ss} of these drugs. This was true especially in the case of some cationic-amphiphilic bases for which their V_{ss} was largely under predicted (Table 3).

DISCUSSION

Distribution and clearances are input parameters in empirical PK compartment models and physiologically based PK models (PBPK).^{3,30} For efficient screening efforts with these models it is essential to obtain rough predictions of V_{ss} and clearances from compound-specific determinants routinely determined in early drug discovery. In the present study, a mechanism-based method for predicting rat and human V_{ss} under *in vivo* conditions has been developed and validated for 123 structurally unrelated drugs. The compound-specific determinants used for predicting V_{ss} with this method could be generated in drug discovery without conducting *in vivo* studies. During the drug development process it is evident that the quantity and quality of experimentally determined data on V_{ss} *in vivo* increase with time, which should permit comparison to the predicted V_{ss} . The present prediction method of V_{ss} built on a mechanism-based framework would permit the understanding of unexpected large deviations that may be observed between sets of predicted and experimental V_{ss} data. In general, a significant under-prediction of V_{ss} *in vivo* by the present method may suggest a drug accumulation into tissues due to additional specific binding and/or uptake processes, whereas an over-prediction of V_{ss} may suggest a limited permeation into cells of the peripheral compartment. Figure 1C presents large under- and over-predictions of V_{ss} especially for 25 drugs.

The drugs investigated in the present study have been classified into two groups. The first group represents 98 drugs for which the predicted and experimental V_{ss} 's differ by a factor of less than two (Tables 2 and 4), whereas the second group represents 25 drugs for which the predicted and experimental V_{ss} 's differ by a factor larger than two (Table 3). However, it appears that the current assumptions on tissue distribution used to predict V_{ss} are not violated especially for the first group of drugs. In other words, relevant additional distribution processes have probably been neglected in predicting V_{ss} of the second group of drugs. For the early drug discovery, V_{ss} was judged as accurately predicted solely for drugs of the first group. An obvious matter of concern with the present study is the uncertainty associated with each input parameter of the series of equations used to predict V_{ss} . The mean data of these parameters on tissue composition as well as drug lipophilicity and protein binding were

