

Population Modelling of Dexmedetomidine Pharmacokinetics and Haemodynamic Effects After Intravenous and Subcutaneous Administration

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Abstract

Background and Objective Dexmedetomidine is a potent agonist of α_2 -adrenoceptors causing dose-dependent sedation in humans. Intravenous dexmedetomidine is commonly used perioperatively, but an extravascular route of administration would be favoured in palliative care. Subcutaneous infusions provide desired therapeutic plasma concentrations with fewer unwanted effects as compared with intravenous dosing. We aimed to develop semi-mechanistic population models for predicting pharmacokinetic and pharmacodynamic profiles of dexmedetomidine after intravenous and subcutaneous dosing. **Methods** Non-linear mixed-effects modelling was performed using previously collected concentration and haemodynamic

effects data from ten (eight in the intravenous phase) healthy human subjects, aged 19–27 years, receiving 1 μ g/kg of intravenous or subcutaneous dexmedetomidine during a 10-min infusion.

Results The absorption of dexmedetomidine from the subcutaneous injection site, and distribution to local subcutaneous fat tissue was modelled using a semi-physiological approach consisting of a depot and fat compartment, while a two-compartment mammillary model explained further disposition. Dexmedetomidine-induced reductions in plasma norepinephrine concentrations were accurately described by an indirect response model. For blood pressure models, the net effect was specified as hyper- and hypotensive effects of dexmedetomidine due to vasoconstriction on peripheral arteries and sympatholysis mediated via the central nervous system, respectively. A heart rate model combined the dexmedetomidine-induced sympatholytic effect, and input from the central nervous system, predicted from arterial blood pressure levels. Internal evaluation confirmed the predictive performance of the final models, as well as the accuracy of the parameter estimates with narrow confidence intervals.

Conclusions Our final model precisely describes dexmedetomidine pharmacokinetics and accurately predicts dexmedetomidine-induced sympatholysis and other pharmacodynamic effects. After subcutaneous dosing, dexmedetomidine is taken up into subcutaneous fat tissue, but our simulations indicate that accumulation of dexmedetomidine in this compartment is insignificant.

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1 Introduction

Patients in palliative care frequently present with pain and anxiety. These symptoms are commonly managed with opioids, benzodiazepines, antidepressants and antipsychotic agents [1, 2], all of which have clinically significant adverse effects. Alternatives to the current regimes are therefore needed to achieve clinically adequate analgesia and anxiolysis, while avoiding undesired drug effects.

Dexmedetomidine is a selective and potent sedative used in anaesthesia and pain medicine for its anxiolytic, sedative and analgesic properties [3]. Dexmedetomidine mediates its effects via α_2 -adrenoceptors to inhibit the release of

Key Points

Using previously collected data and a sequential modelling approach, we were able to describe dexmedetomidine absorption kinetics and further disposition together with its sympatholytic and haemodynamic effects.

Accurate models for haemodynamic effects were obtained by combining norepinephrine-dependent sympatholysis with either dexmedetomidine-evoked hypertension or changes in central neural activity.

Our final models precisely describe dexmedetomidine pharmacokinetics and accurately predicts dexmedetomidine-induced sympatholysis and other pharmacodynamic effects.

norepinephrine (NE) from presynaptic nerve terminals and to hyperpolarise postsynaptic neurons [4]. Its unique mechanism of action causes dose-dependent sedation without a significant risk of respiratory depression, making dexmedetomidine a promising agent for palliative sedation.

Although dexmedetomidine is only registered for intravenous (IV) administration, many alternative routes of administration have been documented for off-label use [3-5]. Our previous work indicates that subcutaneous (SC) infusion of dexmedetomidine is a feasible alternative to the IV route, especially when venous access is unattainable [6]. In our article, we have documented that the plasma concentrations of dexmedetomidine increase to clinically adequate sedative concentrations after SC administration, while the sympatholytic effects are significantly attenuated as compared with IV dosing [6]. The sedative profile of dexmedetomidine has been very well characterised [7, 8], but only a few studies have evaluated the pharmacokinetic-pharmacodynamic (PK-PD) relationship and quantitatively described the various clinically observed effects of dexmedetomidine [9-13], especially after extravascular dosing [9].

Population-based PK analysis of dexmedetomidine has been reported in the literature, with a range of study subjects from healthy adult volunteers [9, 13] to hospitalised children [14–16], and critically ill adult patients [17, 18]. Most of these studies were conducted using data collected after IV dosing, and PD data were included only in a few studies [9–13]. To our knowledge, population-based PK-PD analysis of subcutaneously administered dexmedetomidine has not been previously described.

Our aim was to develop semi-mechanistic population models using healthy volunteers' data to characterise the pharmacokinetics of dexmedetomidine and its sympatholytic and haemodynamic effects after IV and SC dosing. A suitable dataset was available from our previously published clinical study [6]. We hypothesised that our population-based PK-PD models would be able to: (1) accurately describe absorption of subcutaneously administered dexmedetomidine, (2) account for the disposition kinetics of dexmedetomidine, and (3) predict the effects of dexmedetomidine on plasma NE levels, heart rate (HR), and systolic and diastolic arterial blood pressure (SAP and DAP, respectively) after IV and SC dosing.

2 Materials and Methods

2.1 Study Protocol

We have reanalysed our data from a previously published clinical trial [6] (registered as ClinicalTrials.org identifier NCT02724098). The study protocol (EudraCT 2015-004698-34) was approved by the Ethics Committee of the Hospital District of Southwest Finland (184/1800/2015) and by the Finnish National Agency for Medicines (3/2016). Detailed information about the study protocol has been provided in the Electronic Supplementary Material (ESM). Briefly, a two-period randomised crossover study was conducted to investigate the PK and pharmacological effects of subcutaneously and intravenously administered dexmedetomidine in ten (two dropouts for the IV arm of the study) healthy human volunteers. We administered either a single IV or SC dose of 1 µg/kg of dexmedetomidine during a 10-min infusion in a randomised order. Arterial blood samples were collected pre-dose and then at 5, 10, 15, 20, 30 and 45 min and 1, 1.5, 2 and 3 h after the start of the infusions into EDTA tubes for determination of concentrations of dexmedetomidine, NE and epinephrine in plasma. Additional blood samples were collected at 4, 5, 6, 8 and 10 h for determination of dexmedetomidine. Heart rate, SAP, DAP and peripheral oxygen saturation were recorded at times of blood sampling. Plasma concentrations of dexmedetomidine were determined with high-performance liquid chromatography-tandem mass spectrometry and concentrations of NE and epinephrine in plasma were determined with high-performance liquid chromatography and coulometric electrochemical detection as described previously [6, 19].

2.2 Data Analysis

Non-linear mixed effects modelling was conducted using the NONMEM[®] (version 7.3.0) software package (ICON Development Solutions, Ellicott City, MD, USA) [20] and supported by the Perl-Speaks-NONMEM toolkit (PsN version 4.6.0) [21], for model development, execution and evaluation. RStudio (version 1.0.153) using R (version 3.6.0) [22] was used for diagnostic model evaluation and graphical visualisations.

2.3 Model Development

Population models were specified with differential equations that were solved with ADVAN13 subroutine and parameters estimated using first-order conditional estimation with interaction. A sequential PK-PD approach was used as described previously [23, 24]. In the first step, a PK model was developed using IV administration data only, followed by the implementation of a semi-mechanistic absorption model for SC administration. Next, a PD model was developed for NE/epinephrine concentration–time data. Finally, effect models for the PD parameters, i.e. HR, SAP and DAP were developed.

2.3.1 Structural Models

For dexmedetomidine pharmacokinetics, we started with a simple one-compartment empirical model, and nested models were built in a stepwise manner using two- and three-compartmental models. Zero-order input of IV dexmedetomidine, and linear pharmacokinetics with first-order elimination clearance were assumed. A semi-mechanistic model describing dexmedetomidine absorption from the SC site of administration during a SC infusion were added to the disposition model. The inhibitory effect of dexmedetomidine on the spill-over of NE and epinephrine into plasma was modelled using indirect response models. Decreases in catecholamine concentrations and their effects were then used as surrogate measures of the changes in subsequent pharmacodynamic endpoints.

2.3.2 Stochastic Model

The between-subject variability on the fixed-effect parameters in the PK and PD models was specified using exponential model,

$$\Phi_i = \theta \bullet \mathrm{e}^{\eta_i}$$

where Φ_i is the fixed effect parameter for an individual, θ is the population estimate of the parameter and η_i describes the between-subject variability of the *i*th individual from the population estimate.

Between-subject variability was added to the fixed-effect parameters in the models gradually in nested models and was retained if it led to a significant reduction in the objective function value (OFV), did not cause numerical instability in the models, and if the shrinkage of the added betweensubject variability was less than or equal to 25%. For the PK models of intravenously and subcutaneously administered dexmedetomidine, the residual variability between observations was coded into the models with a constant coefficient of variation error model,

$$C_{ij,x} = \widehat{C}_{ij,x} \bullet (1 + \varepsilon_{ij,x}).$$

An additive error model was used for NE and epinephrine concentrations, and for the remaining PD variables,

$$C_{ij,x} = \hat{C}_{ij,x} + \varepsilon_{ij,x,x}$$

where x is one of the dexmedetomidine/NE/epinephrine concentrations or one of the remaining PD endpoints, $C_{ij,x}$ is the *j*th observation for the *i*th individual at the xth time point, $\hat{C}_{ij,x}$ is the corresponding model-predicted value and $\varepsilon_{ij,x}$ is a normally distributed random variable to quantify the residual variability in the values of x. η and ε were assumed to be normally distributed random-effect parameters.

2.3.3 Model Evaluation

The primary criterion for discerning amongst nested models was the difference in the OFV (Δ OFV) at a significance level of 0.05, which equates to a reduction of 3.84 points in the OFV for every added degree of freedom. Akaike information criteria (AIC) was used for non-nested models. Relative standard errors and retrieval of covariance matrices were used to check model precision and parameter identifiability. Standard goodness-of-fit plots and visual predictive checks [25] were routinely used to test model appropriateness and predictive performance. The precision in the final model parameters was estimated using the sampling importance resampling (SIR) procedure [26]. To achieve robustness in the final SIR results, the procedure was run with 40,000 samples and 2000 resamples. Covariance matrices from the final models were used as a sample proposal distribution. Output from these SIR runs were evaluated using the 95% confidence intervals in the parameter estimates, OFV distribution and graphical output from the SIR package in PsN.

