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Exploratory Translational Modeling Approach in Drug Development to Predict Human Brain
Pharmacokinetics and Pharmacologically Relevant Clinical Doses

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Running Title: Predicting Human Brain Pharmacokinetics

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d) Nonstandard abbreviations: CNS, central nervous system; BBB, blood brain barrier; BCB, blood cerebrospinal fluid barrier; bECF, brain extracellular fluid; NE, norepinephrine; DHPG, 3,4-dihydroxyphenylglycol; CSF, cerebrospinal fluid; PK, pharmacokinetic(s); PD, pharmacodynamic(s); hNET K_i , human norepinephrine transporter inhibition constant; hSERT K_i , human serotonin transporter inhibition constant.

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Abstract

The central nervous system (CNS) pharmacokinetics (PK) of drugs that have pharmacological targets in the brain are not often understood during drug development and this gap in knowledge is a limitation in providing a quantitative framework for translating nonclinical pharmacologic data to the clinical patient population. A focus of translational sciences is to improve the efficiency of clinical trial design via a more judicious selection of clinical doses based on nonclinical data. We hypothesize that this can be achieved for CNS-acting drugs based on knowledge of CNS PK and brain target engagement obtained in nonclinical studies. Translating CNS PK models from rat-to-human can allow for the prediction of human brain PK and the human dose – brain exposure relationship, which can provide insight on the clinical dose(s) having potential brain activity and target engagement. In this manuscript, we explored the potential utility of this translational approach using rat brain microdialysis and PK modeling techniques to predict human brain extracellular fluid PK of atomoxetine and duloxetine. The results show that this translational approach merits consideration as a means to support the clinical development of CNS-mediated drug candidates by enhancing the ability to predict pharmacologically relevant doses in humans in the absence of or in association with other biomarker approaches.

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Introduction

Drug development archetypes for transitioning central nervous system (CNS) molecules into clinical testing involve identification of target brain activity in nonclinical pharmacology models, but often the pharmacokinetics (PK) of such drugs in the brain are not understood. The lack of knowledge of brain PK, which influences the centrally mediated effect at the target, is a limitation in providing a quantitative framework for translating nonclinical pharmacologic data to the clinical patient population (Westerhout et al., 2011). Owing to the effectiveness of the blood-brain barrier (BBB) and blood-cerebrospinal fluid barrier (BCB) in limiting access of drugs to the CNS, the use of plasma concentrations to predict brain concentrations for drugs can be misleading. Furthermore, nonhomogeneous drug distribution within the brain, as a result of drug binding to brain tissue and the presence of influx and efflux carrier-mediated transport across the BBB and BCB and brain cell membranes, complicates the prediction of drug concentrations in the brain extracellular fluid (bECF), the site of action for CNS drugs whose targets are in the extracellular space. In order to address these complexities of CNS drug development, approaches such as positron emission tomography (PET) or single photon emission computed tomography (SPECT) imaging to measure receptor occupancy (Tauscher and Kapur, 2001) and measurement of drug concentrations in cerebrospinal fluid (CSF) (de Lange and Danhof, 2002; Shen et al., 2004; Lin, 2008) have been used; however, limitations exist preventing consistent application of these approaches. Notable among these limitations is the lack of a suitable tracer ligand to conduct imaging studies and the technical complexities associated with CSF sampling in humans. Application of biomarker approaches that measure CNS target engagement, which are based on proteomics or metabolomics, have received much

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attention; however, for novel targets, development of a suitable biomarker often proceeds in parallel with drug development, thus lessening the degree to which the biomarker can impact development.

Pharmacokinetic modeling plays an essential role in the process of drug development, yet cogent arguments have been made regarding the need to be more intentional about expanding its role to one of a continuous model-based approach applied across the drug development continuum to increase the efficiency and effectiveness of drug development (Zhang et al., 2008).

Incorporation of pharmacokinetic-pharmacodynamic (PK-PD) principles into animal models of CNS drug action has been advocated as a comprehensive approach to guide the selection of clinical candidates with a high probability of target engagement (de Lange et al., 2005). An additional goal of this translational-sciences approach is to improve the efficiency of clinical trial design through a more rational selection of doses based on knowledge of the relationship between dose, PK and target engagement obtained in nonclinical studies. We are interested in the application of quantitative brain microdialysis as a translational tool to support clinical development of CNS drugs. Brain microdialysis in rats can be used to measure drug concentrations in bECF and these data can support the development of brain PK models that quantitatively relate effect-site concentrations to plasma concentrations. Translating preclinical brain PK models to humans using allometric principles allows for the prediction of human brain PK and the human dose-brain exposure relationship, which adds perspective on potential brain activity and target engagement when used in context with information on drug-receptor potency. To explore the potential utility of this translational approach, we evaluated duloxetine and atomoxetine because the targets for these molecules are in the brain (centrally-acting), and the human plasma PK and clinical effective doses are known. Atomoxetine [(-)-N-methyl-3-phenyl-

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3-(o-tolyloxy)-propylamine] is a selective norepinephrine transporter inhibitor (Bymaster et al., 2002) approved as a therapeutic agent for the treatment of attention deficit/hyperactivity disorder (ADHD). Duloxetine [(+)-(S)-N-Methyl-3-(naphthalen-1-yloxy)-3-(thiophen-2-yl)propan-1-amine] is an inhibitor of both serotonin and norepinephrine reuptake transporters (Pitsikas, 2000; Wong, 1998) approved for the treatment of major depressive disorder, generalized anxiety disorder, fibromyalgia, chronic musculoskeletal pain and diabetic neuropathy.

We previously reported on a CNS PK model of atomoxetine in rats using brain microdialysis (Kielbasa et al., 2009). In this manuscript, we applied a rat-to-human translational approach to predict human bECF PK of atomoxetine and duloxetine from microdialysis studies and CNS PK models developed from rats. The investigational approaches, results and applications are discussed herein.

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Methods

Methods for atomoxetine were described previously (Kielbasa et al., 2009). Briefly, methods for duloxetine experiments are described below.

Drugs and Chemicals

Duloxetine HCl and $^{13}\text{CD}_3$ -duloxetine, the retrodialysis calibrator for PK experiments and stable label of duloxetine, were synthesized at Eli Lilly and Company (Indianapolis, IN). The perfusion solution consisted of microdialysis buffer (Kielbasa et al., 2009) with bovine serum albumin (BSA, 0.5%) and 100 ng/ml $^{13}\text{CD}_3$ -duloxetine. The intravenous (iv) dosing solution consisted of 0.5 mg/ml duloxetine in 5% dextrose in water.