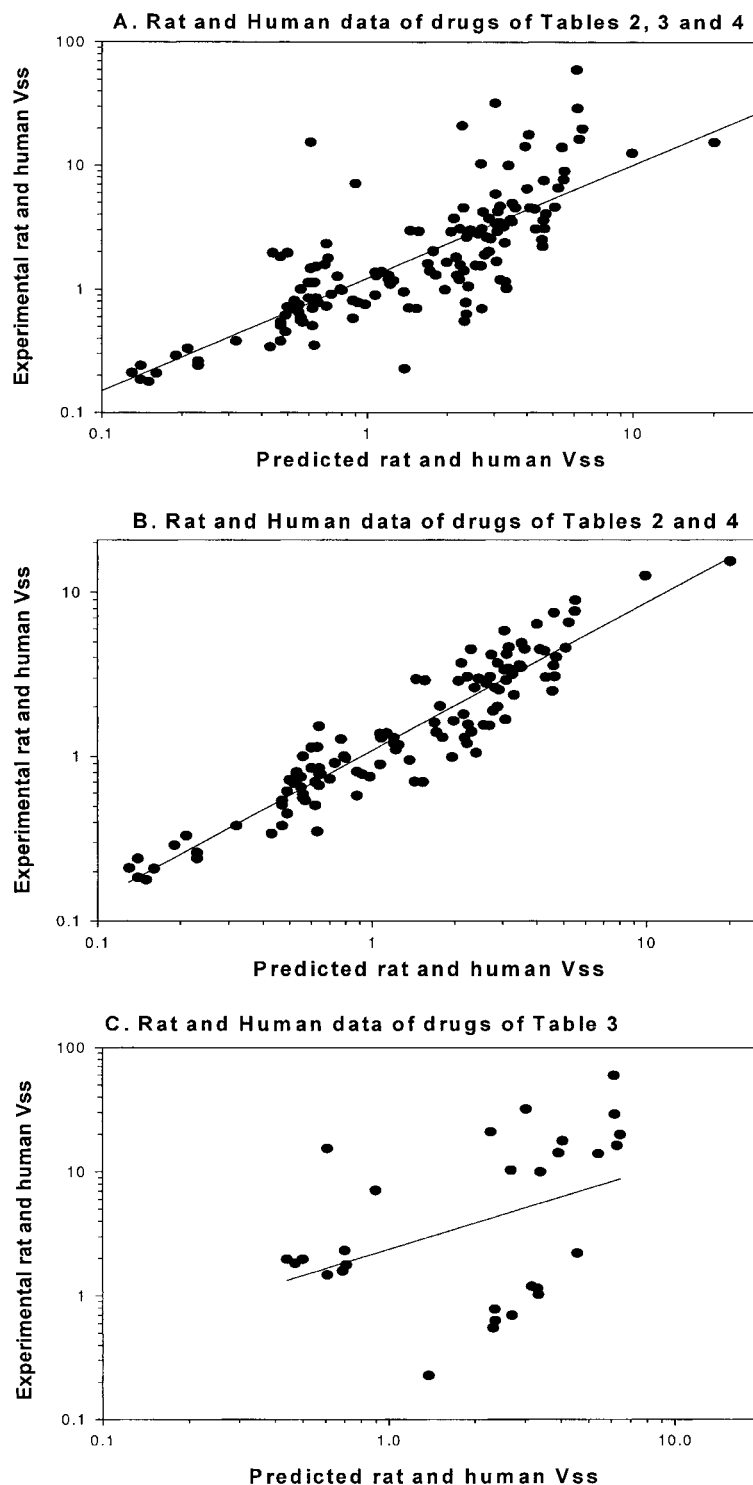


Figure 1. Relationship between predicted and experimentally determined V_{ss} of 123 drugs of Tables 2–4 (A), of 98 drugs of Tables 2 and 4 (B), and of 25 drugs of Table 3 (C). The line indicates the best fit for (A) [$\log y = 0.91383 \cdot \log x + 0.0892$, $r = 0.78$, $n = 147$, average ratio predicted-to-experimental $V_{ss} = 1.06 \pm 0.82$], (B) [$\log y = 0.90180 \cdot \log x + 0.03561$, $r = 0.93$, $n = 118$, average ratio predicted-to-experimental $V_{ss} = 1.01 \pm 0.39$], and (C) [$\log y = 0.70273 \cdot \log x + 0.37492$, $r = 0.42$, $n = 29$, average ratio predicted-to-experimental $V_{ss} = 1.32 \pm 1.74$]. Sigmaplot[®] version 4.0 was used for plotting.

obtained from various literature sources. Elaborated statistical and sensitivity exercises need to be done, but this was not necessary at this step because the main objective of the present study was only to develop a method to get first rough estimates of V_{ss} *in vivo* as required in drug discovery.

Predictions of V_{ss} *in vivo* within the given accuracy criteria were particularly obtained for the 89 drugs of Table 2, for which a homogeneous distribution into tissues mainly by passive diffusion can be assumed, and for the nine other drugs of Table 4, for which a tissue distribution limited to extracellular space can be assumed. An explanation for such accurate predictions for most drugs may be that muscle and fat are probably two main determinants of V_{ss} (based on their volume and/or lipid content) for which the $P_{t:p}$ of several drugs can reasonably be predicted with tissue composition-based equations.^{6,7} Nevertheless, the results for these 98 drugs (Tables 2 and 4) suggest that V_{ss} is mainly governed by partitioning into total lipids and water (or partitioning into extracellular space only) and a reversible binding to common proteins present in plasma and interstitial space (i.e., albumin, globulins, lipoproteins as HDL and LDL). However, some drugs of this study (e.g., disopyramide, imipramine, methadone, mibefradil, propranolol, quinidine) can bind predominantly to other plasma proteins (e.g., alpha-glycoprotein).