2.3.4 Simulations

We also assessed model performance with simulations using the final parameter estimates for the fixed and random effects in 1000 virtual subjects with NONMEM. The first set of simulations was conducted with the final PK model only, to evaluate model output for dexmedetomidine concentration–time profiles for short- and long-term infusions using the IV and SC routes. Two clinically relevant infusion time scales of 8 h and 48 h were used at dexmedetomidine doses of 0.25, 0.5, 1, 2, 3 and 4 μ g/kg/h. These simulations were conducted to evaluate the time required for achieving steady-state plasma concentrations and to compare relative areas under the curve achieved with SC administration as compared to the IV route. Additionally, we evaluated the deposition of dexmedetomidine in the SC fat tissue after SC dosing to investigate possible accumulation after SC drug administration for possible clinical implications.

Next, the same dosing schemes were used to simulate the effect of dexmedetomidine on the NE concentration-time course in plasma. These simulations were used to evaluate decreases in NE levels during continuous infusions in a dose-dependent manner. The aim was to check a model's performance in dynamically predicting the inhibition of NE spill-over over increasing doses and in a temporal manner, and to identify the dose required for a near-complete cessation of NE spill-over into the plasma.

Third, different dexmedetomidine dosing schemes and the resulting NE profiles were used to evaluate drug effects on haemodynamic parameters. Because dexmedetomidine influences blood pressure by two distinct mechanisms, model simulations were used to provide information on the development of NE-associated hypotension (sympatholysis) and dexmedetomidine-evoked hypertension (direct vasoconstriction) over an increasing dose scale. Simultaneously, HR model simulations were conducted to evaluate the dose dependency of HR inhibition by dexmedetomidine, as well as the interplay between the inhibition of NE release and other central nervous system (CNS) effects contributing to dexmedetomidine-evoked bradycardia. Last, the ncappc package of PsN [27] was used as an internal evaluation method to perform a non-compartmental analysis (NCA)based comparison between the observed values in the dataset to model-derived predictions for the PK parameters.

3 Results

A schematic of the final PK–PD model is shown in Fig. 1 and Table 1 shows the comparison between different models tested during model development. A more detailed description of the model development including the set of mass transfer differential equations is provided in the ESM. The model estimates in all the final PK and PD models were computed without bias, and numerical and graphical diagnostics demonstrated adequate precision and robustness of the PK (Table 2, Fig. 2) and PD models (Table 3, Figs. 2, 3, 4).

3.1 Pharmacokinetic Model

Pharmacokinetic modelling of dexmedetomidine concentration-time data began with an empirical one compartment model (OFV = -306), but the model fit was poor and therefore two- and three-compartment mammillary models were considered. Although the OFV reduced significantly for a three-compartment model (Δ OFV = -348) as compared with a two-compartment model (Δ OFV = -325), bootstrap analysis demonstrated wide confidence intervals for the added model parameters, and therefore the two compartmental system was retained as the final structural PK model.

A first-order absorption process was initially tested to describe the absorption of dexmedetomidine after a short SC infusion ($\Delta OFV = -659$, AIC = -1276), as previously reported for nasal mucosal bioavailability [9], but visual predictive checks demonstrated inconsistent output. Multiple different solutions were tested (see the ESM, p. 3) without success. Finally, a semi-mechanistic biphasic absorption model was constructed (AIC = -1436), such that a fast absorption process ($k_{a,FAST}$) governed the influx of drug from a SC depot into the systemic circulation and a fat layer permeation constant (k_{FAT}) was added to describe drug distribution into the local SC fat tissue. The addition of bioavailability parameters in the final PK model for the SC dosing phase improved the model fit ($\Delta OFV = -201$, AIC = -1469).

Fractional permeation of subcutaneously administered dexmedetomidine was estimated in the PK model $(F_{\text{FAT}} = 0.55)$, while the fast absorption process from the depot $(K_{a \text{ FAST}})$ was linked with a bioavailability parameter $(F_{\text{DEPOT}} = 0.79)$ that governed the systemic procurement of dexmedetomidine only via the fast process. A slow absorption process $(k_{a \text{ SLOW}})$ was included to describe drug release from the SC fat tissue. Although this scenario represented a biphasic absorption process, no strict criterion was coded into the model to describe the temporal shift between the two processes, and both were modelled simultaneously to represent a dynamic interplay between fast and slow absorption processes from the injection site. An attempt to model a bioavailability parameter associated with slow absorption from a fat compartment failed with numerical instability and over-parametrisation. Net bioavailability of dexmedetomidine (0.89) through SC administration was estimated assuming complete recovery of the drug from the fat layer into the blood circulation.

The final PK model produced biologically plausible parameter estimates (Table 2). Our results demonstrate that inclusion of a fat compartment into the model allowed adequate description of the data. The rate constants governing SC absorption were estimated with high precision (Table 2) and the visual predictive checks for the final PK model (Fig. 2) show good predictive performance. In addition, the results indicate that the ratio of fast to slow absorption of dexmedetomidine from the local injection depot and from the SC fat tissue compartment is approximately 1:9, while the capacity of permeation into the SC fat tissue compartment is about three-fold compared to the capacity of the simultaneous fast absorption process from the depot compartment into the systemic circulation (Table 2).





Fig. 1 Schematic of the final semi-mechanistic model for dexmedetomidine pharmacokinetics (**a**), its effect on norepinephrine (NE) release (**b**), systolic and diastolic blood pressures (**c**), and heart rate (**d**). A amount, C_I dexmedetomidine concentration on central compartment, C_E amount of (heart rate/systolic or diastolic blood pressure), *CNS* central nervous system, *EC* effect compartment, EC_{50} concentration causing 50% effect from E_{MAX} , E_{MAX} maximum effect,

The final PK model was used to calculate the area under the curve (AUC_{0-∞} = (DOSE × *F*)/CL) levels for the study population, and the levels of AUC_{0-∞} were also simulated using the NCA package in PsN [21]. The model calculated levels of the AUC_{0-∞} were 2.01 and 1.82 ng · h/mL for the IV and SC phase, respectively, while the simulated values of these variables were synchronous with calculated results, at 1.94 and 1.78 ngh/mL respectively using the NCA package. Although our results demonstrate a lower relative bioavailability of dexmedetomidine in the SC dosing phase, comparable values of AUC_{0-∞} establish the strong clinical utility of the SC route of dexmedetomidine administration.

 F_{spill} fraction on NE spilled out from the release compartment, *HPN* hypotension, *HR* heart rate, *HTN* hypertension, *I*(*t*) dosing event, *IV* intravenous, k_{1e} , k_{5e} and k_{7e} rate constants for distribution to the effect compartment, $k_{IN,R}$ rate constant for NE release, k_a absorption rate constant, k_{e0} elimination rate constant for the effect, $k_{OUT,P}$ rate constant for NE elimination from the plasma, *P* plasma, *R* release, *SC* subcutaneous, γ hill parameter

3.2 Norepinephrine Model

An indirect response model of NE was used, and baseline levels of NE were estimated for each individual in the final model with an inter-individual variability parameter. Plasma levels of NE represent an approximately 15% spill-over fraction of the amount of NE released from sympathetic nerve endings [28], and our NE model was specified as a composite of two separate compartments, representing sympathetic nerves (release compartment) and plasma (circulatory compartment), respectively. Thereafter, the initial conditions for the NE release compartment were specified with the rate constants for the resting-state release and spill-over of NE from the release compartment. Because dexmedetomidine is known to reduce the rate of the resting-state release of NE from sympathetic nerve endings in vivo, an inhibitory effect

Model	Degrees of freedom	OFV ^a	Change in OFV	Akaike information criterion
Dexmedetomidine PK				
One-compartment model	2	- 306	_	_
Two-compartment model	6	- 631	- 325	-
Three-compartment model ^b	8	- 654	- 348	-
PK + absorption				
2 CMT + first-order absorption (with K_a)	7	- 1290	_	- 1276
2 CMT + first-order absorption + absorption lag time	8	- 1433	- 143	- 1417
2 CMT + single Weibull function	8	- 1310	- 20	- 1294
2 CMT + biphasic absorption	8	- 1430	- 140	- 1414
2 CMT + transit compartment	8	- 1350	- 70	- 1334
2 CMT + Michaelis-Menten absorption	8	- 1416	- 126	- 1400
2 CMT + fat compartment	9	- 1454	- 164	- 1436
2 CMT + fat compartment + bioavailability	11	- 1491	- 201	- 1469
Norepinephrine				
One-compartment model	6	- 1641	-	- 1629
Two-compartment model (with F _{SPILL})	8	- 2229	- 588	- 2213
Systolic blood pressure				
NE-based sympatholysis	8	- 567	_	- 551
NE + dexmedetomidine-induced hypertension	13	- 622	- 55	- 596
Diastolic blood pressure				
NE-based sympatholysis	7	- 828	_	- 814
NE + dexmedetomidine-induced hypertension	11	- 920	- 92	- 898
Heart rate				
NE-based sympatholysis	6	- 1106	_	- 1094
NE + dexmedetomidine-induced hypertension ^c	12	389	_	413

Models with major structural changes have been reported in this table to highlight the modelling steps. Smaller changes, e.g. addition of interindividual variation, and changes in the residual variability model have not been included. Final models are shown in bold

^a*CMT* compartment, F_{SPILL} fraction on norepinephrine spilled out from the release compartment, K_a absorption rate constant, *NE* norepinephrine, *OFV* objective function value, *PK* pharmacokinetics

^bThe two-compartment model was chosen because the three-compartment model showed model instability and a high relative standard error in added parameters

^cFor the final heart rate model, the data file was modified to include population values of the final systolic blood pressure model

on the release rate constant $(k_{IN,R})$ at time = t was coded into the model using a indirect response model,

$$k_{\text{IN,R}} = k_{\text{IN,R}}^{t=0} \bullet \left(1 - \frac{C_1}{\text{EC}_{50,\text{DEX}} + C_1}\right),$$

where $k_{\text{IN,R}}^{t=0}$ defines the resting-state rate of NE release at t = 0, C_1 is the concentration of dexmedetomidine in the central compartment and EC_{50,DEX} is the concentration of dexmedetomidine that causes 50% of the maximal inhibitory effect on NE release in vivo.

Dexmedetomidine EC_{50} for NE release was estimated at approximately 0.33ng/mL (Table 2), which is numerically close to the maximum concentrations achieved in the study subjects with SC dosing of 1 µg/kg, and much lower than the

drug concentrations achieved after IV dosing. The values of the rate constant for NE spill-over from the release compartment ($k_{\text{OUT,R}}$) and the rate constant for NE elimination from the plasma ($k_{\text{OUT,P}}$) were 9.03 h⁻¹ and 11.4 h⁻¹, respectively, which are in accordance with a recent report modelling the inhibition of NE release following dexmedetomidine challenge in humans [9].

3.3 Epinephrine Model

Our data for epinephrine concentrations included multiple below limit of quantification values. All attempts to model epinephrine failed, indicating that these models were overparametrised to provide a numerically stable predictive model output.