Animal Preparation

Male Sprague-Dawley rats weighing between 275 and 325 g were surgically prepared at Charles River Laboratories, Inc. (Wilmington, MA) to support iv administration and arterial blood collection via femoral vein and artery cannulation, respectively. Under isoflurane anesthesia, a guide cannula (BAS, Bioanalytical Systems, West Lafayette, IN) was implanted in the medial-prefrontal cortex (coordinates: anteroposterior, + 3.2 mm; mediolateral, + 0.6 mm; and dorsoventral, +2.2 mm) and rats were allowed to recover from surgery for 3 days. One day before an experiment, a CMA-12 probe 3 mm in length (CMA Microdialysis AB, Solna, Sweden) was inserted into the guide cannula under isoflurane anesthesia and subsequently perfused with dialysis buffer for 18 to 24 hours (h) at a flow rate of 0.2 $\mu\text{l}/\text{min}$. All activities were performed humanely under a protocol approved by the Eli Lilly and Company Institutional

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Animal Care and Use Committee in an animal facility accredited by the Association for Assessment and Accreditation of Laboratory Animal Care, International.

Evaluation of Duloxetine Plasma and Brain Binding

The duloxetine unbound plasma fraction (f_{uP}) and unbound brain fraction (f_{uB}) were determined in a 96-well equilibrium dialysis apparatus (HTDialysis, Gales Ferry, CT) using a method reported previously (Kalvass and Maurer, 2002).

Sample Collection, Preparation, and Analysis

The live phase portion of the experiment was conducted as described previously (Kielbasa et al., 2009). Briefly, a Culex Automated Pharmacology System (BAS, Bioanalytical Systems, West Lafayette, IN) was used that enabled timed blood and dialysate collections in awake and freely moving animals. Blood was collected in disodium ethylenediaminetetraacetic acid (EDTA) tubes and centrifuged to generate plasma within 30 min of collection. At the conclusion of the live phase animals were euthanized by CO₂ asphyxiation and whole brain and CSF from the cisterna magnum were collected from each animal. Duloxetine containing blood was removed from brain via carotid arterial perfusion with cold saline prior to collection. Plasma, dialysate and CSF samples were stored in a 96-well plate at -70°C before bioanalysis. Whole brains were homogenized in acetonitrile at a ratio of 2 ml/g of brain and centrifuged at 3000 rpm for 10 min at ambient temperature. Supernatant (50 µl) was placed into a 96-well plate for analysis. All matrices were analyzed by liquid chromatography tandem mass spectrometry (LC/MS/MS) using a Shimadzu high performance liquid chromatograph and a Sciex API 4000 mass spectrometer. Duloxetine and ¹³CD₃-duloxetine concentrations were determined against standard curves prepared in the corresponding matrix (perfusion buffer for CSF). The major metabolite in

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humans, 4-hydroxy-duloxetine, was used as an internal standard. The lower limit of quantitation of duloxetine and $^{13}\text{CD}_3$ -duloxetine was 1 ng/ml.

Duloxetine Pharmacokinetic Study Design

The perfusion solution (containing $^{13}\text{CD}_3$ -duloxetine) was initiated through the microdialysis probe at 1 $\mu\text{L}/\text{min}$ for 0.5 hours prior to and during the 8 h iv administration of duloxetine. An iv loading dose of 3 mg/kg duloxetine was given over 0.5 h followed by a 4.1 mg/hr/kg infusion for 7.5 hours to achieve a targeted duloxetine plasma concentration of 600 ng/ml. Microdialysis samples (30 μL) were collected every 0.5 h beginning from the start of the loading dose. At 0.25 h, blood-sample (0.15 ml) collections were initiated and taken every 0.5 h thereafter. At 8 h, the iv infusion and the perfusion solution were stopped and animals were euthanized by CO_2 asphyxiation.

Rat Pharmacokinetic Analysis of Duloxetine.

The loss (L) of $^{13}\text{CD}_3$ -duloxetine across the microdialysis probe at each dialysate collection time was calculated using the equation,

$$L = \frac{C_{in} - C_{out}}{C_{in}} \quad (1)$$

where C_{in} is the concentration of $^{13}\text{CD}_3$ -duloxetine in the perfusate (100 ng/ml) and C_{out} is the concentration of $^{13}\text{CD}_3$ -duloxetine in the dialysate. At each dialysate collection period, the duloxetine bECF concentration was calculated using the equation,

$$C_{ECF} = \frac{C_{Dt}}{R} \quad (2)$$

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where, C_{Dt} is the measured concentration of duloxetine in the dialysate at time t and R is the recovery of duloxetine, which is assumed to be equal to the loss of the retrodialysis calibrator, $^{13}\text{CD}_3$ -duloxetine. Prior to PK experiments, separate in vitro and in vivo experiments were conducted to confirm similar loss of duloxetine and $^{13}\text{CD}_3$ -duloxetine across the microdialysis probe (data not shown).

The duloxetine concentration associated with brain cells (C_{BC}) was calculated using the following equation,

$$C_{BC} = \frac{C_B \times V_B - C_{ECF} \times V_{ECF}}{V_{BC}} \quad (3)$$

where C_B is the whole brain concentration following duloxetine infusion, C_{ECF} is the brain extracellular fluid concentration, V_{ECF} is the brain extracellular fluid volume, V_{BC} is the brain cell volume and V_B is the total brain volume. Values for V_{ECF} and V_{BC} were assumed to be 0.00029 L and 0.00099 L, respectively (Mahar Doan and Boje, 2000). The sum of V_{ECF} and V_{BC} was assumed to be equal to V_B (Scism et al., 2000).

The unbound duloxetine plasma concentration (C_{uP}) was determined by multiplying the measured duloxetine plasma concentration (C_P) by f_{uP} . Duloxetine unbound brain cell concentration (C_{uBC}) was determined by multiplying C_{BC} by f_{uB} . The duloxetine CSF concentration (C_{CSF}) and C_{ECF} were considered to be unbound as duloxetine binding to proteins in CSF and bECF was assumed to be negligible. The C_P , C_{uP} and C_{ECF} from each rat were determined by averaging the data collected from 4 h to 8 h after duloxetine administration, and the C_{CSF} and C_B were determined at 8 hour after duloxetine administration. The following ratios were calculated in each rat when applicable: C_B/C_P , C_B/C_{ECF} , C_{BC}/C_{ECF} , C_{uBC}/C_{ECF} , C_{ECF}/C_{uP} and C_{CSF}/C_{uP} .