¹³ The rat and human V_{ss} 's of these drugs are also accurately predicted except for disopyramide and imipramine (Tables 2 and 3; for imipramine, another explanation for the under-prediction of its V_{ss} is presented later in the Discussion section). Therefore, a binding to alpha-glycoprotein can potentially be accounted for in predicting V_{ss} . The rationale for that can be the ratio fu_p/fu_t of eq. 2 used to calculate the $P_{t:p}$'s. This ratio was estimated from data on tissue interstitial fluid-to-plasma concentration ratios of albumin, globulins, and lipoproteins⁶; similar ratios may also be found for alpha-glycoprotein. This cannot be confirmed in the absence of experimental data for alpha-glycoprotein.

Specific binding to cell macromolecules other than lipids has been neglected in predicting V_{ss} . Regarding the results of Table 2, it may be suggested that specific binding of drugs to cell macromolecules, if present, does not influence on a quantitative basis the prediction of V_{ss} for 89 drugs. This is also supported by the accurate predictions of rat and human V_{ss} of cyclosporin,

quinidine, and methotrexate (Table 2) by neglecting a saturable specific binding to particular cell components reported for these drugs.^{14,132} An explanation for this may be the much higher content of lipids and plasma proteins in tissues compared to cell macromolecules. Consequently, this may be a factor explaining that nonspecific binding to lipids and specific binding to plasma proteins (this study) might be the main "apparent" determinants of tissue distribution of most drugs. This needs to be verified, but the results in Table 2 suggest that specific binding to cell macromolecules can be neglected in predicting V_{ss} *in vivo* of several drugs. However, there are exceptions. For instance, for a drug binding extensively to DNA (e.g., doxorubicin),¹⁵ V_{ss} was largely under-predicted (Table 3). In general, a significant specific binding to cell macromolecules may potentially explain unexpected under-predictions of V_{ss} *in vivo* observed for other drugs of Table 3. In this context, additional *in vitro* data on macromolecular binding are needed to predict V_{ss} accurately.