Table 2	Parameter estimates	s from the final ph	armacokinetic mo	odel for dexr	nedetomidine	(DEX) and nore	pinephrine (NE)	with median	and 95%
confider	ce intervals (CIs) fr	om the sampling i	mportance resam	oling (SIR)	procedure with	n 40,000 final sar	mples and 2000 i	resamples	

Parameter	Description	Parameter estimate		SIR results	
		Mean	%RSE	Median	95% CI
DEX					
Cl ₁	Elimination clearance (L/h)	31.1	8.74	31.1	27.5-34.6
V_1	Volume of the central compartment (L)	15.0	64.2	14.8	9.34-22.2
Cl_2	Intercompartmental clearance (L/h)	82.4	8.49	82.7	73.2–92.9
V_2	Volume of the peripheral compartment (L)	58.6	13.0	58.8	52.7-65.3
k _{a,FAST}	Rate constant for fast absorption (h^{-1})	0.983	36.5	0.978	0.673-1.363
k _{a,SLOW}	Rate constant for slow absorption (h^{-1})	0.151	41.6	0.149	0.105-0.199
k _{FAT}	Rate constant for partitioning to subcutaneous fat (h^{-1})	3.36	39.8	3.28	2.21-4.64
FDEPOT	Bioavailability of the fraction left in the depot after fat permeation	0.788	15.2	0.792	0.671-0.941
F _{FAT}	Fraction of drug that enters the fat layer	0.552	29.0	0.581	0.390-0.813
$\eta_{1,\text{CL1}}$	Inter-individual variability on CL ₁	0.124	76.1	0.137	0.005-0.045
$\eta_{2,V1}$	Inter-individual variability on V ₁	0.456	53.9	0.448	0.068-0.431
$\eta_{3.F.DEPOT}$	Inter-individual variability on F _{DEPOT}	0.131	57.0	0.145	0.004-0.048
$\eta_{4,\mathrm{F,FAT}}$	Inter-individual variability on F _{FAT}	0.337	78.8	0.371	0.021-0.358
$\sigma_{\mathrm{PROP,IV}}$	Proportional RE for intravenous dosing	0.129	17.7	0.131	0.013-0.022
$\sigma_{\mathrm{PROP,SC}}$	Proportional RE for subcutaneous dosing	0.189	7.2	0.185	0.027-0.042
NE					
$E_{\rm MAX}$	Scaling factor for E _{MAX}	1 (fixed)			
EC _{50,NE}	DEX concentration at 50% inhibition in NE release (nM)	0.334	14.5	0.333	0.252-0.426
$C_{\rm E0}$	Baseline NE concentration in effect compartment (nM)	0.814	20.5	0.816	0.634-1.018
k _{OUT,R}	Rate constant for NE spill-over from release compartment (h ⁻¹)	9.03	14.3	9.218	7.14–11.5
$f_{\rm SPILL}$	Spill-over fraction to plasma	0.15 (fixed)			
k _{OUT,P}	Rate constant for NE elimination from plasma (h ⁻¹)	11.4	7.53	11.5	10.3-12.8
$\eta_{1,\text{CE0}}$	Interindividual variability on C _{E.0}	0.446	20.7	0.230	0.120-0.401
$\eta_{2,kOUT,R}$	Interindividual variability on k _{OUT,R}	0.237	32.7	0.072	0.022-0.143
$\sigma_{ m ADD,NE,IV}$	Additive RE for NE model after intravenous dosing	0.097	11.0	0.100	0.084–0.117
$\sigma_{ m ADD,NE,SC}$	Additive RE for NE model after subcutaneous dosing	0.125	6.6	0.126	0.11-0.141

The subjects were given a single $1-\mu g/kg$ DEX infusion in two phases either intravenously or subcutaneously in 10 min *RE* residual error, *RSE* relative standard error

3.4 Systolic/Diastolic Arterial Blood Pressure Models

Decreases in SAP and DAP after dexmedetomidine administration were assumed to result from reduced NE levels in the release compartment, and a sigmoidal $E_{\rm MAX}$ model with a biophase was utilised to estimate individual SAP and DAP values. The hypertensive effect of dexmedetomidine through α 2-adrenoceptors has been documented before [13], and therefore we assumed that a binomial response, i.e. combined, and dose-dependent hypertensive (α 2-adrenoceptor mediated) and hypotensive (NE mediated) effect was applicable to both SAP and DAP. The hypotensive effects of NE ($E_{\rm BP,HPN$) concentration fluctuations and hypertensive blood pressure effects due to plasma dexmedetomidine concentrations ($E_{\rm BP,HTN}$) on SAP and DAP were defined in the model as,

$$E_{\rm BP,HPN} = 1 + \left(\frac{E_{\rm MAX} \bullet C_{\rm E,NE}{}^{\gamma_{\rm HPN}}}{{\rm EC}_{50,\rm NE}{}^{\gamma_{\rm HPN}} + C_{\rm E,NE}{}^{\gamma_{\rm HPN}}}\right),$$

$$E_{\rm BP,HTN} = 1 + \left(\frac{E_{\rm MAX} \bullet C_{\rm E,DEX}^{\gamma_{\rm HTN}}}{\rm EC_{50,DEX}^{\gamma_{\rm HTN}} + C_{\rm E,DEX}^{\gamma_{\rm HTN}}}\right),$$

where E_{MAX} is the maximum effect on SAP or DAP and was fixed to 1 for both of these circumstances, C_E is the apparent concentration of either NE or dexmedetomidine in the effect compartments, and EC₅₀ is the concentration of NE or dexmedetomidine resulting in half-maximal effects for hypotension and hypertension, respectively. γ is the hill coefficient that signifies the slope of the concentration–response curve for the development of hypotensive or hypertensive effects.

7.5

2

10.0

3



Fig. 2 Visual predictive checks based on 1000 simulations showing dexmedetomidine and norepinephrine after a 1-µg/kg intravenous (a, c) or subcutaneous (b, d) dexmedetomidine infusion. The black solid and dashed lines are the observed percentiles (10th, 50th and 90th

Then, the combined hypotensive and hypertensive effect of NE and dexmedetomidine on SAP or DAP, respectively, was modelled as,

$$E_{\rm BP} = E_{\rm MIN,BP} \bullet \left(E_{\rm BP,HPN} + E_{\rm BP,HTN} \right),$$

where $E_{\text{MIN,BP}}$ is the minimum blood pressure estimated as a model parameter.

A binomial response model for SAP and DAP data was clearly preferable over the models that used NE as the only surrogate measure of these PD endpoints. Models using NE as the only measure of blood pressure development produced final OFV values of -567 and -828, respectively. A binomial response model for SAP and DAP data with hypertensive action of dexmedetomidine resulted in a significant OFV reduction (- 622 and - 920 for SAP and DAP, respectively).



В

0.5

0.4

0.3

0.2

0.1

Subcutaneous

The effect-site rate constant for hypotensive NE effect on SBP levels ($k_{e0 \text{ HPN}}$) was estimated at 22.1 h⁻¹ and for DBP at 24.3 h^{-1} (Table 3), showing a fast equilibration between the release compartment and the effect site, which is biologically plausible considering the action of NE on blood pressure in humans. The model-predicted EC_{50.NE} for SAP and DAP was 4.56 ng/mL and 4.32 ng/mL, respectively, and the values of γ_{HPN} were 6.01 and 4.99 for SAP and DAP, respectively, showing that small changes in NE levels can have marked changes in blood pressure.

In comparison, the values of effect-site rate constants for a dexmedetomidine-induced hypertensive effect $(k_{e0,HTN})$ on SAP and DAP were estimated at 22.1 h^{-1} and 30 h^{-1} , respectively, which indicates rapid equilibration between the plasma and effect compartment but also signifies very short half-lives for these processes in vivo. The values of these rate constants were fixed to their respective population