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Rat PK Model Development of Duloxetine

Pharmacokinetic modeling was performed using NONMEM software (Version V, level 1.1; GloboMax LLC, Hanover, MD). The PK model (Figure I) consisted of a systemic compartment and a CNS compartment, the latter of which was divided into CSF, bECF and brain cell compartments, described using differential equations as follows:

$$\frac{dA_{uP}}{dt} = -\frac{CL_u}{V_u} f_{uP}A_P - \frac{Q_{BCB}}{V_u} f_{uP}A_P - \frac{Q_{BBB}}{V_u} f_{uP}A_P + \frac{Q_{BCB}}{V_{CSF}} A_{CSF} + \frac{Q_{BBB}}{V_{ECF}} A_{ECF} \quad (4)$$

$$\frac{dA_{CSF}}{dt} = -\frac{Q_{BCB}}{V_{CSF}} A_{CSF} + \frac{Q_{BCB}}{V_u} f_{uP}A_P + \frac{CL_{ECF-CSF}}{V_{ECF}} A_{ECF} \quad (5)$$

$$\frac{dA_{ECF}}{dt} = -\frac{Q_{BBB}}{V_{ECF}} A_{ECF} - \frac{CL_{ECF-BC}}{V_{ECF}} A_{ECF} - \frac{CL_{ECF-CSF}}{V_{ECF}} A_{ECF} + \frac{Q_{BBB}}{V_u} f_{uP}A_P + \frac{CL_{BC-ECF}}{V_{BC}} f_{uB}A_{BC} \quad (6)$$

$$\frac{dA_{uBC}}{dt} = -\frac{CL_{BC-ECF}}{V_{BC}} f_{uB}A_{BC} + \frac{CL_{ECF-BC}}{V_{ECF}} A_{ECF} \quad (7)$$

The unbound duloxetine plasma concentrations were used for modeling; therefore, amounts, clearances and volume parameters are described as unbound terms. Time was represented as t , A_{uP} was the amount in plasma, A_{CSF} was the amount in CSF, A_{ECF} was the amount in ECF, A_{uBC} was the amount associated with the brain cell, CL_u was the plasma clearance, Q_{BCB} was the distributional clearance at the BCB, Q_{BBB} was the distributional clearance at the BBB, $CL_{ECF-CSF}$ was the clearance from the ECF to the CSF, CL_{ECF-BC} was the clearance from the ECF to the brain cell, CL_{BC-ECF} was the clearance from the brain cell to the ECF and V_u was the plasma volume of distribution. Model terms V_{BC} (0.00099 L), V_{CSF} (0.00025 L) and V_{ECF} (0.00029 L) represented the volume of the brain cell; CSF and ECF, respectively, and were fixed in the PK model (Mahar Doan and Boje, 2000; Segal 1993; Shen et al., 2004).

The model incorporated the unidirectional transport from ECF to CSF ($CL_{ECF-CSF}$). In rats, the first-order rate constant for flow from ECF to CSF ($k_{ECF-CSF}$) was estimated to be 0.084 h^{-1}

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(Szentistványi et al., 1984). A calculated value for $CL_{ECF-CSF}$ is 0.000024 L/h ($CL_{ECF-CSF} = k_{ECF-CSF} \times V_{ECF}$). During model development, $CL_{ECF-CSF}$ was either fixed to the calculated value or estimated. Fixing $CL_{ECF-CSF}$ provided a better fit of the data to the model (data not shown).

Inter-individual variability in PK parameters were included in the model as described by the following equation,

$$P_{ij} = P_j \times \exp(\eta_{ij}) \quad (8)$$

where P_{ij} is the j^{th} parameter for the i^{th} individual, P_j is the typical population parameter estimate for the j^{th} parameter and η_{ij} is the deviation of P_{ij} from P_j in the j^{th} parameter for the i^{th} individual. For η_{ij} , it is assumed that the parameter is normally distributed with a mean zero and a variance (ω) to be estimated. Residual error was estimated using a proportional error model as described by the following equation,

$$C_{ik} = \text{Pred}_{ik} \times (1 + \sigma_{\text{prop}}), \quad (9)$$

where C_{ik} and Pred_{ik} are the measured and model-predicted concentration at the k^{th} sampling time in the i^{th} individual, respectively. The residual error, σ , is a random variable normally distributed with mean zero and estimated variance σ^2 . The residual error describes errors arising from assay errors, sampling inaccuracies, and model misspecification.

The modeling data set consisted of 207, 166, 8 and 13 data points for unbound plasma, bECF, CSF and unbound brain cell concentrations, respectively, from 13 rats. The criteria used for the model evaluation were the fit between observed and predicted concentrations, the parameter's percentage standard error of the estimate (% SEE), the randomization of weighted residual

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concentrations versus time between observed and predicted concentrations and the objective function.

Prediction of Human Atomoxetine and Duloxetine bECF Pharmacokinetics

Simulations were conducted (NONMEM software Version VII, level 1.2; GloboMax LLC, Hanover, MD) in a population consisting of 1000 humans to generate plasma and bECF PK profiles following oral administration of 40-mg and 80-mg atomoxetine once daily (QD) and 5-mg and 60-mg duloxetine QD. The human unbound plasma binding for atomoxetine ($f_{up} = 0.013$) (Sauer et al., 2003) and duloxetine was incorporated in the modeling such that only unbound drug in plasma was available to cross the BBB and BCB.

Rat PK parameters were scaled to humans using allometric principles based on the following equation,

$$P_h = P_r \times \left(\frac{W_{th}}{W_{tr}} \right)^{0.75} \quad (10)$$

where P_h is the scaled human parameter, P_r is the model-predicted parameter in rat, W_{th} is the average human brain weight (1.35 kg) and W_{tr} is the average rat brain weight (0.0015 kg).

Based on an adult brain volume of 1.35 L, the estimated human V_{ECF} was 0.31 L and V_{BC} was 1.04 L (Segal, 1993). The human V_{CSF} was fixed in the model to be 0.16 L (Shen et al., 2004).

Statistical Analysis

Unless otherwise indicated, all data shown are represented as the mean and standard deviation.

For graphical presentation, total (bound + unbound) plasma concentrations are shown.

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Results

Plasma and Brain Binding Assessment

At 3 μM , duloxetine was highly bound to plasma and brain with an unbound fraction of 0.047 ± 0.002 ($n=3$) and 0.00283 ± 0.00012 ($n=3$), respectively.

Duloxetine Pharmacokinetics in Rats

Figure II shows the duloxetine total plasma, CSF, bECF and whole brain concentrations following duloxetine administration ($n=13$). The C_P and C_{ECF} were 762 ± 421 ng/mL ($n=13$) and 19 ± 9 ng/mL ($n=13$), respectively. The C_B and C_{CSF} were 3467 ± 755 ng/mL ($n=13$) and 29 ± 8 ng/mL ($n=9$). Duloxetine concentration ratios were $C_B/C_P=6 \pm 3$ ($n=13$), $C_{ECF}/C_P=0.027 \pm 0.012$ ($n=13$) and $C_{CSF}/C_P=0.043 \pm 0.017$ ($n=9$). The C_{ECF}/C_{uP} and C_{CSF}/C_{uP} were 0.6 ± 0.3 ($n=13$) and 0.9 ± 0.4 ($n=9$), respectively. The C_B/C_{ECF} and C_B/C_{CSF} were 233 ± 142 ($n=13$) and 128 ± 76 ($n=9$), respectively. The C_{BC} was 4477 ± 976 ng/mL ($n=13$) and the C_{BC}/C_{ECF} and C_{uBC}/C_{ECF} were 301 ± 183 ($n=13$) and 0.9 ± 0.5 ($n=13$), respectively.