Ionic interactions between drugs and charged lipids of cell membranes as well as a pH gradient between intracellular and extracellular space have also been neglected in predicting V_{ss} . In fact, each tissue has been considered as a mixture of lipids, water, and plasma proteins having a global pH of 7.4. Only the hydrophobic interactions between drugs and tissue lipids have currently been considered for predicting V_{ss} . Consequently, the potential ionic interactions with charged lipids have been neglected. Regarding the predicted and experimental V_{ss} 's provided in Table 2, the present description of tissues is acceptable for several kinds of drugs. However, this appears not to be the case for drugs in Table 3, for which inaccurate predictions of their V_{ss} 's are observed. In fact, an under-prediction of V_{ss} by a factor of at least three is especially observed for some cationic-amphiphilic compounds (amiodarone, amlodipine, biperiden, chlorpromazine, exaprolol, haloperidol, imipramine; $\log P_{o:w} > 3$, $pK_a > 8.6$). This may be explained by the fact that ionic interactions between drugs and charged tissue lipids were neglected in predicting their V_{ss} as compared to hydrophobic interactions. As an example, amiodarone has an *in vivo* V_{ss} value three times larger than the one predicted (Table 3). This drug is extremely lipophilic, amphiphilic, and cationic, properties for which anomalous pharmacokinetic behaviors are reported.¹⁶ Its $B:P$ ratio is independent of drug

concentration, which is not the case for V_{ss} , suggesting that distribution into peripheral tissues accounts for the concentration-dependence of V_{ss} .¹⁶ Therefore, it may be speculated that some lipid-rich inclusions of tissues (e.g., cells) act as deep traps considering the high membrane:buffer ratio of amiodarone probably caused by the presence of hydrophobic and ionic interactions.

For other cationic-amphiphilic bases [e.g., alprenolol, carvedilol, chlorpheniramine, fentanyl, lorcaïnide, mefloquine (S^+), methadone, pentazocine, phencyclidine, propafenone, propranolol, quinidine, verapamil; $\log P_{o:w} > 3$, $pK_a > 8.4$], V_{ss} can be more reasonably predicted (Table 2). This is in clear contrast with other bases of Table 3, for which V_{ss} 's are largely under-predicted. It is then suspected that the ionic interactions with charged lipids of cell membranes (a nonspecific binding) are less relevant for bases of Table 2 compared to those of Table 3. This has been observed from preliminary *in vitro* studies (not shown). Therefore, this is now under investigation. If significant saturable nonspecific ionic binding is present in cell membranes of tissues for specific bases, V_{ss} can therefore be dependent on drug concentration, which can be an obvious matter of concern in predicting V_{ss} as a function of dose.

A presence of lysosomal ionic trapping in certain tissues (e.g., muscle, liver, lung)^{6,17} may also be a potential source of explanation for the large under-prediction of V_{ss} in the case of cationic-amphiphilic bases of Table 3. The most pronounced effect of the lysosomal ionic trapping may represent 20 to 40% of the total tissue accumulation of a base such as thioridazine under *in vitro* conditions.²⁰⁶ However, the lysosomes (where the pH is 4.7) represent only 1% of the total tissue volume under *in vivo* conditions.⁶⁵ Nevertheless, a significant lysosomal ionic trapping may potentially affect the first passage of a base through the lung, a tissue rich in lysosomes, particularly after intravenous bolus administrations. This may result in a slow release into the plasma leaving this tissue. In other words, the estimation of V_{ss} *in vivo* from data on plasma concentration-time profiles might be biased. Furthermore, a problem of drug solubility in the lung following the intravenous administration of an inadequate formulation may also result in a slow release into the plasma leaving this tissue. At present, however, mechanistic studies are still needed to explain why V_{ss} of cationic-amphiphilic bases can or cannot be accurately

predicted with the present method (i.e., Table 2 vs. Table 3).

Enterohepatic recirculation and/or active uptake processes may lead to a contribution of the intestinal lumen to the distribution space. This would be significant particularly if the active uptake processes become important compared to the passive diffusion process. In this case, the *in vivo* V_{ss} would be under-predicted by the present approach, which does not consider any accumulation into the intestinal lumen. It is observed that V_{ss} is under-predicted for ciprofloxacin, enoxacin, and trimethoprim (Table 3), but is more reasonably predicted for cyclosporin, propranolol, and verapamil (Table 2). It is known that these drugs may accumulate into the intestinal lumen to different degrees. In another context, the prediction of V_{ss} is species variant especially for three drugs. For thiopental, its predicted and experimental V_{ss} 's differ by a factor of 3.27 and 1.44 in rat and human, respectively. For digoxin and alfentanil the predicted and experimental V_{ss} differ by a factor of 0.61 and 2 in the rat, respectively. In the human, the factor is 0.13 and 6.1, respectively (Tables 2 and 3). Definitive explanations cannot be provided. However, the disposition of digoxin is governed either by active transport, biliary clearance, and/or Na/K ATPase activity,^{18,19,207} which may affect rat and human V_{ss} to different degrees.