Table 3 Pharmacodynamic parameters for blood pressure and heart rate (HR) models

$\begin{tabular}{ c c c c c c c c c c c c c c c c c c c$	Parameter	Description	Parameter estimate		SIR results	
$ \begin{array}{cccc} SAP \\ k_{ALRPN} & Biophase rate constant for hypotensive NE effect (h^{-1}) 22.1 27.7 22.9 17.7-30.4 \\ k_{ALRPN} & Biophase Rife or causing 50% maximal SAP (nM) 4.56 16.7 4.57 3.49-5.80 \\ First Hill coefficient for hypotension 6.01 15.9 6.29 4.93-7.90 \\ k_{MISAP} & SAP at 100% NE inhibition (nmHg) 136 8.7 134 123-145 \\ E_{SAAP} & Baseline SAP (mMHg) 136 8.7 134 123-145 \\ E_{MAX} & Maximum SAP response 11 (fixed) \\ k_{ALTN} & Biophase rate constant for hypertensive effect (ng/mL) 2.37 59.1 2.11 0.829-4.06 \\ THE-CADER & Diophase DEX for 50% hypertensive effect (ng/mL) 2.37 59.1 2.11 0.829-4.06 \\ T_{RESONE} & Inter-individual variability on EC_{SONE} 0.445 39.9 0.521 0.346-0.718 \\ T_{RESONE} & Inter-individual variability on EC_{SONE} 0.104 70.8 0.137 0.067-0.200 \\ T_{RESONE} & Additive residual error for nancenous dosing 0.53 14.4 9.50 & 8.22-11.0 \\ GADDN & Additive residual error for subcutaneous dosing 10.8 10.7 10.6 9.23-11.0 \\ CADDN & Additive residual error for subcutaneous dosing 10.8 10.7 10.6 9.23-11.0 \\ CADDN & Additive residual error for nancenous dosing 10.8 10.7 10.6 9.23-11.0 \\ CADDN & CADDN & Additive residual error for nancenous dosing 10.8 10.7 10.6 9.23-11.0 \\ CADDN & CADDN & DAP at 100% NE hilbition (nmHg) 30.2 2.1 30.3 29.5-31.2 \\ CADDN & CADDN & Biophase rate constant for hypotensive NE effect (h^{-1}) 30 (fixed) C_{SINE} Biophase rate constant for hypotensive effect (ng/mL) 0.761 40.3 0.812 0.482-1.25 \\ TAR & Hill coefficient of hypotensive effect (ng/mL) 0.761 40.3 0.812 0.482-1.25 \\ TAR & Maximum DAP response 1 (fixed) C_{SINE} Biophase rate constant for hypotensive effect (ng/mL) 0.761 40.3 0.812 0.482-1.25 \\ TAR & Hill coefficient or hypotensive effect (ng/mL) 0.761 40.3 0.812 0.482-1.25 \\ TAR & Maximum DAP response 1 (fixed)$			Mean	%RSE	Median	95% CI
k_{clIPN} Biophase NE for causing 50% maximal SAP (nM) 4.56 16.7 4.57 3.49-5.80 P_{MNN} Bill coefficient for hypotension 6.01 1.59 6.29 4.93.7.90.4 E_{MNSAP} SAP at 100% NE inhibition (mmHg) 49.6 1.6 49.4 48.2-50.8 E_{MASAP} Baseline SAP (mmHg) 136 8.7 1.23 1.23-145 E_{MASN} Biophase rate constant for hypertensive effect (h ⁻¹) 2.21 (fixed) - - - E_{GLDEN} Biophase tract for 50% hypertensive effect (h ⁻¹) 2.21 (fixed) - - - - T_{RENOR} Inter-individual variability on EC _{SADE} 0.104 7.08 0.32 0.346-0.718 T_{EESORE} Inter-individual variability on EC _{SADE} 0.77 62.5 1.13 0.0692-1.57 σ_{ADDSC} Additive residual error for intravenous dosing 10.8 10.7 10.6 9.23-1.10 σ_{ADDSC} Additive residual error for intravenous dosing 10.8 10.7 10.6 9.23-1.10 σ_{ADDSC}	SAP					
$\begin{array}{ccccccc} EC_{SUNE} & Biophase NE for causing 50% maximal SAP (nM) & 4.56 & 16.7 & 4.57 & 3.49-5.80 \\ r_{PRM} & Ihil coefficient for hypotension & 6.01 & 15.9 & 6.29 & 4.307-780 \\ E_{SAP} & Baseline SAP (mmHg) & 49.6 & 1.6 & 49.4 & 48.2-50.8 \\ E_{SAP} & Baseline SAP (mmHg) & 136 & 8.7 & 134 & 123-145 \\ E_{MAX} & Maximum SAP response & 1 (fixed) & - & - & - \\ EC_{GaDRX} & Biophase the constant for hypertensive effect (n-1) & 2.1 (ixed) & - & - & - \\ EC_{GaDRX} & Biophase DEX for 50% hypertensive effect (ng/mL) & 2.37 & 59.1 & 2.11 & 0.829-4.06 \\ PHN & Ihil coefficient for hypertensive effect (ng/mL) & 2.37 & 59.1 & 0.346-0.718 \\ \eta_{ECSABP} & Inter-individual variability on EC_{SANE} & 0.445 & 39.9 & 0.521 & 0.346-0.718 \\ \eta_{ECSABP} & Inter-individual variability on EC_{SANE} & 0.777 & 62.5 & 1.13 & 0.092-1.57 \\ \sigma_{ADDX} & Additive residual error for intravenous dosing & 9.53 & 14.4 & 9.50 & 8.22-11.0 \\ \sigma_{ADDX} & Additive residual error for subcutaneous dosing & 10.8 & 10.7 & 10.6 & 9.23-11.0 \\ DAP & - & - & - \\ k_{GUBN} & Biophase rate constant for hypotensive NE effect (h-1) & 24.3 & 61.1 & 29.4 & 17.4-50.0 \\ C_{NRB} & Biophase rate constant for hypotensive NE effect (h-1) & 24.3 & 61.1 & 29.4 & 17.4-50.0 \\ C_{NRB} & Biophase rate constant for hypotensive RE effect (h-1) & 30.1 & 30.3 & 29.5-31.2 \\ E_{MAX} & Maximum DAP response & 1 (fixed) - & - & - \\ EC_{SADEX} & Biophase rate constant for hypotensive effect (h-1) & 30 (fixed) - & - & - \\ EC_{SADEX} & Biophase rate constant for hypotensive effect (h-1) & 30 (fixed) - & - & - \\ EC_{SADEX} & Biophase rate constant for hypotensive effect (h-1) & 30 (fixed) - & - & - \\ EC_{SADEX} & Biophase rate constant for hypertensive effect (h-1) & 30 (fixed) - & - & - \\ EC_{SADEX} & Biophase rate constant for hypertensive effect (h-1) & 30 (fixed) - & - & - \\ EC_{SADEX} & Inter-individual variability on EC_{SANE} & 0.382 & 65.5 & 0.434 & 0.295-0.633 \\ \eta_{AECSANE} & Inter-individual variability on EC_{SANE} & 0.382 & 65.5 & 0.434 & 0.295-0.633 \\ \eta_{AECSA$	$k_{\rm e0,HPN}$	Biophase rate constant for hypotensive NE effect (h ⁻¹)	22.1	27.7	22.9	17.7-30.4
$\gamma_{\rm HPR}$ Hill coefficient for hypotension6.0115.96.294.937.790 $E_{\rm MINSAP}$ SAP at 100% NE inhibition (mmHg)49.61.649.44.932.708 $E_{\rm SAAP}$ Baseline SAP (mmHg)1368.7134123-145 $E_{\rm SAAV}$ Biophase rate constant for hypertensive effect (ng/mL)2.3759.12.110.829-4.06 $P_{\rm RESOBIE}$ Inter-individual variability on $E_{\rm SAAP}$ 0.1047.080.1370.036-0.718 $\gamma_{\rm RESOBIE}$ Inter-individual variability on $E_{\rm SAAP}$ 0.1047.080.1370.067-0.200 $\gamma_{\rm EESOAPE}$ Additive residual error for intravenous dosing9.531.449.508.22-11.0 $\sigma_{\rm ADDSC}$ Additive residual error for intravenous dosing0.1681.0710.69.23-1.10 $\sigma_{\rm ADDSC}$ Additive residual error for subcutaneous dosing1.081.0710.69.23-1.10 $\sigma_{\rm ADDSC}$ Additive residual error for subcutaneous dosing0.322.130.32.92-3.12.0 $\sigma_{\rm ADDSC}$ Biophase rate constant for hypotensive effect4.992.525.073.27-7.16 $F_{\rm MINDAP}$ Baseline DAP (mmHg)30.22.130.32.95-3.12 $F_{\rm MINDAP}$ DAP (nom Hg)3.022.130.32.95-3.12 $F_{\rm ADDSC}$ Biophase rate constant for hypetensive effect (h ⁻¹)3.0 (fixed) $F_{\rm ADDSC}$ Biophase rate constant for hypetensive effect (h ⁻¹)3.01 (fixed) <td>EC_{50.NE}</td> <td>Biophase NE for causing 50% maximal SAP (nM)</td> <td>4.56</td> <td>16.7</td> <td>4.57</td> <td>3.49-5.80</td>	EC _{50.NE}	Biophase NE for causing 50% maximal SAP (nM)	4.56	16.7	4.57	3.49-5.80
E_{MRXAP} SAP at 100% NE inhibition (mmHg) 49.6 1.6 49.4 48.2-50.8 E_{MAX} Bascline SAP (mmHg) 136 8.7 1.34 123-145 E_{MAX} Biophase rate constant for hypertensive effect (1^{-1}) 2.11 (fixed) - - E_{MAX} Biophase DEX for 50% hypertensive effect (1^{-1}) 2.37 59.1 2.11 0.829-4.06 YHTN Hill coefficient for hypertensive effect (1^{-1}) 2.37 59.1 0.21 0.346-0.71 PLESCORE Inter-individual variability on EC _{50.NE} 0.77 6.25 1.13 0.062-1.57 $\sigma_{\text{ADD,OC}}$ Additive residual error for intravenous dosing 9.53 1.44 9.50 8.22-11.0 $\sigma_{\text{ADD,OC}}$ Additive residual error for subcutaneous dosing 0.8 10.7 10.6 9.23-11.0 $\sigma_{\text{ADD,OC}}$ Additive residual error for subcutaneous dosing 1.8 0.61-5.64 3.0 9.27-1.70 $K_{0.01W}$ Biophase rate constant for waximal DAP (nM) 4.32 2.3.5 4.54 3.61-5.64 P_{100}	$\gamma_{\rm HPN}$	Hill coefficient for hypotension	6.01	15.9	6.29	4.93-7.90
E_{03AP} Baseline SAP (mmHg) 136 8.7 134 123-145 E_{MAX} Maximum SAP response 1 (fixed) - - - E_{03DEX} Biophase ruc constant for hypertensive effect (n^{-1}) 2.21 (fixed) - - - R_{LEOOR} Biophase DEX for 50% hypertensive effect (n_2/mL) 2.37 59.1 2.11 0.829-4.06 R_{LEOOR} Inter-individual variability on E_{05AP} 0.104 7.08 0.137 0.067-0.200 R_{LEOOR} Inter-individual variability on E_{05AP} 0.104 7.08 0.137 0.067-0.200 R_{LEOOR} Inter-individual variability on E_{05AP} 0.77 6.25 1.13 0.067-0.200 R_{LEOOR} Inter-individual variability on E_{05AP} 0.77 6.25 1.13 0.067-0.200 $\sigma_{ADD,V}$ Additive residual error for intravenous dosing 9.53 14.4 9.50 8.22-11.0 DAP Varth Matiour varthize	E _{MIN.SAP}	SAP at 100% NE inhibition (mmHg)	49.6	1.6	49.4	48.2-50.8
$ \begin{array}{ccccc} E_{AAX} & Maximum SAP response & 1 (fixed) & - & - & - & - & - & - & - & - & - & $	$E_{0,\text{SAP}}$	Baseline SAP (mmHg)	136	8.7	134	123-145