The rat PK model incorporates both physiologically-based parameters and compartmental PK parameters. We used compartmental PK estimation of clearance at the BBB to inform the nature (passive, active) and extent of drug transport across the BBB, which also accounts for potential transporter expression or activity at the BBB. The CNS PK data obtained for atomoxetine (Kielbasa 2009) or duloxetine fit better to a model parameterized in terms of distributional clearance at the BBB (Q_{BBB}) rather than bidirectional clearances at the BBB (CL_{PL-ECF} and CL_{ECF-PL}), where CL_{PL-ECF} represents BBB influx clearance and CL_{ECF-PL} represents BBB efflux

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clearance. This result supported the passive transport process for atomoxetine and duloxetine. This may not be the case of other drugs that have non-passive transport at the BBB.

The rat duloxetine PK data supported estimation of model parameters with acceptable precision (Table I). Diagnostic plots (not shown) of the model population and individual predicted concentrations versus measured concentrations were scattered close to a line of unity, indicating an acceptable fit of the data to the model. Together, these results indicate that the model adequately characterized the disposition of unbound duloxetine. In Table I, rat atomoxetine PK parameters determined previously (Kielbasa et al., 2009) are shown to support the translational modeling approach for atomoxetine herein.

Prediction of Atomoxetine and Duloxetine Human bECF Pharmacokinetics

Table II shows the atomoxetine and duloxetine human PK parameters used as inputs in the translational models to generate the human simulations (Lobo et al., 2009). Atomoxetine exposure is substantially influenced by cytochrome P450 2D6 (CYP2D6) polymorphism (Farid et al., 1985). The clinical atomoxetine PK parameters shown in Table II were estimated from subjects that were extensive metabolizers of the CYP2D6 enzyme.

Figure III illustrates the predicted PK profiles following administration of 40-mg and 80-mg atomoxetine QD and the in vitro human norepinephrine transporter inhibition constant (hNET K_i = 1.1 ng/ml). At 80-mg atomoxetine, the median plasma and bECF maximal concentrations (C_{max}) were about 561 ng/ml and 5 ng/ml, respectively. The median bECF concentration was about 4-5 times greater than the hNET K_i at t_{max} (1-2 h) and 1.5 ng/ml at 24 h postdose, which was slightly above the hNET K_i . At 40-mg atomoxetine, the median plasma and bECF C_{max}

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were about 280 ng/ml and 2.4 ng/ml, respectively. The median bECF concentration was about 2-3 times greater than the hNET K_i at t_{max} and at 16 h and 24 h postdose, the median bECF concentrations were 1.1 ng/ml and 0.74 ng/ml, respectively. The hNET K_i value is lower than or contained within the 90% prediction interval of the bECF concentrations for both doses of atomoxetine. The $t_{1/2}$ of atomoxetine in plasma and bECF were about 4 h and 13 h, respectively. The plasma-to-bECF ratios for AUC_{τ} ($AUC_{\tau, PL}/AUC_{\tau, ECF}$) and C_{max} ($C_{max, PL}/C_{max, ECF}$) were about 55 and 116, respectively, irrespective of the dose. These predictions show that 40-mg and 80-mg atomoxetine would be expected to have bECF concentrations in the target range of the hNET K_i , with 80 mg having a greater bECF exposure and duration above the hNET K_i . Figure IV illustrates the predicted PK profiles following administration of 5-mg and 60-mg duloxetine QD and the hNET $K_i = 2$ ng/ml and the in vitro human serotonin transporter inhibition constant (hSERT $K_i = 0.07$ ng/ml). At 60-mg duloxetine, the median plasma and bECF C_{max} were about 70 ng/ml and 2 ng/ml, respectively. The bECF concentration was essentially unchanged during the dosing interval and about 29 times greater than the hSERT K_i and similar to the hNET K_i . At 5-mg duloxetine, the plasma and bECF C_{max} were about 6 ng/ml and 0.17 ng/ml, respectively. The bECF concentration was about 2-3 times greater than the hSERT K_i , but the hNET K_i was about 12 times greater than the bECF concentration over the 24h dosing interval. The $t_{1/2}$ of duloxetine in plasma and bECF were about 16 hours and 92 hours, respectively. The $AUC_{\tau, PL}/AUC_{\tau, ECF}$ and $C_{max, PL}/C_{max, ECF}$ ratios were about 28 and 34, respectively, irrespective of the dose. These predictions show that 5-mg and 60-mg duloxetine would be expected to have bECF concentrations substantially above the hSERT K_i and in the target range of the hNET K_i at 60 mg only. At 5 mg, duloxetine bECF concentrations are substantially below the hNET K_i .

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Discussion

Pharmacokinetics of Duloxetine

Based on the $C_{ECF}/C_{uP} = 0.6$ and $C_{CSF}/C_{uP} = 0.9$ and being within 3 times of unity (Maurer et al., 2005; Kalvass et al., 2007; Doran et al., 2005) we conclude that duloxetine transport across the BBB and BCB is primarily passive. Similar values for duloxetine f_{uP}/f_{uB} and C_B/C_P of 17 and 23, respectively, supports the same conclusion at the BBB (Maurer et al., 2005). Estimations of C_B , C_{BC} and C_{uBC} and their comparison to C_{ECF} provide insight regarding the brain distribution of duloxetine. The C_B/C_{ECF} and C_{BC}/C_{ECF} ratios of 233 and 301, respectively, indicate that duloxetine does not preferentially reside in bECF, which is suggestive of nonspecific binding to brain tissue. A $C_{uBC}/C_{ECF} = 0.9$ indicates that duloxetine passively distributes within brain parenchyma.