In Figures 2 and 3 tridimensional mesh and contour plots are presented for the relationships between the calculated V_{ss} and the compound-specific determinants fu_p (ranging from 0 to 1) and $\log P_{o:w}$ (ranging from -2 to 8). These theoretical exercises are mechanism-based relationships between potential determinants of rat and human V_{ss} . For a hypothetical monoprotic base of pK_a equal to 3 or 9, sigmoid relationships between V_{ss} , $\log P_{o:w}$, and fu_p are observed. Similar types of relationships were also observed for monoprotic acids and neutral compounds (not shown). The resulting relationships demonstrate that V_{ss} of a drug is enhanced by increasing its lipophilicity, but is reduced by increasing its plasma protein binding. The relevance of ionization on V_{ss} becomes apparent when comparing a base with pK_a of 3 or 9. In other words, the upper profile of the tridimensional plots observed in Figures 2 and 3 is not exactly similar between a base of pK_a equal to 3 or 9 (Figures 2A1 and 3A1 vs. Figures 2A2 and 3A2). The rationale for that is the contribution of $P_{t,p}$ of adipose tissue in V_{ss} . In fact, $P_{t,p}$ of adipose tissue calculated with eq. 3 is

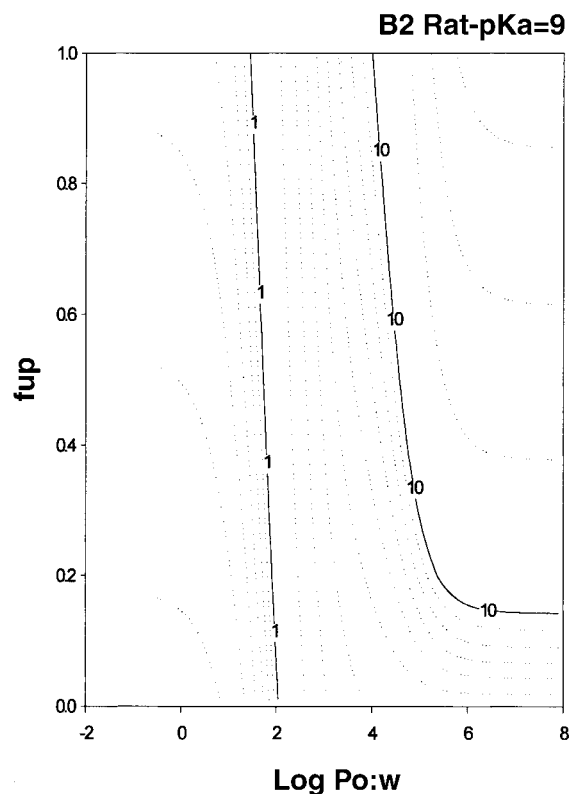
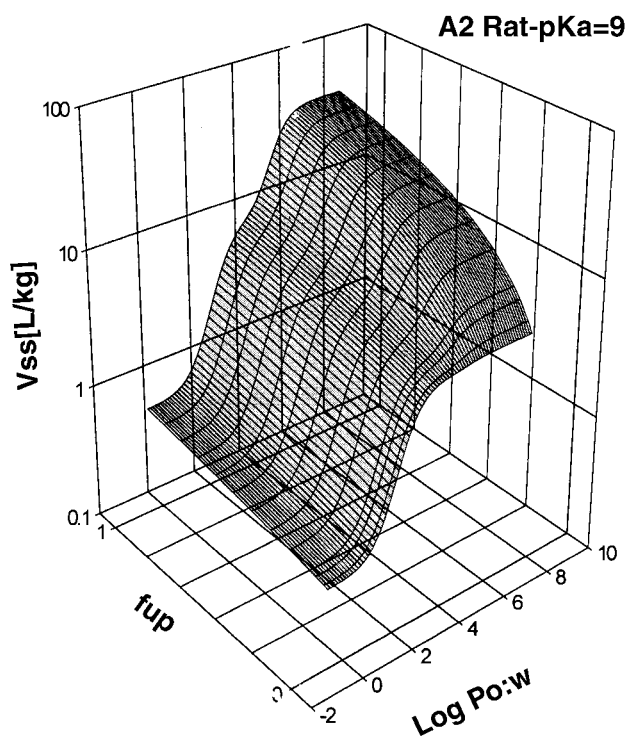
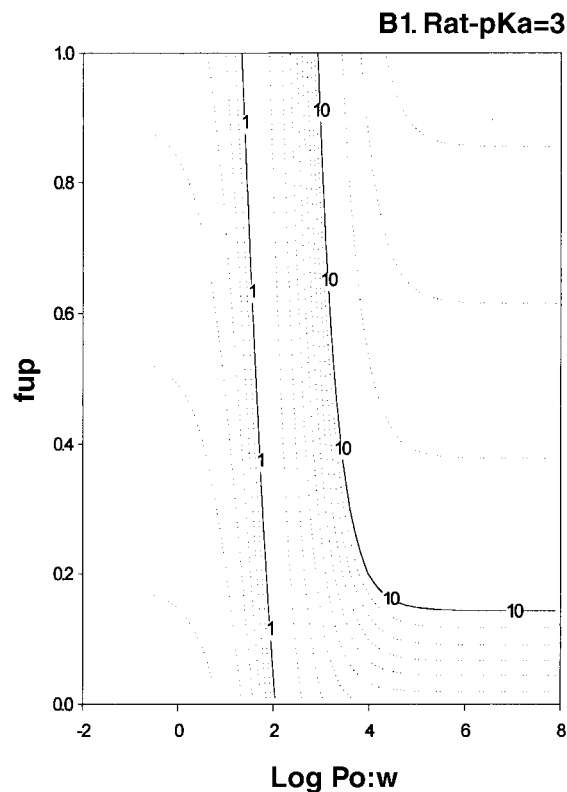
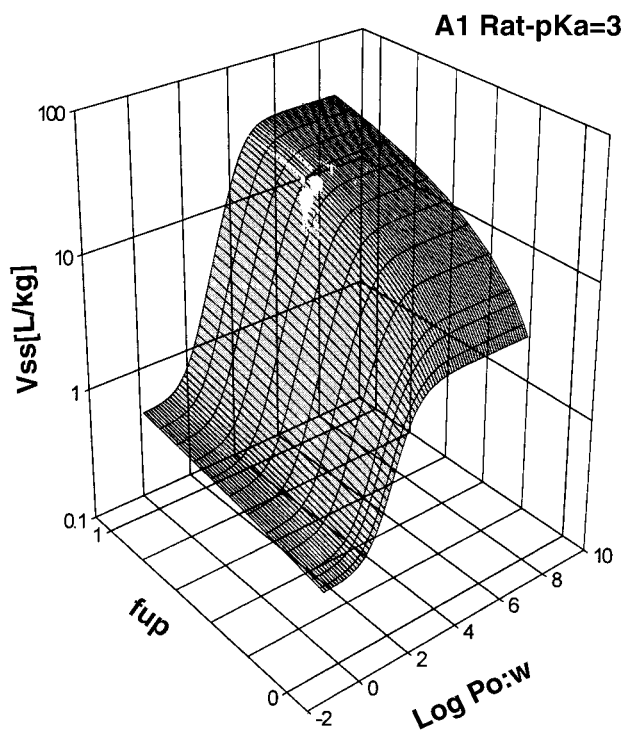