$k_{\rm MITN}$ Biophase rate constant for hypertensive effect (h ⁻¹) 22.1 (fixed) - - - $\rm EC_{SDEX}$ Biophase DEX for 50% hypertensive effect (g/mL) 2.37 59.1 2.11 0.829-4.06 $\gamma_{\rm HTN}$ Hill coefficient for hypertensive effect 1.52 29.9 0.521 0.346-0.718 $\eta_{\rm LESDAR}$ Inter-individual variability on EC _{SDEX} 0.77 62.5 1.13 0.0692-1.57 σ_{ADDN} Additive residual error for intravenous dosing 9.53 14.4 9.50 8.22-11.0 σ_{ADDN} Additive residual error for subcutaneous dosing 10.8 10.7 10.6 9.232-11.0 σ_{ADDN} Additive residual error for subcutaneous dosing 10.8 10.7 10.6 9.232-11.0 σ_{ADDN} Additive residual error for subcutaneous dosing 10.8 10.7 10.6 9.22-1.10 σ_{ADDN} Biophase rate constant for hypotensive NE effect (h ⁻¹) 24.3 61.1 29.4 17.4-50.0 EC_SODE Biophase rate constant for hypertensive effect (h ⁻¹) 30.2 2.1 30.3 29	E _{MAX}	Maximum SAP response	1 (fixed)	-	_	_
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$k_{\rm e0.HTN}$	Biophase rate constant for hypertensive effect (h^{-1})	22.1 (fixed)	-	_	_
THTM Hill coefficient for hypertensive effect 1.52 29.9 1.37 0.896–1.95 $\eta_{LECSORE}$ Inter-individual variability on EC _{SORE} 0.445 39.9 0.521 0.346–0.718 $\eta_{LECSORE}$ Inter-individual variability on EC _{SOREX} 0.77 62.5 1.13 0.067–0.200 $\eta_{LECSOREX}$ Additive residual error for intravenous dosing 9.53 1.44 9.50 8.22–11.0 $\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing 10.8 1.0.7 10.6 9.23–11.0 DAP Statistical error for subcutaneous dosing 10.8 1.0.1 29.4 17.4–50.0 EC_SORE Biophase R1 for causing 50% maximal DAP (nM) 4.32 2.3.5 4.54 3.61–5.64 γ_{IPN} Hill coefficient for hypotensive effect (h ⁻¹) 24.3 30.3 22–5.71.6 E_{MNNAP} DAP at 100% NE inhibition (mmHg) 30.2 2.5 3.3 24–5.71.6 k_{MIN} Maximum DAP response 1 (fixed) - - - E_{ADAP} Biophase	EC _{50.DEX}	Biophase DEX for 50% hypertensive effect (ng/mL)	2.37	59.1	2.11	0.829-4.06
η_{LESONE} Inter-individual variability on EC_{SONE} 0.445 39.9 0.521 0.346-0.718 η_{LESONEX} Inter-individual variability on EC_{SONEX} 0.77 62.5 1.13 0.067-0.200 η_{LESONEX} Inter-individual variability on EC_{SONEX} 0.77 62.5 1.13 0.692-1.57 $\sigma_{\text{ADD,SC}}$ Additive residual error for intravenous dosing 9.33 14.4 9.50 82.2-11.0 DAP 10.6 9.23-11.0 $\sigma_{\text{ADD,SC}}$ Biophase rate constant for hypotensive NE effect (h ⁻¹) 24.3 61.1 29.4 1.7.4-50.0 EG_SONE Biophase rate constant for hypotensive NE effect (h ⁻¹) 24.3 5.1 3.03 29.5-31.2 EMINDAP DAP at 100% NE inhibition (mmHg) 30.2 2.1 30.3 29.5-31.2 EMINDAP Biophase Tate constant for hypertensive effect (h ⁻¹) 30 61.5 0.7 83.4 79.6-87.1 EMINDAP DAP at 1000% NE inhibition (mmHg) 302 6.5 0.434 0.295-31.2 <t< td=""><td>γ_{HTN}</td><td>Hill coefficient for hypertensive effect</td><td>1.52</td><td>29.9</td><td>1.37</td><td>0.896-1.95</td></t<>	γ _{HTN}	Hill coefficient for hypertensive effect	1.52	29.9	1.37	0.896-1.95
η_{LESAP} Inter-individual variability on $\mathbb{E}_{0,\text{SOPEX}}$ 0.104 70.8 0.137 0.067-0.200 η_{LECONDEX} Inter-individual variability on $\mathbb{E}_{0,\text{SOPEX}}$ 0.77 62.5 1.13 0.692-1.57 $\sigma_{\text{ADD,XC}}$ Additive residual error for intravenous dosing 9.53 14.4 9.50 8.22-11.0 DAP .	$\eta_{1 EC50 NE}$	Inter-individual variability on EC _{50 NE}	0.445	39.9	0.521	0.346-0.718
matrix Inter-individual variability on EC _{SODEX} 0.77 62.5 1.13 0.692-1.57 $\sigma_{ADD,IV}$ Additive residual error for intravenous dosing 9.53 14.4 9.50 8.22-11.0 $\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing 10.8 10.7 10.6 9.23-11.0 DAP	$\eta_{2 E0 SAP}$	Inter-individual variability on $E_{0 SAP}$	0.104	70.8	0.137	0.067-0.200
or appLiv AppLic Additive residual error for intravenous dosing 9.53 14.4 9.50 8.22-11.0 $\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing 10.8 10.7 10.6 9.23-11.0 DAP	$\eta_{3 EC50 DEX}$	Inter-individual variability on $EC_{50 DEX}$	0.77	62.5	1.13	0.692-1.57
Additive residual error for subcutaneous dosing 10.8 10.7 10.6 9.23-11.0 $A_{0,0HPN}$ Biophase rate constant for hypotensive NE effect (h ⁻¹) 24.3 61.1 29.4 17.4-50.0 $EC_{50,NE}$ Biophase NE for causing 50% maximal DAP (nM) 4.32 23.5 4.54 3.61-5.64 γ_{HPN} Hill coefficient for hypotensive effect 4.99 25.2 5.07 3.27-7.16 $E_{0,DAP}$ Baseline DAP (mHg) 83.5 5.7 83.4 79.6-87.1 $E_{0,0AP}$ Baseline DAP (mHg) 83.5 5.7 83.4 79.6-87.1 $E_{0,0AP}$ Biophase rate constant for hypertensive effect (ng/mL) 0.761 40.3 0.812 0.482-1.25 γ_{HTN} Hill coefficient for hypertensive effect (ng/mL) 0.761 40.3 0.812 0.482-1.25 γ_{HTN} Hill coefficient for inpertensive effect (ng/mL) 0.761 40.3 0.812 0.482-1.25 $\gamma_{LCSO,NE}$ Inter-individual variability on EC_{50,NEX} 0.541 66.8 0.635 0.364-0.979 $\sigma_{ADD,NC}$ Additive residual error for intravenous dosing 6.30 8.7 6.32 <t< td=""><td>$\sigma_{\rm ADDIV}$</td><td>Additive residual error for intravenous dosing</td><td>9.53</td><td>14.4</td><td>9.50</td><td>8.22-11.0</td></t<>	$\sigma_{\rm ADDIV}$	Additive residual error for intravenous dosing	9.53	14.4	9.50	8.22-11.0
DAP Normal State Normal	$\sigma_{\rm ADD,SC}$	Additive residual error for subcutaneous dosing	10.8	10.7	10.6	9.23-11.0
$k_{e0,HPN}$ Biophase rate constant for hypotensive NE effect (h ⁻¹)24.361.129.417.4-50.0EC_{50,NE}Biophase NE for causing 50% maximal DAP (nM)4.3223.54.543.61-5.64 γ_{HPN} Hill coefficient for hypotensive effect4.9925.25.073.27-7.16 $E_{MIN,DAP}$ DAP at 100% NE inhibition (mmHg)30.22.130.329.5-31.2 E_{0DAP} Baseline DAP (mmHg)83.55.783.479.6-87.1 E_{MAX} Maximum DAP response1 (fixed) E_{0DAF} Biophase rate constant for hypertensive effect (ng/mL)0.76140.30.8120.482-1.25 γ_{HTN} Hill coefficient for hypertensive effect (ng/mL)0.54166.80.6350.364-0.979 $\sigma_{ADD,JV}$ Additive residual error for intravenous dosing6.308.76.325.61-7.04 $\sigma_{ADD,JC}$ Additive residual error for subcutaneous dosing6.1525.76.283.68-8.97 $E_{0,HR}$ Biophase rate constant for NE in HR effect (h ⁻¹)6.1525.76.283.68-8.97 $F_{0,HR}$ Biophase NE at 50% of the maximal HR (nM)3.1225.93.20 <td< td=""><td>DAP</td><td>-</td><td></td><td></td><td></td><td></td></td<>	DAP	-				
EC_{SUNE} CSUNE Biophase NE for causing 50% maximal DAP (nM)4.3223.54.543.61-5.64 γ_{HPN} Hill coefficient for hypotensive effect4.9925.25.073.27-7.16 $E_{MIN,DAP}$ 	$k_{\rm e0\;HPN}$	Biophase rate constant for hypotensive NE effect (h^{-1})	24.3	61.1	29.4	17.4-50.0
THENHill coefficient for hypotensive effect4.9925.25.073.27-7.16 $E_{MIN,DAP}$ DAP at 100% NE inhibition (mmHg)30.22.130.329.5-31.2 $E_{0,DAP}$ Baseline DAP (mmHg)83.55.783.479.6-87.1 E_{MAX} Maximum DAP response1 (fixed) $k_{0,HTN}$ Biophase rate constant for hypertensive effect (ng/mL)0.76140.30.8120.482-1.25 Y_{HTN} Hill coefficient for hypertensive effect1.9920.71.991.50-2.60 $\eta_{LECS0,NE}$ Inter-individual variability on EC _{S0,DEX} 0.54166.80.6350.644-0.979 $\sigma_{ADD,SC}$ Additive residual error for intravenous dosing6.308.76.325.61-7.04 $\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing8.732.78.828.26-9.39HR $k_{cl,HR_{NE}}$ Biophase Rate constant for NE in HR effect (h ⁻¹)6.1525.76.283.68-8.97 $E_{0,LR}$ Baseline HR effect (beats/min)62.04.1762.252.26-3.29HRHill coefficient or NE in HR effect (h ⁻¹)6.1525.76.283.69-2.65.2 $K_{0,LR}$ Biophase NE at 50% of the maximal HR (nM)3.1225.93.202.26-4.289 Y_{NE} Hill coefficient for NE effect2.852.262.791.879-3.73 $E_{0,LR}$ Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $\kappa_{0,NR}$ Baseline RR at physi	EC _{50 NE}	Biophase NE for causing 50% maximal DAP (nM)	4.32	23.5	4.54	3.61-5.64
The second se	γ _{HPN}	Hill coefficient for hypotensive effect	4.99	25.2	5.07	3.27-7.16
MARK Baseline DAP (mmHg)83.55.783.479.6–87.1 E_{0DAP} Maximum DAP response1 (fixed) k_{0HTN} Biophase rate constant for hypertensive effect (n^{-1})30 (fixed) $EC_{50,DEX}$ Biophase DEX for 50% hypertensive effect (ng/L)0.76140.30.8120.482–1.25 Y_{HTN} Hill coefficient for hypertensive effect1.9920.71.991.50–2.60 $n_{LECS0,NE}$ Inter-individual variability on $EC_{50,DEX}$ 0.54166.80.6350.364–0.979 $\sigma_{ADD,IV}$ Additive residual error for intravenous dosing6.308.76.325.61–7.04 $\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing8.732.78.828.26–9.39HR6.1525.76.283.68–8.97 $E_{0,HR}$ Biophase rate constant for NE in HR effect (h^{-1})6.1525.76.283.68–8.972.6–4.289 $K_{0,HR_{ME}}$ Biophase Rate soft from the effect2.8522.62.791.879–3.732.7 $E_{0,SNE}$ Biophase NE at 50% of the maximal HR (nM)3.1225.93.202.26–4.289 γ_{NE} Hill coefficient for NE effect2.8522.62.791.879–3.73 $E_{CS_{0,SNF}}$ Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{0,NR}$ Baseline NR at physiological SAP levels0.0001 (fixed)	E _{MIN DAP}	DAP at 100% NE inhibition (mmHg)	30.2	2.1	30.3	29.5-31.2