Translational Pharmacokinetic Modeling of Duloxetine and Atomoxetine

The derived translational human PK model was used to simulate human PK in plasma and bECF, the matrix in direct contact with the hNET and hSERT. Based on the free drug hypothesis, it is a commonly accepted assumption that unbound drug is the entity available for interaction with drug targets. It is also assumed that unbound drug in the brain is in direct contact or in equilibrium with the site of action (de Lange and Danhof, 2002). Several key assumptions were used for this translational modeling approach. First, both drugs studied do not have active metabolite(s), or active metabolites in sufficient concentrations that cross the BBB and BCB and interact with the target transporters to elicit pharmacologic activity. Although extensive metabolism occurs for atomoxetine (Sauer et al., 2003) and duloxetine (Lantz et al., 2003), their pharmacologic activity is not believed to be due to target engagement from metabolites. Second,

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hNET and hSERT K_i can be used as target concentrations that would be expected to have appreciable interaction at the target transporter. Third, the mechanism of transport across the BBB and BCB in rat and human is similar. In our evaluations, we show that duloxetine and atomoxetine (Kielbasa et al., 2009) transport in rats is primarily passive; therefore, we assume that this remains the case for humans. Lastly, we assume that rat CNS PK model parameters can be translated accurately to humans using allometric principals based on differences in brain weight between species. To corroborate this assumption, PK parameters were compared from the scaling approach and clinical PK data following 80 mg QD atomoxetine (Table II). In theory, bidirectional clearances (CL_{PL-CSF} and CL_{CSF-PL}) and Q_{BCB} are similar if the drug primarily crosses the BCB by passive diffusion. The rat atomoxetine Q_{BCB} as shown in Table I was scaled to humans using equation 10, resulting in a human Q_{BCB} prediction of 0.015 L/h. This value was similar to the actual human CL_{PL-CSF} of 0.00825 L/h and CL_{CSF-PL} of 0.0205 L/h shown in Table 2 (the mean of CL_{PL-CSF} and CL_{CSF-PL} = 0.014 L/h) and suggests that the translational scaling approach has merit for predicting the human CSF PK from rats.

Ideally, such translational CNS PK models could be used to evaluate the influence of the model parameters on the predicted human PK by simulating various scenarios with altered PK parameters (and variability). For example, the impact of efflux clearance at the human BBB (CL_{ECF-PL} , clearance of drug from the bECF to plasma) in relation to plasma concentration may be explored if data exist suggesting the drug may be a substrate for P_{gp} in human. Simulation can lead to optimized study designs and effective management of clinical plans and strategies.

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The translational approach for predicting human bECF PK is not high-throughput methodology and is not well-suited for incorporation into a project flow scheme in early drug discovery.

Alternatively, it is applicable in supporting candidate drugs selected for clinical investigation or to discriminate between a smaller set of compounds that have been identified as potential clinical candidates. For atomoxetine and duloxetine, a wealth of clinical PK data was available to validate the translational approach. In the context of drug development when human PK have yet to be determined, *in vitro* or allometric techniques could be used initially to support model development. When human systemic PK data are obtained from a clinical study, the model can then be updated.

At 40-mg and 80-mg atomoxetine, the predicted human bECF concentrations are suggestive of target inhibition at the NET (Figure III). Putting these results in clinical context, according to the atomoxetine product label in the US, dosing adults, children and adolescents over 70-kg body weight should be initiated at 40 mg QD and increased after a minimum of 3 days to approximately 80 mg QD. After 2 to 4 additional weeks, the dose may be increased to a maximum of 100 mg QD in patients who have not achieved an optimal response. Biochemical evidence for norepinephrine (NE) reuptake inhibition can be provided through evaluation of the NE metabolite 3,4-dihydroxyphenylglycol (DHPG) and assessment of DHPG can provide insight of a drug's effect on NET function. In humans given 80-mg atomoxetine QD, both plasma and CSF DHPG concentrations were reduced indicating that atomoxetine has activity at the NET (Kielbasa et al., 2006). Also, children with ADHD given atomoxetine exhibited changes in the urine DHPG/NE ratio (Montoya et al., 2011). The prediction of human bECF PK at 40 mg and 80 mg inhibiting hNET transport is consistent with clinical dosing recommendations and human atomoxetine biomarker data.

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Human bECF concentrations of duloxetine were predicted at 5 mg QD and at a clinically efficacious dose of 60 mg QD. Since the duloxetine mechanism of action is to act at the hNET and hSERT, the human bECF PK predictions relative to in vitro potency at those transporters were examined. The predicted bECF concentrations were below the hNET K_i during the dosing interval suggesting that 5-mg duloxetine QD would be expected to have no or minimal activity at the hNET. Conversely, at 60-mg duloxetine QD, the predicted bECF concentrations were similar to the hNET K_i during the entire dosing interval of 24 hours. These predictions are corroborated with published reports where target activity of duloxetine was assessed by measurements of NE and DHPG. At 60-mg duloxetine QD, reductions of DHPG in CSF, plasma and urine were observed (Quinlan et al., 2009). It was also demonstrated that the plasma DHPG/NE ratio was reduced significantly at 2 weeks of treatment with 80-mg duloxetine QD, the lowest dose tested in the study (Vincent et al., 2004). Similar results were obtained when assessing NE metabolites in urine (Chalon et al., 2003). No reports of the effects of duloxetine on NE and DHPG appear to exist at 5-mg duloxetine. In rat ECF, concentrations of atomoxetine or duloxetine that were comparable to or greater than their respective rat NET K_i resulted in increased NE concentrations (data not shown).

There is currently no validated PET tracer to measure brain NET occupancy, but brain SERT blockade by duloxetine was measured in humans using PET (Takano et al., 2006). In that study, the SERT receptor occupancy (RO) at 6 h postdose was about 44% at 5 mg and 82% at 60 mg when duloxetine was given as a single dose. After 60-mg duloxetine QD, the SERT RO at 6 hours postdose was 84% and estimated to be about 78% about 24 h later indicating little change in SERT RO during the dosing interval at steady state. Serotonin RO at 60-mg duloxetine is consistent with clinically efficacious doses of SSRIs providing over 80% occupancy (Meyer et

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al., 2004). The predicted duloxetine bECF PK agree with the dose-dependent SERT RO findings clinically, including the ability to discriminate a 5-mg dose from the clinically effective 60-mg dose. The data show that duloxetine total plasma and predicted bECF PK profiles are distinct. The plasma $t_{1/2}$ was about 16 hours, as expected, and the bECF $t_{1/2}$ was about 92 hours demonstrating a relatively stable profile and lesser peak to trough fluctuation in bECF concentrations over the dosing interval. Similar results were obtained for atomoxetine, where the plasma $t_{1/2}$ was about 4 hours, as expected, and the predicted bECF $t_{1/2}$ was about 13 hours. Interestingly, in the study by Takano et. al. (2006), the duloxetine plasma concentrations declined appreciably while SERT RO did not decrease to any reasonable extent during the dosing interval at steady state, which is consistent with the predicted PK profile in bECF. These data indicate that bECF predictions for drugs with appreciable binding, like atomoxetine and duloxetine, may be a better surrogate matrix than plasma for estimating the time-course of brain target engagement. Furthermore, prediction of bECF PK in humans may be insightful when considering the clinical dose and frequency of administration.