Figure 2. Tridimensional mesh (A) and contour (B) plots for the relationships between f_{up} , $\log P_{o:w}$, and rat V_{ss} calculated. The plots were performed for a hypothetical monoprotic base of pK_a equal to 3 (A1, B1) or 9 (A2, B2). The rat V_{ss} was calculated with eqs. 1–3 as demonstrated in the Methods section by using theoretical data on f_{up} and $\log P_{o:w}$ ranging from 0 to 1 and -2 to 8, respectively. A value of blood:plasma ratio of 1 was used. The pK_a was used to calculate $D_{vo:w}^*$ of eq. 3 from data on $P_{o:w}$. A homogeneous tissue distribution was assumed. Sigmaplot[®] version 4.0 was used for plotting.

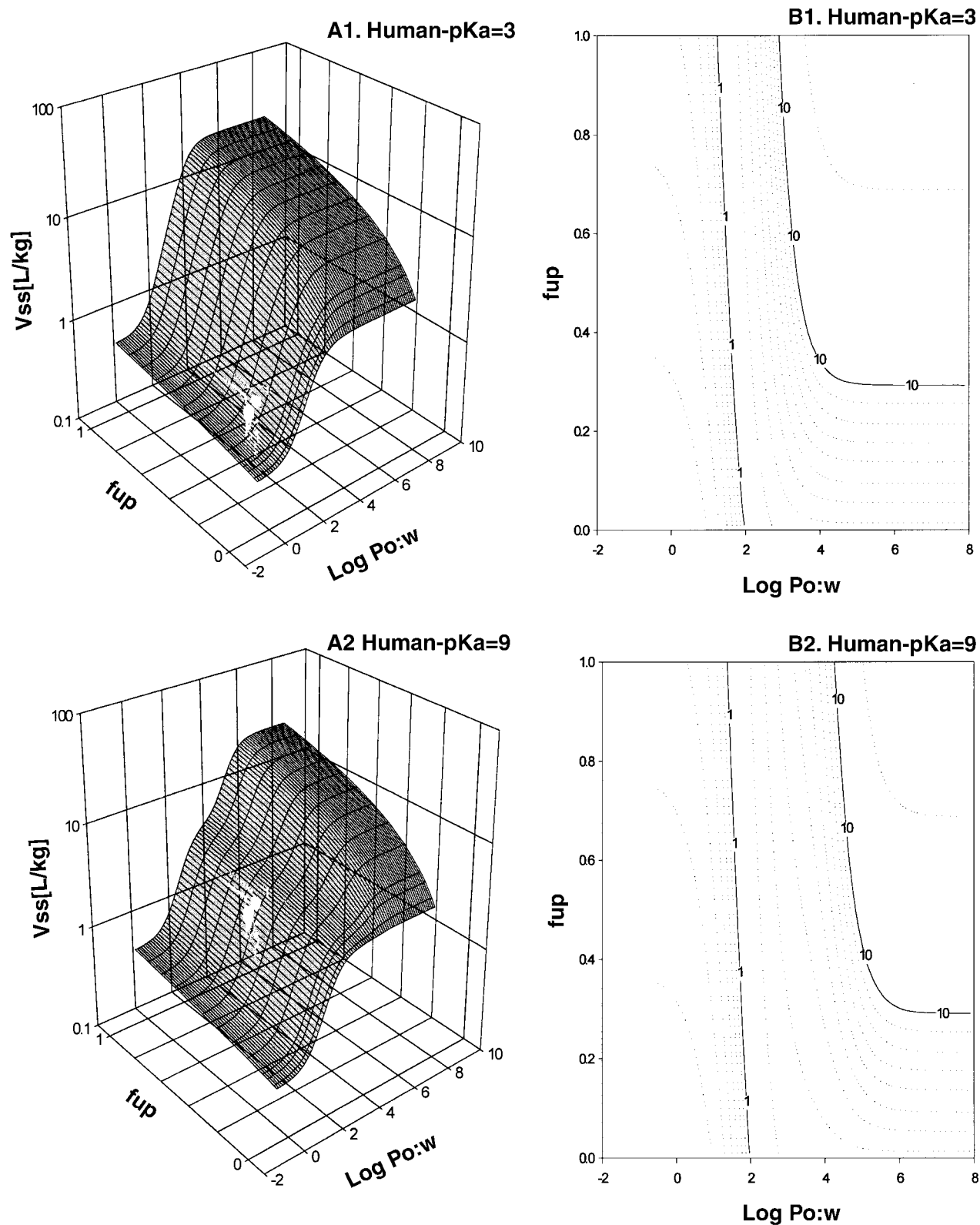


Figure 3. Tridimensional mesh (A) and contour (B) plots for the relationships between $f u_p$, $\log P_{o:w}$, and human V_{ss} calculated. The plots were performed for a hypothetical monoprotic base of pK_a equal to 3 (A1, B1) or 9 (A2, B2). The human V_{ss} was calculated with eqs. 1–3 as demonstrated in the Methods section by using theoretical data on $f u_p$ and $\log P_{o:w}$ ranging from 0 to 1 and -2 to 8, respectively. A value of blood:plasma ratio of 1 was used. The pK_a was used to calculate $D_{vo:w}^*$ of eq. 3 from data on $P_{o:w}$. A homogeneous tissue distribution was assumed. Sigmaplot[®] version 4.0 was used for plotting.