b_{MAX} Maximum DAP response1 (fixed) $k_{e0,HTN}$ Biophase rate constant for hypertensive effect (n^{-1})30 (fixed) $EC_{50,DEX}$ Biophase DEX for 50% hypertensive effect (ng/mL)0.76140.30.8120.482-1.25 γ_{HTN} Hill coefficient for hypertensive effect1.9920.71.991.50-2.60 $\eta_{1,EC50,NE}$ Inter-individual variability on $EC_{50,NE}$ 0.38265.50.4340.295-0.633 $\eta_{1,EC50,DEX}$ Inter-individual variability on $EC_{50,DEX}$ 0.54166.80.6350.364-0.979 $\sigma_{ADD,IV}$ Additive residual error for intravenous dosing6.308.76.325.61-7.04 $\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing8.732.78.828.26-9.39HR $k_{e0,HR_{NE}}$ Biophase rate constant for NE in HR effect (h^{-1})6.1525.76.283.68-8.97 $E_{0,HR}$ Baseline HR effect (beats/min)62.04.1762.259.2-65.2EC $_{50,NE}$ Biophase NE at 50% of the maximal HR (nM)3.1225.93.202.26-4.289 γ_{NE} Hill coefficient for NE effect2.8522.62.791.879-3.73 $E_{0,SR}$ Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{0,NR}$ Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{0,NR}$ Biophase rate constant for NR in HR effect (h^{-1})0.4387.390	$E_{0 \text{ DAP}}$	Baseline DAP (mmHg)	83.5	5.7	83.4	79.6-87.1
MAX $\epsilon_{0,HTN}$ Biophase rate constant for hypertensive effect (h ⁻¹)30 (fixed)EC_{S0,DEX}Biophase DEX for 50% hypertensive effect (ng/mL)0.76140.30.8120.482-1.25 γ_{HTN} Hill coefficient for hypertensive effect1.9920.71.991.50-2.60 $\eta_{1,EC50,NE}$ Inter-individual variability on EC _{50,NE} 0.38265.50.4340.295-0.633 $\eta_{2,EC50,DEX}$ Inter-individual variability on EC _{50,DEX} 0.54166.80.6350.364-0.979 $\sigma_{ADD,NC}$ Additive residual error for intravenous dosing6.308.76.325.61-7.04 $\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing8.732.78.828.26-9.39HR $k_{e0,HR_{NE}}$ Biophase rate constant for NE in HR effect (h ⁻¹)6.1525.76.283.68-8.97 $E_{0,HR}$ Baseline HR effect (beats/min)62.04.1762.259.2-65.2EC_{50,NE}Biophase NE at 50% of the maximal HR (nM)3.1225.93.202.26-4.289 γ_{NE} Hill coefficient for NE effect2.8522.62.791.879-3.73 $E_{0,IRR}$ Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396-0.478 $E_{50,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396-0.478 $E_{0,NR}$ Biophase rate constant for NR in HR effect (h ⁻	E _{MAX}	Maximum DAP response	1 (fixed)	-	_	_
The big base of the field o	k _{e0 HTN}	Biophase rate constant for hypertensive effect (h^{-1})	30 (fixed)	-	_	_
JHTNHil coefficient for hypertensive effect1.9920.71.991.50–2.60 $\eta_{LECS0NE}$ Inter-individual variability on EC _{50.NE} 0.38265.50.4340.295–0.633 $\eta_{2ECS0.DEX}$ Inter-individual variability on EC _{50.DEX} 0.54166.80.6350.364–0.979 $\sigma_{ADD,IV}$ Additive residual error for intravenous dosing6.308.76.325.61–7.04 $\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing8.732.78.828.26–9.39HR $k_{e0,HR_{NE}}$ Biophase rate constant for NE in HR effect (h ⁻¹)6.1525.76.283.68–8.97 $E_{0,HR}$ Baseline HR effect (beats/min)62.04.1762.259.2–65.2EC _{50,NE} Biophase NE at 50% of the maximal HR (nM)3.1225.93.202.26–4.289 γ_{NE} Hill coefficient for NE effect2.8522.62.791.879–3.73 $E_{50,SAP}$ Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{0,NR}$ Baseline NR at physiological SAP levels0.0001 (fixed) $k_{0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4286.700.4780.396–0.482 $EC_{50,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396–0.482 $EC_{50,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396–0.482 $EC_{50,NR}$ NR at 50% of the maximal HR0.0001 (fixed)	EC _{50 DEX}	Biophase DEX for 50% hypertensive effect (ng/mL)	0.761	40.3	0.812	0.482-1.25
HIR $\eta_{1,EC50,NE}$ Inter-individual variability on EC_{50,DEX}0.38265.50.4340.295-0.633 $\eta_{2,EC50,DEX}$ Inter-individual variability on EC_{50,DEX}0.54166.80.6350.364-0.979 $\sigma_{ADD,IV}$ Additive residual error for intravenous dosing6.308.76.325.61-7.04 $\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing8.732.78.828.26-9.39HR $k_{e0,HR_{SE}}$ Biophase rate constant for NE in HR effect (h ⁻¹)6.1525.76.283.68-8.97 $E_{0,HR}$ Baseline HR effect (beats/min)62.04.1762.259.2-65.2EC_{50,NE}Biophase NE at 50% of the maximal HR (nM)3.1225.93.202.26-4.289 γ_{NE} Hill coefficient for NE effect2.8522.62.791.879-3.73 $E_{MIN,HR}$ HR at 100% NE inhibition (beats/min)24.281.4324.323.7-24.9EC_{50,SAP}Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{0,NR}$ Baseline NR at physiological SAP levels0.0001 (fixed) $k_{0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396-0.482 $E_{0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396-0.482 $E_{0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396-0.482 $E_{0,NR}$ Biophase rate constant for NR in HR effect (h	Y _{HTN}	Hill coefficient for hypertensive effect	1.99	20.7	1.99	1.50-2.60
The stand $\eta_{2,ECS0,DEX}$ Inter-individual variability on EC $s_{0,DEX}$ 0.54166.80.6350.364-0.979 $\sigma_{ADD,IV}$ Additive residual error for intravenous dosing6.308.76.325.61-7.04 $\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing8.732.78.828.26-9.39HR </td <td>$\eta_{1 EC50 NE}$</td> <td>Inter-individual variability on EC_{50 NE}</td> <td>0.382</td> <td>65.5</td> <td>0.434</td> <td>0.295-0.633</td>	$\eta_{1 EC50 NE}$	Inter-individual variability on EC _{50 NE}	0.382	65.5	0.434	0.295-0.633
$\sigma_{ADD,IV}$ Additive residual error for intravenous dosing6.308.76.325.61–7.04 $\sigma_{ADD,SC}$ Additive residual error for intravenous dosing8.732.78.828.26–9.39HR $k_{e0,HR_{NE}}$ Biophase rate constant for NE in HR effect (h ⁻¹)6.1525.76.283.68–8.97 $E_{0,HR}$ Baseline HR effect (beats/min)62.04.1762.259.2–65.2 $E_{S0,NE}$ Biophase NE at 50% of the maximal HR (nM)3.1225.93.202.26–4.289 γ_{NE} Hill coefficient for NE effect2.8522.62.791.879–3.73 $E_{MIN,HR}$ HR at 100% NE inhibition (beats/min)24.281.4324.323.7–24.9 $E_{S0,SAP}$ Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{oUT,NR}$ Dissociation rate constant for NR (h ⁻¹)0.4286.700.4780.396–0.478 $E_{0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396–0.482 $E_{0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396–0.482 $E_{0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396–0.482 $E_{0,NR}$ Hill coefficient for NR effect on HR12.456.711.75.92–20.7 $\eta_{1,ECS0,NR}$ Inter-individual variability on EC _{50,CNS} 0.49740.80.6660.135–0.666 σ_1 Additive residual error for intravenous dosing5.4615.0<	$\eta_{2 EC50 DEX}$	Inter-individual variability on $EC_{50,DEX}$	0.541	66.8	0.635	0.364-0.979
$\sigma_{ADD,SC}$ Additive residual error for subcutaneous dosing8.732.78.828.26–9.39HR $k_{e0,HR_{NE}}$ Biophase rate constant for NE in HR effect (h ⁻¹)6.1525.76.283.68–8.97 $E_{0,HR}$ Baseline HR effect (beats/min)62.04.1762.259.2–65.2EC_{50,NE}Biophase NE at 50% of the maximal HR (nM)3.1225.93.202.26–4.289 γ_{NE} Hill coefficient for NE effect2.8522.62.791.879–3.73 $E_{MIN,HR}$ HR at 100% NE inhibition (beats/min)24.281.4324.323.7–24.9EC_{50,SAP}Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{oUT,NR}$ Dissociation rate constant for NR (h ⁻¹)0.4286.700.4780.396–0.478 $E_{0,NR}$ Baseline NR at physiological SAP levels0.0001 (fixed) $k_{e0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396–0.482EC_{50,NR}NR at 50% of the maximal HR0.0001 (fixed) γ_{NR} Hill coefficient for NR effect on HR12.456.711.75.92–20.7 $\eta_{1,EC50,NR}$ Inter-individual variability on EC _{50,CNS} 0.49740.80.66660.135–0.666 σ_1 Additive residual error for subcutaneous dosing5.4615.06.564.88–6.56 σ_1 Additive residual error for subcutaneous dosing5.5113.66.294.89–6.29 <td>$\sigma_{ADD IV}$</td> <td>Additive residual error for intravenous dosing</td> <td>6.30</td> <td>8.7</td> <td>6.32</td> <td>5.61-7.04</td>	$\sigma_{ADD IV}$	Additive residual error for intravenous dosing	6.30	8.7	6.32	5.61-7.04
HR End of the term of the term of the term of the term of term	$\sigma_{ADD,SC}$	Additive residual error for subcutaneous dosing	8.73	2.7	8.82	8.26-9.39