The exploratory translational approach aims to predict the human dose-bECF concentration relationship to identify clinical doses that should engage the brain target and provide insight into early drug development about the dose and regimen to test the clinical hypotheses. Regardless of the mechanism of a centrally acting drug or the indication being investigated, without significant activity at the CNS target a positive clinical outcome would not be expected. Based on the results obtained, we believe this translational approach merits consideration as a robust tool to support the clinical development of CNS-mediated drug candidates. This approach may enhance the ability to predict pharmacologically relevant doses in the absence of, or in association with, other biomarker approaches.

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Authorship Contributions

Participated in research design: Kielbasa and Stratford

Conducted experiments: Kielbasa and Stratford

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Footnotes

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There are no numbered footnotes in this manuscript.

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Figure Legends

Figure I. Schematic diagram of the rat duloxetine neuropharmacokinetic model

Figure II. Duloxetine concentrations in plasma, bECF, CSF and whole brain during an intravenous infusion maintenance dose of duloxetine at 4.2 mg/kg/h in rats

Figure III. Model-predicted human plasma and bECF concentrations of atomoxetine following once daily administration 40 mg (upper panel A) and 80 mg (lower panel B). *Shown are the predicted median and 90% confidence interval of plasma and bECF atomoxetine concentrations. The dashed horizontal line represents the atomoxetine inhibition constant determined for the human norepinephrine reuptake transporter.*

Figure IV. Model-predicted human plasma and bECF concentrations of duloxetine following once daily administration of 5 mg (upper panel A) and 60 mg (lower panel B). *Shown are the predicted median and 90% confidence interval of plasma and bECF duloxetine concentrations. The solid and dashed horizontal lines represent the duloxetine inhibition constant determined for the human norepinephrine and serotonin reuptake transporters, respectively.*

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Tables

Table I. Model-estimated Rat Pharmacokinetic Parameters of Atomoxetine^x and Duloxetine

Parameter	Atomoxetine		Duloxetine	
	Estimate (% SEE)	Variability (% SEE)	Estimate (% SEE)	Variability (% SEE)
CL _u (L/h)	14.5 (8.69)	38.6 (23.7)	35.6 (17.6)	64.6 (35.5)
V _u (L)	6.92 (30.8)	65.4 (80.1)	33.2 (15.7)	55.7 (46.3)
Q _{BCB} (L/h)	0.0000909 (29.9)	---	0.0000527 (31.7)	---
Q _{BBB} (L/h)	0.00110 (34.5)	---	0.000160 (15.8)	55.4 (39.9)
CL _{ECF-CSF} (L/h)	0.000129 (20.2)	---	0.000024 (fixed)	---
CL _{ECF-BC} (L/h)	0.00216 (25.1)	---	0.00168 (41.8)	---
CL _{BC-ECF} (L/h)	0.000934 (25.7)	---	0.00249 (50.6)	60.6 (42.8)
V _{BC} (L)	0.000990 (fixed)	---	0.000990 (fixed)	---
V _{CSF} (L)	0.000250 (fixed)	---	0.000250 (fixed)	---
V _{ECF} (L)	0.000290 (fixed)	---	0.000290 (fixed)	---
σ _{prop}	34.7 (9.09)	---	28.8 (14.6)	---

% SEE, percentage S.E. of the estimate; σ_{prop}, proportional residual error

^xKielbasa W, Kalvass JC, and Stratford R (2009) Microdialysis evaluation of atomoxetine brain penetration and central nervous system pharmacokinetics in rats. *Drug Metab Dispos* **37**: 137-142.

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Table II. Pharmacokinetic Parameters of Atomoxetine and Duloxetine^x Used in Translational Models to Predict Human Brain Extracellular Pharmacokinetics

Parameter	Atomoxetine		Duloxetine	
	Clinical	Scaled from Rat	Clinical	Scaled from Rat
k_a (h^{-1})	3.1	---	0.168	---
CL/F(L/h)	20.6	---	45.1	---
V/F (L)	121	---	814	---
V_{CSF} (L)	0.16	---	0.16	---
V_{ECF} (L)	0.31	---	0.31	---
V_{BC} (L)	1.04	---	1.04	---
CL_{PL-CSF} (L/h)	0.00825	---	NE	---
CL_{CSF-PL} (L/h)	0.0205	---	NE	---
Q_{BCB} (L/h)	NE	0.015	---	0.009
Q_{BBB} (L/h)	---	0.181	---	0.026
$CL_{ECF-CSF}$ (L/h)	---	0.021	---	0.004
CL_{ECF-BC} (L/h)	---	0.355	---	0.279
CL_{BC-ECF} (L/h)	---	0.153	---	0.412
ω -CL/F (%)	90.8	---	58.9	---
ω -V/F (%)	65.6	---	96.6	---

ω , inter-subject variability; NE, not estimated

^xLobo ED, Quinlan T, O'Brien L, Knadler MP, and Heathman M (2009) Population pharmacokinetics of orally administered duloxetine in patients: implications for dosing recommendation. *Clin Pharmacokinet* **48**: 189-97.