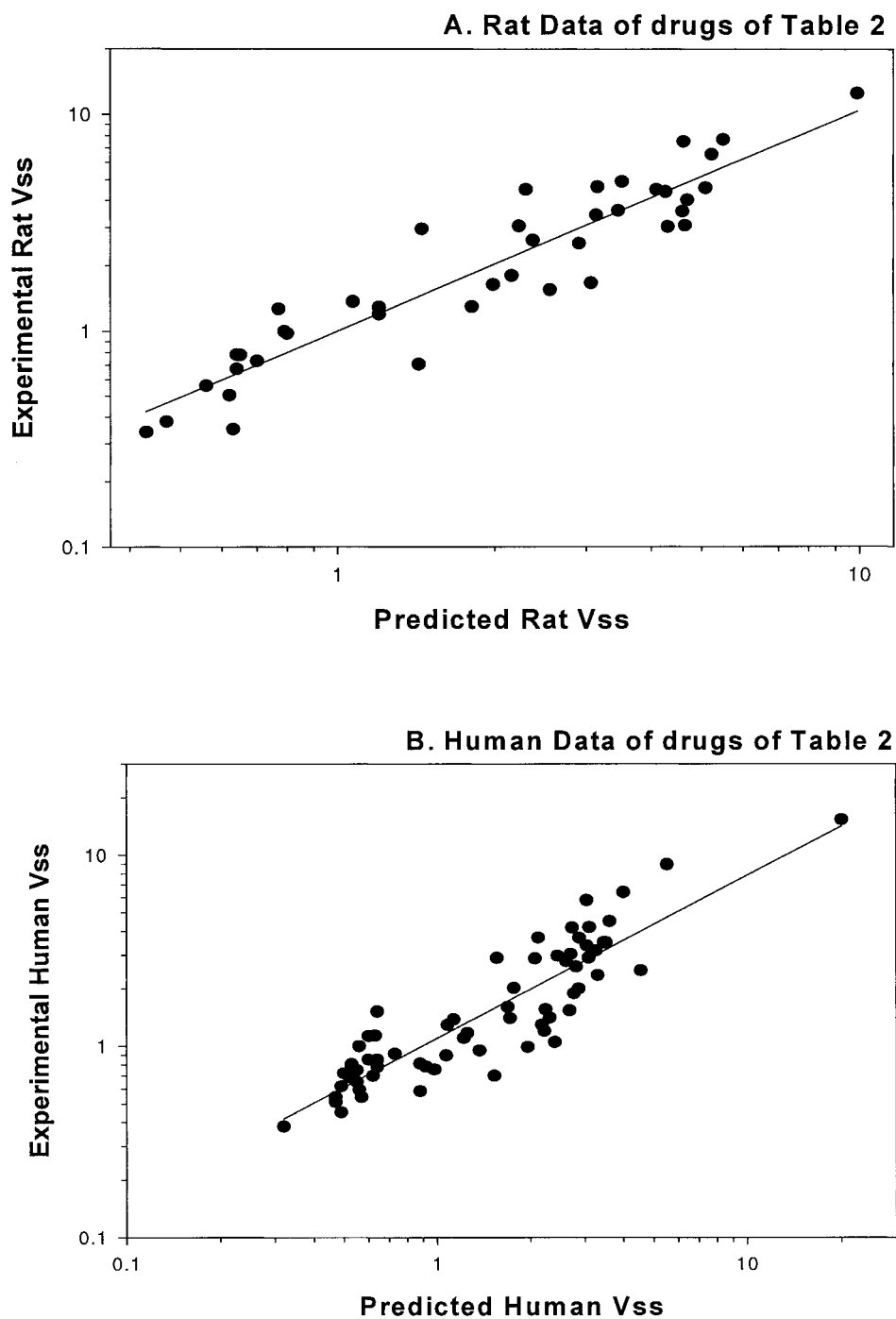


Figure 4. Relationships between the predicted and experimentally determined V_{ss} of rats (A) and humans (B) for the 89 drugs listed in Table 2. The line indicates the best fit for (A) [$\log y = 1.01895 \cdot \log x + 0.00101$, $r = 0.93$, $n = 41$, average ratio predicted-to-experimental $V_{ss} = 1.03 \pm 0.37$] and (B) [$\log y = 0.85347 \cdot \log x + 0.04118$, $r = 0.89$, $n = 68$, average ratio predicted-to-experimental $V_{ss} = 1.03 \pm 0.42$]. Sigmaplot[®] version 4.0 was used for plotting.

reduced when the input parameter $D_{vo:w}^*$ decreases by increasing pK_a .

Potential interspecies differences in V_{ss} are demonstrated in Figures 2 and 3. On a theoretical basis, the interspecies differences in V_{ss} between rat and human are differences in tissue lipid composition and f_{up} for similar drug lipophilicity. Therefore, the minimal theoretical value of V_{ss} that can be calculated from eq. 1 is similar for rat and human (≈ 0.31), which it is not the case for the maximal value that is higher in rat than in human (≈ 46 vs. 28; Figures 2 and 3). For drugs listed in Table 2, the regression analyses comparing the calculated and experimental V_{ss} are similar for both species (Figure 4).

Eq. 1 can predict V_{ss} 's of drugs distributed homogeneously or nonhomogeneously (i.e., extracellular space only) into tissues. However, approaches are still needed to identify by which of these two scenarios a novel drug candidate will be distributed into tissues. An *in vitro* technique using tissue slices can potentially be used.²⁰⁸ The drugs residing in the extracellular space are mainly acids that are highly hydrophilic or lipophilic as well as highly bound compounds (Table 4), properties that may reduce cell permeability. Eq. 1 calculates a maximal value of 0.31 L/kg for rat V_{ss} of drugs residing mainly in extracellular space, which corresponds to the minimal calculated value of V_{ss} of drugs distributing homogeneously into tissues. This calculated value of 0.31 is similar to the extracellular water space in the rat body.²⁰⁵

In conclusion, the present work suggests that V_{ss} *in vivo* of rat and human can be reasonably predicted prior to *in vivo* studies for most of the drugs investigated. However, the predicted and experimental V_{ss} 's may deviate by a factor larger than two (under- or over-predictions) for other drugs such as some cationic-amphiphilic bases, for which the V_{ss} 's are largely under-predicted. Additional relevant processes may have been neglected in the tissue composition-based equations explaining the less accurate predictions. These processes may refer to ionic interactions with charged lipids of cell membranes, specific macromolecular binding to cell components, active transport processes, or limitation for membrane permeation. These processes need to be described in tissue composition-based equations for a broader applicability. It is essential to adapt eq. 1 to predict V_{ss} according to age, sex, and disease state. Changing data on tissue volume and composition, as well as on f_{up} , may do this.

The present study is the first attempt to develop and validate a mechanistic distribution model for predicting rat and human V_{ss} *in vivo* of structurally unrelated drugs. Predictions of V_{ss} were made solely from compound-specific data determined *in vitro* ($P_{o:w}$, pK_a , f_{up}), as well as tissue composition data obtained from the literature. Thus, the present study may open opportunities for a reduction of animal experiments.