$k_{e0,HR_{NE}}$ Biophase rate constant for NE in HR effect (h^{-1})6.1525.76.283.68-8.97 $E_{0,HR}$ Baseline HR effect (beats/min)62.04.1762.259.2-65.2EC_{50,NE}Biophase NE at 50% of the maximal HR (nM)3.1225.93.202.26-4.289 γ_{NE} Hill coefficient for NE effect2.8522.62.791.879-3.73 $E_{MIN,HR}$ HR at 100% NE inhibition (beats/min)24.281.4324.323.7-24.9EC_{50,SAP}Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{OUT,NR}$ Dissociation rate constant for NR (h^{-1})0.4286.700.4780.396-0.478 $E_{0,NR}$ Baseline NR at physiological SAP levels0.0001 (fixed) $k_{e0,NR}$ Biophase rate constant for NR in HR effect (h^{-1})0.4387.390.4820.396-0.482EC_{50,NR}NR at 50% of the maximal HR0.0001 (fixed) γ_{NR} Hill coefficient for NR effect on HR12.456.711.75.92-20.7 $\eta_{1,EC50,NR}$ Inter-individual variability on EC_{50,CNS}0.49740.80.6660.135-0.666 σ_1 Additive residual error for intravenous dosing5.4615.06.564.88-6.56 σ_1 Additive residual error for subcutaneous dosing5.5113.66.294.89-6.29	HR	č				
$E_{0,HR}$ Baseline HR effect (beats/min)62.04.1762.259.2-65.2 $E_{0,HR}$ Biophase NE at 50% of the maximal HR (nM)3.1225.93.202.26-4.289 γ_{NE} Hill coefficient for NE effect2.8522.62.791.879-3.73 $E_{MIN,HR}$ HR at 100% NE inhibition (beats/min)24.281.4324.323.7-24.9 $EC_{50,SAP}$ Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{OUT,NR}$ Dissociation rate constant for NR (h ⁻¹)0.4286.700.4780.396-0.478 $E_{0,NR}$ Baseline NR at physiological SAP levels0.0001 (fixed) $k_{e0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396-0.482 $EC_{50,NR}$ NR at 50% of the maximal HR0.0001 (fixed) γ_{NR} Hill coefficient for NR effect on HR12.456.711.75.92-20.7 $\eta_{1,EC50,NR}$ Inter-individual variability on $EC_{50,CNS}$ 0.49740.80.6660.135-0.666 σ_1 Additive residual error for intravenous dosing5.4615.06.564.88-6.56 σ_1 Additive residual error for subcutaneous dosing5.5113.66.294.89-6.29	k _{e0 HR}	Biophase rate constant for NE in HR effect (h^{-1})	6.15	25.7	6.28	3.68-8.97
EC 50.NEBiophase NE at 50% of the maximal HR (nM) 3.12 25.9 3.20 $2.26-4.289$ γ_{NE} Hill coefficient for NE effect 2.85 22.6 2.79 $1.879-3.73$ $E_{MIN,HR}$ HR at 100% NE inhibition (beats/min) 24.28 1.43 24.3 $23.7-24.9$ EC $50,SAP$ Biophase SAP at 50% NR inhibition (mmHg) 0.0001 (fixed) $ k_{OUT,NR}$ Dissociation rate constant for NR (h ⁻¹) 0.428 6.70 0.478 $0.396-0.478$ $E_{0,NR}$ Baseline NR at physiological SAP levels 0.0001 (fixed) $ k_{e0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹) 0.438 7.39 0.482 $0.396-0.478$ EC $50,NR$ NR at 50% of the maximal HR 0.0001 (fixed) $ \gamma_{NR}$ Hill coefficient for NR effect on HR 12.4 56.7 11.7 $5.92-20.7$ $\eta_{1,EC50,NR}$ Inter-individual variability on EC $50,CNS$ 0.497 40.8 0.666 $0.135-0.666$ σ_1 Additive residual error for intravenous dosing 5.46 15.0 6.56 $4.88-6.56$ σ_1 Additive residual error for subcutaneous dosing 5.51 13.6 6.29 $4.89-6.29$	$E_{0 \text{ HR}}$	Baseline HR effect (beats/min)	62.0	4.17	62.2	59.2-65.2
$\gamma_{\rm NE}$ Hill coefficient for NE effect2.8522.62.791.879–3.73 $E_{\rm MIN,HR}$ HR at 100% NE inhibition (beats/min)24.281.4324.323.7–24.9EC_{50,SAP}Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{\rm OUT,NR}$ Dissociation rate constant for NR (h ⁻¹)0.4286.700.4780.396–0.478 $E_{0,NR}$ Baseline NR at physiological SAP levels0.0001 (fixed) $k_{e0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396–0.478EC_{50,NR}NR at 50% of the maximal HR0.0001 (fixed) γ_{NR} Hill coefficient for NR effect on HR12.456.711.75.92–20.7 $\eta_{1,EC50,NR}$ Inter-individual variability on EC_{50,CNS}0.49740.80.6660.135–0.666 σ_1 Additive residual error for intravenous dosing5.4615.06.564.88–6.56 σ_1 Additive residual error for subcutaneous dosing5.5113.66.294.89–6.29	EC _{50 NE}	Biophase NE at 50% of the maximal HR (nM)	3.12	25.9	3.20	2.26-4.289
HRHR at 100% NE inhibition (beats/min)24.281.4324.323.7-24.9 $E_{50,SAP}$ Biophase SAP at 50% NR inhibition (mmHg)0.0001 (fixed) $k_{OUT,NR}$ Dissociation rate constant for NR (h ⁻¹)0.4286.700.4780.396-0.478 $E_{0,NR}$ Baseline NR at physiological SAP levels0.0001 (fixed) $k_{e0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396-0.482 $E_{50,NR}$ NR at 50% of the maximal HR0.0001 (fixed) γ_{NR} Hill coefficient for NR effect on HR12.456.711.75.92-20.7 $\eta_{1,EC50,NR}$ Inter-individual variability on EC _{50,CNS} 0.49740.80.6660.135-0.666 σ_1 Additive residual error for intravenous dosing5.4615.06.564.88-6.56 σ_1 Additive residual error for subcutaneous dosing5.5113.66.294.89-6.29	y _{NE}	Hill coefficient for NE effect	2.85	22.6	2.79	1.879-3.73
EC 50.SAPBiophase SAP at 50% NR inhibition (mmHg) 0.0001 (fixed) $ k_{OUT,NR}$ Dissociation rate constant for NR (h ⁻¹) 0.428 6.70 0.478 $0.396-0.478$ $E_{0,NR}$ Baseline NR at physiological SAP levels 0.0001 (fixed) $ k_{e0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹) 0.438 7.39 0.482 $0.396-0.482$ EC $50,NR$ NR at 50% of the maximal HR 0.0001 (fixed) $ \gamma_{NR}$ Hill coefficient for NR effect on HR 12.4 56.7 11.7 $5.92-20.7$ $\eta_{1,EC50,NR}$ Inter-individual variability on EC $50,CNS$ 0.497 40.8 0.666 $0.135-0.666$ σ_1 Additive residual error for intravenous dosing 5.46 15.0 6.56 $4.88-6.56$ σ_1 Additive residual error for subcutaneous dosing 5.51 13.6 6.29 $4.89-6.29$	E _{MIN HP}	HR at 100% NE inhibition (beats/min)	24.28	1.43	24.3	23.7-24.9
$k_{OUT,NR}$ Dissociation rate constant for NR (h ⁻¹)0.4286.700.4780.396-0.478 $E_{0,NR}$ Baseline NR at physiological SAP levels0.0001 (fixed) $k_{e0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹)0.4387.390.4820.396-0.478 $EC_{50,NR}$ NR at 50% of the maximal HR0.0001 (fixed) γ_{NR} Hill coefficient for NR effect on HR12.456.711.75.92-20.7 $\eta_{1,EC50,NR}$ Inter-individual variability on EC _{50,CNS} 0.49740.80.66660.135-0.666 σ_1 Additive residual error for intravenous dosing5.4615.06.564.88-6.56 σ_1 Additive residual error for subcutaneous dosing5.5113.66.294.89-6.29	EC _{50 SAP}	Biophase SAP at 50% NR inhibition (mmHg)	0.0001 (fixed)	_	_	_
$E_{0,NR}$ Baseline NR at physiological SAP levels 0.0001 (fixed) $ k_{e0,NR}$ Biophase rate constant for NR in HR effect (h ⁻¹) 0.438 7.39 0.482 $0.396-0.482$ EC _{50,NR} NR at 50% of the maximal HR 0.0001 (fixed) $ \gamma_{NR}$ Hill coefficient for NR effect on HR 12.4 56.7 11.7 $5.92-20.7$ $\eta_{1,EC50,NR}$ Inter-individual variability on EC _{50,CNS} 0.497 40.8 0.666 $0.135-0.666$ σ_1 Additive residual error for intravenous dosing 5.46 15.0 6.56 $4.88-6.56$ σ_1 Additive residual error for subcutaneous dosing 5.51 13.6 6.29 $4.89-6.29$	k _{OUT NR}	Dissociation rate constant for NR (h^{-1})	0.428	6.70	0.478	0.396-0.478
0.1K Biophase rate constant for NR in HR effect (h ⁻¹) 0.438 7.39 0.482 $0.396-0.482$ EC _{50,NR} NR at 50% of the maximal HR 0.0001 (fixed) $ \gamma_{\text{NR}}$ Hill coefficient for NR effect on HR 12.4 56.7 11.7 $5.92-20.7$ $\eta_{1,\text{EC50,NR}}$ Inter-individual variability on EC _{50,CNS} 0.497 40.8 0.6666 $0.135-0.666$ σ_1 Additive residual error for intravenous dosing 5.46 15.0 6.56 $4.88-6.56$ σ_1 Additive residual error for subcutaneous dosing 5.51 13.6 6.29 $4.89-6.29$	$E_{0 \text{ NP}}$	Baseline NR at physiological SAP levels	0.0001 (fixed)	_	_	_
EC.50,NRNR at 50% of the maximal HR 0.0001 (fixed) $ \gamma_{NR}$ Hill coefficient for NR effect on HR 12.4 56.7 11.7 $5.92-20.7$ $\eta_{1,EC50,NR}$ Inter-individual variability on EC _{50,CNS} 0.497 40.8 0.666 $0.135-0.666$ σ_1 Additive residual error for intravenous dosing 5.46 15.0 6.56 $4.88-6.56$ σ_1 Additive residual error for subcutaneous dosing 5.51 13.6 6.29 $4.89-6.29$	k _{o0 NIP}	Biophase rate constant for NR in HR effect (h^{-1})	0.438	7.39	0.482	0.396-0.482
$\gamma_{\rm NR}$ Hill coefficient for NR effect on HR12.456.711.75.92–20.7 $\eta_{1,\rm EC50,\rm NR}$ Inter-individual variability on EC _{50,CNS} 0.49740.80.6660.135–0.666 σ_1 Additive residual error for intravenous dosing5.4615.06.564.88–6.56 σ_1 Additive residual error for subcutaneous dosing5.5113.66.294.89–6.29	EC _{50 MD}	NR at 50% of the maximal HR	0.0001 (fixed)	_	_	_
$\eta_{1,EC50,NR}$ Inter-individual variability on EC_{50,CNS}0.49740.80.6660.135-0.666 σ_1 Additive residual error for intravenous dosing5.4615.06.564.88-6.56 σ_1 Additive residual error for subcutaneous dosing5.5113.66.294.89-6.29	- 30,ΝΚ γ _{NID}	Hill coefficient for NR effect on HR	12.4	56.7	11.7	5.92-20.7
σ_1 Additive residual error for intravenous dosing5.4615.06.564.88-6.56 σ_1 Additive residual error for subcutaneous dosing5.5113.66.294.89-6.29	$\eta_{1 \text{EC50 ND}}$	Inter-individual variability on $EC_{50,CNS}$	0.497	40.8	0.666	0.135-0.666
σ_1 Additive residual error for subcutaneous dosing 5.51 13.6 6.29 4.89–6.29	σ_1	Additive residual error for intravenous dosing	5.46	15.0	6.56	4.88-6.56
	σ_1	Additive residual error for subcutaneous dosing	5.51	13.6	6.29	4.89-6.29