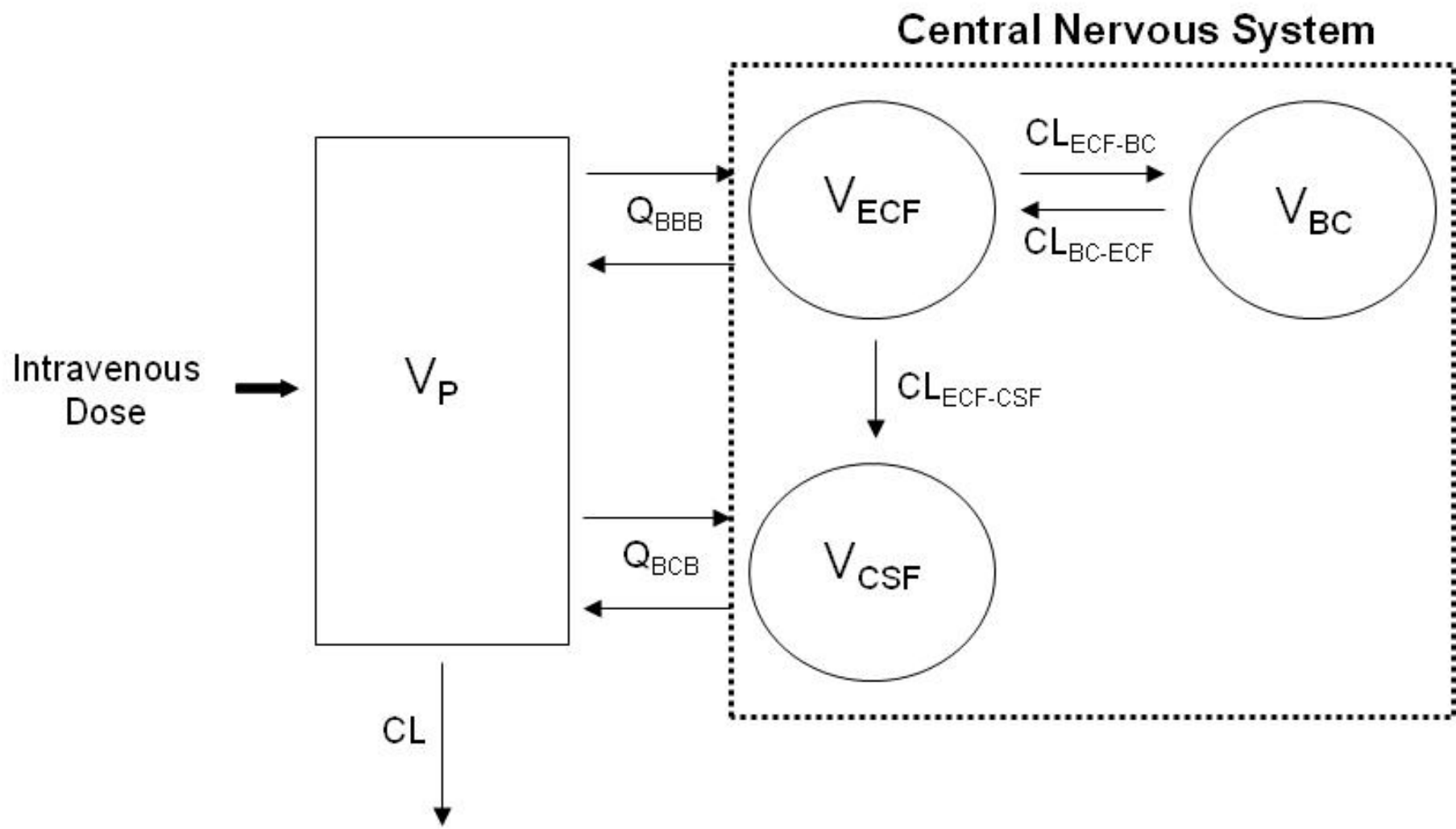


Figure I

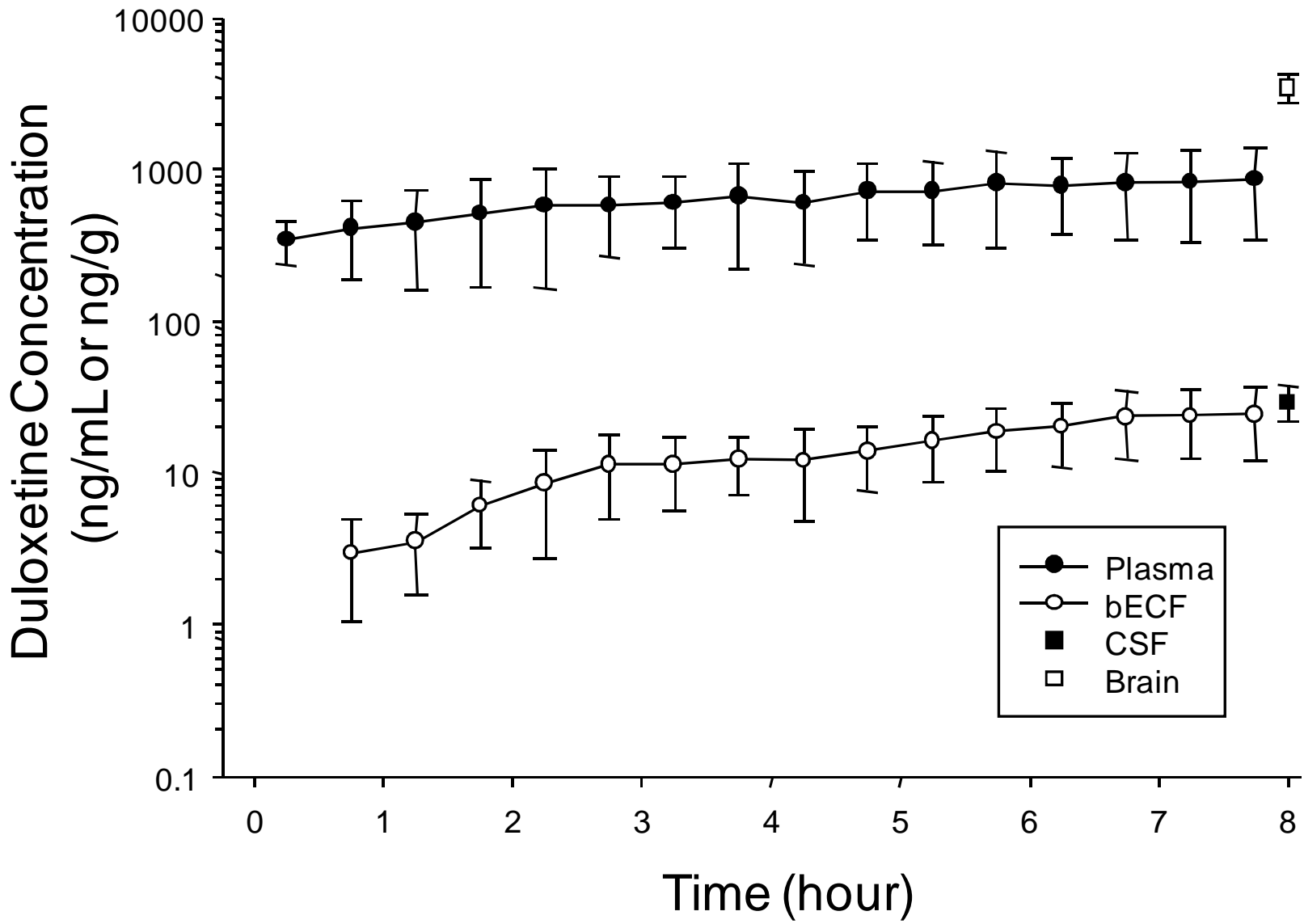


Figure II