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APPENDIX

Estimation of Tissue Volume and Composition of Standard Adult Male Rats (250 g) and Humans (70 kg)

Tissue Volume Data (V_t) Reported in Table 1 for Adult Male Rats and Humans

The compilation of Brown et al.¹⁹⁰ reports the mean values of tissue volume (V) as a fraction of total body weight (L/kg). These authors provide data for adipose tissue (subcutaneous), skeleton + bone marrows, brain, gut, heart, kidney, liver, lung, muscle, skin, and spleen. In converting the values of tissue volume in kilograms to liters, the density was considered for adipose tissue (0.92 kg/L for rat and human) and skeleton + bone marrows (1.76 kg/L for rat and 1.67 kg/L for human).¹⁹⁰ For all the other tissues a density close to 1 was reported¹⁹⁰ or assumed. The overall density for skeleton + bone marrows was recalculated by multiplying the density of the main bone components (cortical, trabular, yellow, and red marrows) with their respective fractional volume of total bone.¹⁹⁰

The arterial and venous blood volumes expressed as a fraction of total body weight (L/kg) were obtained from Igari et al.¹⁹¹ in absence of these data in the compilation of Brown et al.¹⁹⁰ In calculating the volume of plasma and red blood cells, the rat and human blood were assumed to be constituted of 55% of plasma and 45% of erythrocytes (hematocrit) as classical values.

Tissue Composition Data (V_w , V_{nl} , V_{ph}) Reported in Table 1 for Adult Male Rats and Humans

The neutral lipids (nl) refer to the sum of triglycerides, diglycerides, monoglycerides, cholesterol,

and other types of nonpolar lipids. Phospholipids (ph) represent lipids that contain phosphoric acid esterified at one position of the glycerol molecule (e.g., phosphatidylcholine, phosphatidyl ethanolamine, phosphatidylserine, sphingomyelin).

The fractional volume content of total neutral lipids (V_{nl}) in each tissue was obtained from the difference between fractional volume content of total lipids and fractional volume content of total phospholipids (V_{ph}). V_{nl} and V_{ph} are reported on the basis of wet tissue weight. However, V_{nl} and V_{ph} reported on the basis of dry tissue weight were converted on the basis of wet tissue weight as follows: V_{nl} (V_{ph}) for wet tissue weight = V_{nl} (V_{ph}) for dry tissue weight $\times (1 - V_w)$, where V_w is the fractional volume content of water. The mean data on V_w in rat and human tissues were obtained from the compilation of Reinoso et al.,¹⁹² whereas data on V_{nl} and V_{ph} in each rat and human tissue were obtained in the references listed as follows.

Here are the references for V_{nl} and V_{ph} of each tissue: adipose tissue (subcutaneous white fat) of rat and human¹⁹³; bone + marrows of rat¹⁹⁴ and human¹⁹⁵; whole brain of rat and human¹⁹⁶; gut/small intestine of rat¹⁹⁷ and human¹⁹⁸; heart of rat and human¹⁹⁹; whole kidney of rat and human²⁰⁰; liver of rat²⁰⁰ and human¹⁹⁸; lung of rat and human²⁰¹; skeletal muscle of rat¹⁹⁹ and human²⁰²; skin epidermis of rat¹⁹⁷ and human²⁰³; spleen of rat and human²⁰⁰; plasma of rat¹⁹⁸ and human.²⁰⁴

In the literature cited above, the estimation of V_{nl} and V_{ph} was performed as detailed below. It was assumed that cortical and trabecular components of bone contain only traces of lipids.¹⁹⁸ Therefore, V_{nl} and V_{ph} for the overall bone (skeleton) were recalculated by multiplying V_{nl} and V_{ph} of the marrows (yellow and red)^{194,195} with their respective fractional volume content in total bone.¹⁹⁰ For the whole brain, V_{nl} and V_{ph} were recalculated from data on V_{nl} and V_{ph} determined for the gray and white matters¹⁹⁶ by assuming that each of these two components constitutes 50% of the brain. In calculating V_{nl} and V_{ph} for gut, heart, kidney, liver, lung, muscle, and spleen, a conventional factor of 25 was used to convert the level of phospholipid expressed in milligrams phosphorus to milligrams phospholipids.^{198–201} Data on V_{nl} and V_{ph} for gut (small intestine) and skin (epidermis) represent those for a neonatal mouse¹⁹⁷ in the absence of data for adult rats. These data, however, were available in human adults.^{203,204} The composition of human skin

epidermis taken from reference 203 corresponds to the average values of V_{nl} and V_{ph} that have been determined with pieces of summer and winter skin of subjects of 25 to 35 years.

In references 199 and 200 data on fractional volume content of total lipid in muscle and liver were reported in both rat and human. However, these data for muscle (10%) and liver (14%) reported for an individual human appear elevated compared to other literature sources. This may be explained by the fact that the sample of human skeletal muscle used for the determination of total lipids may contain subcutaneous fat, whereas the sample of human liver may correspond to a sick and/or an old individual (but no information is provided). Therefore, values of V_{nl} and V_{ph} in the human skeletal muscle (psoas) and liver were calculated from other references, 198 and 202, respectively.

The values concerning the fractional volume content of extracellular (interstitial) space (V_{eis}) in all rat blood-free tissues were also obtained from the literature.²⁰⁵

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