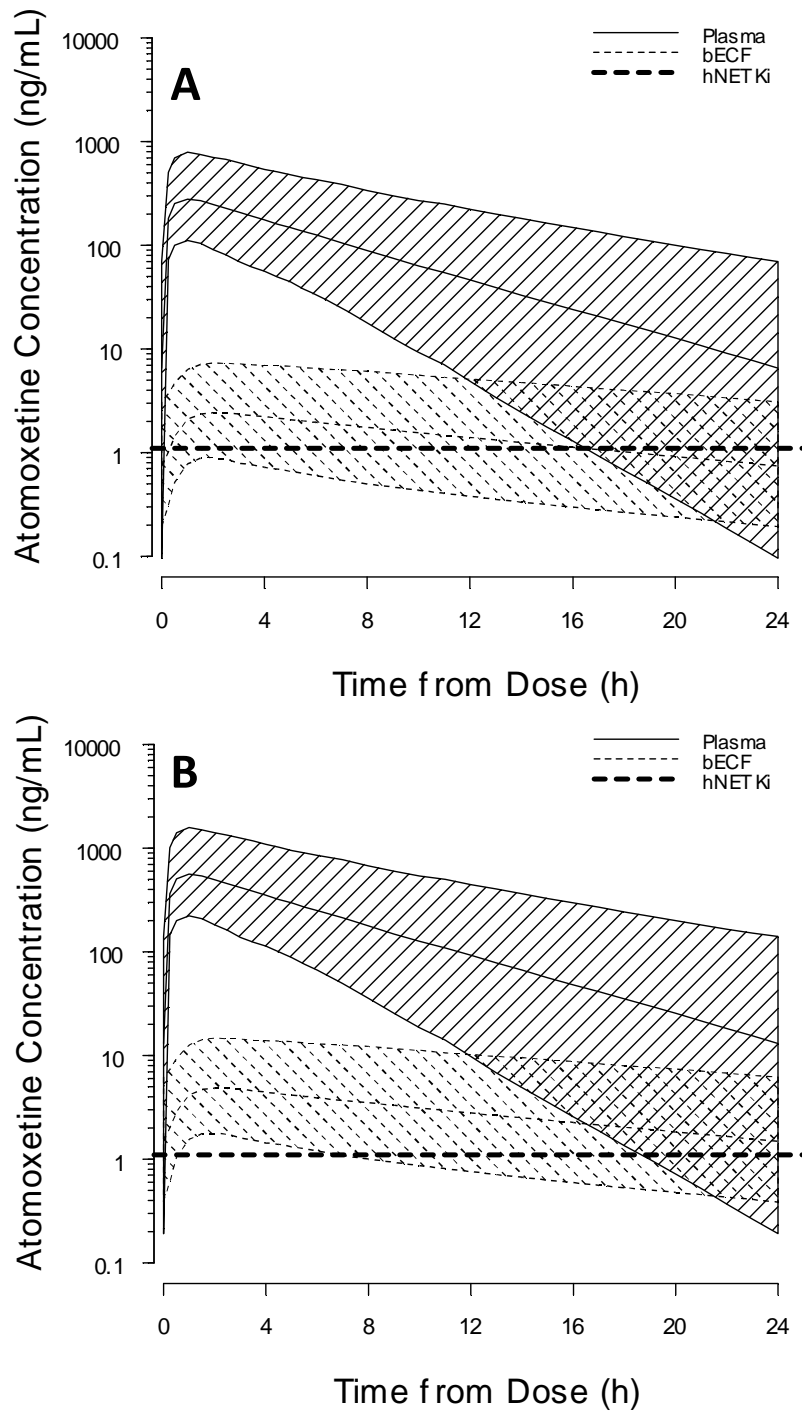


Figure III

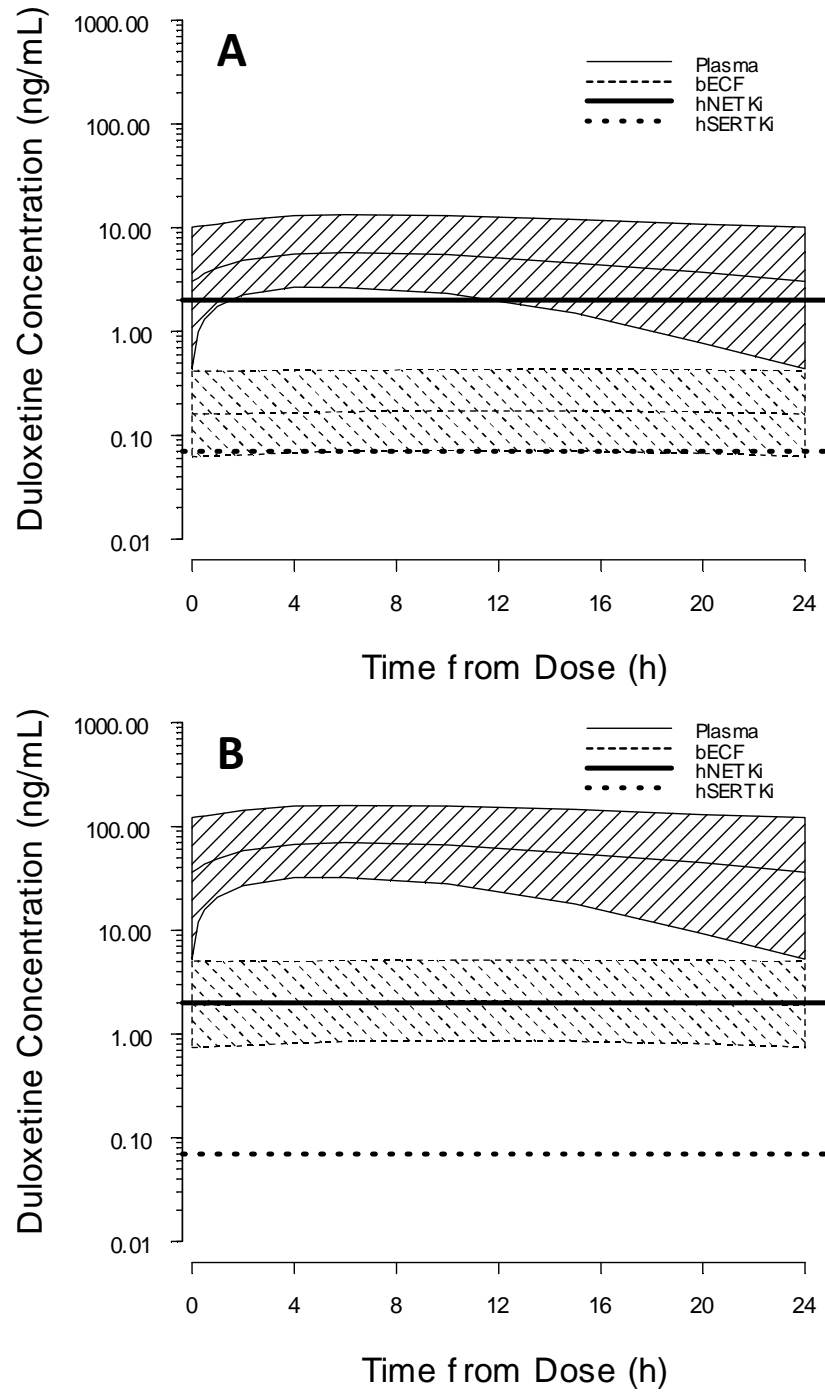


Figure IV