Median and 95% confidence intervals (CIs) from the sampling importance resampling procedure with 40,000 final samples and 2000 resamples. The subjects were given a single 1-µg/kg dexmedetomidine infusion in two phases either intravenously or subcutaneously in 10 min

DAP diastolic arterial blood pressure, DEX dexmedetomidine, HPN hypotensive effect of DEX, HTN hypertensive effect of DEX, NE norepinephrine, NR neural reflex from central nervous system, RSE relative standard error, SAP systolic arterial blood pressure, SIR sampling importance resampling

10.0

10.0

10.0



Fig. 3 Visual predictive checks based on 1000 simulations showing heart rate (upper row), systolic blood pressure (middle row) and diastolic blood pressure (lower row) after a 1- μ g/kg intravenous (**a**, **c**, **e**) or subcutaneous (**b**, **d**, **f**) dexmedetomidine infusion. The black solid

and dashed lines are the observed percentiles (10th, 50th, and 90th percentiles), respectively, and the blue ribbon is the corresponding median predictive interval. Light blue ribbons are predicted percentiles. Black circles are individual observations



Fig. 4 Simulated dexmedetomidine concentration profiles in plasma using the final mechanism-based model and 8-h (upper row) and 48-h (lower row) intravenous (a, c) and subcutaneous (b, d) continuous infusions of varying dexmedetomidine doses

estimates to ease up the estimation routines of the final models, considering that dexmedetomidine-induced vasoconstriction is a documented phenomenon [13]. Interestingly, the EC_{50,DEX} of dexmedetomidine for SAP and DAP was noticed to be considerably different, at 2.37 ng/mL and 0.81 ng/mL, respectively, while $\gamma_{\rm HTN}$ for SAP and DAP was 1.52 and 1.99, respectively, indicating that the hypertensive effect of dexmedetomidine on blood pressure is less pronounced compared with the hypotensive effect mediated by NE inhibition.

3.5 Heart Rate Model

Bradycardia after dexmedetomidine dosing was initially modelled with the sympatholytic effect of NE release inhibition only, as previously documented [9]. Inhibition of NE release was able to account for the general trend in the HR data, but a clear model misspecification was observable in the model predictions, especially after SC dexmedetomidine administration. We added an inhibitory effect of SAP on HR, mediated by a baroreceptor reflex in the CNS, to account for the misfit in the HR model, which resolved the model misspecification and resulted in an unbiased output.

The effect of SAP on CNS neural activity was specified using an indirect response model as follows:

$$k_{\text{in,CNS}} = k_{\text{in,CNS}} t^{t=0} \bullet \left(1 - \left(\frac{\text{IPRED}_{\text{SAP}}}{\text{EC}_{50,\text{SAP}} + \text{IPRED}_{\text{SAP}}} \right) \right)$$

where $k_{in,CNS}$ is the physiological rate of development for CNS neural activity, $k_{in,CNS}^{t=0}$ are the initial conditions for the neural reflex at time = 0, IPRED_{SAP} is the individual predicted SAP levels from the final model, and EC_{50,SAP} is

the level of SAP required to cause a 50% change in central neural activity.

We assumed that the overall HR effect caused by dexmedetomidine is the sum of the HR-modulating reflex ($E_{\rm HR,CNS}$) and sympatholytic ($E_{\rm HR,NE}$) mechanisms as pointed out recently [13]. The former was considered to result from an initial vasoconstrictive hypertension after dexmedetomidine administration, which thereby modulates central neural activity through the baroreceptor reflex, causing a diminished HR effect. The latter driver of the HR effect was coded using NE as a surrogate measure, while reduced NE release and resulting sympatholysis were anticipated to cause bradycardia.

Hence, the two components of the overall effect were estimated simultaneously in the final HR model as follows:

$$E_{\text{HR,CNS}} = 1 - \left(\frac{E_{\text{MAX,HR}} \bullet C_{\text{E,CNS}}^{\gamma_{\text{NR}}}}{\text{EC}_{50,\text{CNS}}^{\gamma_{\text{NR}}} + C_{\text{E,CNS}}^{\gamma_{\text{NR}}}}\right),$$
$$E_{\text{HR,NE}} = 1 + \left(\frac{E_{\text{MAX,HR}} \bullet C_{\text{E,NE}}^{\gamma_{\text{NE}}}}{\text{EC}_{50,\text{NE}}^{\gamma_{\text{NE}}} + C_{\text{E,NE}}^{\gamma_{\text{NE}}}}\right),$$

 $E_{\rm HR} = E_{\rm MIN, HR} \bullet \left(E_{\rm HR, CNS} + E_{\rm HR, NE} \right).$

This approach led to a numerically stable model and resulted in physiologically plausible parameter estimates and corrected the previous misspecification in the HR model (Table 3). Norepinephrine EC_{50,NE} for HR effect was estimated at 3.12 nM, while $\gamma_{\rm NE}$ was 2.85, showing a steep concentration–response relationship for NE on HR development. $k_{\rm e0,HR_{NE}}$ for NE at the HR effect compartment was 6.15 h⁻¹, suggesting that the effect is slower as compared with the NE on SAP/DAP but has a longer half-life.

Additionally, $k_{e0,NR}$ for the effect of neural reflex on HR was 0.438 h⁻¹, which demonstrates a long half-life for this process, which is biologically plausible. The final parameter estimate for γ_{NR} is 12.4, indicating that minute changes in neural responses can have an almost all-or-none effect on HR.

3.6 Simulations

Our simulations with the final PK model indicate that both IV and SC administration of dexmedetomidine doses of $0.5-1 \mu g/kg$ resulted in previously reported therapeutic plasma concentrations [7]. The NCA indicated that the model-predicted peak concentrations and AUCs were consistent with previously reported results (ESM) [6]. The time to achieve steady-state plasma concentrations with a constant-rate SC infusion was 15–20 h, which is considerably longer compared with 3 h after IV dosing (Fig. 4).

Our model results show that SC administration of dexmedetomidine results in a milder inhibition of NE release from the release compartment at the same dose concentrations, which translates into an attenuation of an otherwise pronounced haemodynamic effect with the IV route. A dose concentrations of 4 µg/kg/h or above can result in lowering NE levels in the release compartment below 0.1 nM, leading to very low circulating NE levels, both in IV as well as SC administration (Fig. 5). The simulations with our final SAP and DAP models show a dynamic interplay between the hypotensive and hypertensive effects, depending upon the concentrations of NE and dexmedetomidine. Subcutaneous dosing of 0.25-1 µg/kg/h does not evoke notable hypertensive effects, and a unanimously downward trend can be seen in blood pressure simulations at these dose concentrations. After IV dosing, initial high dexmedetomidine concentrations result in a short hypertensive effect $(E_{\text{BP,HTN}})$, which is swiftly followed by a hypotensive effect $(E_{\text{BP HPN}})$ due to reduced NE release (ESM). Furthermore, our simulations with IV doses suggest that the hypertensive effect is minimal at lower doses but increases considerably with increasing doses and surpasses the hypotensive effect at dose concentrations above 2 µg/kg. However, because of the rapid disappearance of the hypertensive effect of dexmedetomidine, the hypotensive response due to reduced NE release causes blood pressure to fall (ESM).

The addition of an effect from changes in central neural activity following SAP perturbations helped to simulate the dynamic interplay between these effect components on HR over a range of doses and dosing schemes and our simulations suggest a dose-dependent decline. The effects are clearly seen to be more pronounced for the IV route over the SC route and the slope of decline after IV administration is much steeper as compared with SC administration, owing to high initial concentrations of dexmedetomidine (ESM).

4 Discussion

Our primary aim was to develop semi-mechanistic models to characterise dexmedetomidine concentration– and response–time courses after SC and IV dosing. Using previously collected data and a sequential PK–PD modelling approach, we were able to describe dexmedetomidine absorption kinetics and further disposition together with its sympatholytic and haemodynamic effects. Our models were able to capture the data trends with relatively high precision, and the model predictions show good agreement with the actual observations. Internal evaluation confirmed the predictive performance of the models and the accuracy of the final model parameter estimates, with narrow confidence intervals.



Fig. 5 Simulated norepinephrine concentration profiles in plasma using the final mechanism-based model and 8-h (upper row) and 48-h (lower row) intravenous (\mathbf{a}, \mathbf{c}) and subcutaneous (\mathbf{b}, \mathbf{d}) continuous infusions of varying dexmedetomidine doses

A first-order absorption model has previously been used for nasal dexmedetomidine administration [9], but it did not accurately explain the time course and trends in dexmedetomidine absorption after SC dosing. The original data indicated fast absorption after SC dosing, followed by a long equilibration phase. A wide range of absorption models was tested, but a more mechanistic solution was needed. Dexmedetomidine is highly lipid soluble [29], which pointed towards developing an absorption model with an additional SC fat compartment describing a slower absorption process. As the SC fat compartment is perfused with its own blood supply, we assumed direct absorption from the fat layer into peripheral circulation with a slow absorption rate constant. A redistribution model from the SC fat layer into the injection depot before absorption into plasma was also tested. The former approach was chosen for being physiologically more plausible, and it was also supported by the data. The final absorption model showed an adequate fit for the data at hand. Furthermore, results from our NCA simulations indicate that our model predicted similar AUC as found previously [6].

We postulated that SC fat could act as a slow-release reservoir of dexmedetomidine during prolonged exposure. The results from model simulations indicate that at higher SC doses, the peak concentration in plasma is postponed, while the total systemic exposure remains nearly unchanged. Simulations also suggest that plasma concentrations remain higher for longer time after SC dosing compared with IV dosing. Our results propose that constant-rate infusions of 8 and 48 h and at dexmedetomidine dose rates between 0.25 and 0.5 μ g/kg/h would be adequate to reach previously suggested therapeutically effective plasma concentrations of 0.3–0.8 ng/mL [7] after both IV and SC administration. This implies that adequate palliative sedation might be achieved using low dosing rates and the convenience of an extravascular SC route of administration. However, it needs to be highlighted that according to our results, the time needed to achieve steady-state plasma concentrations was higher in SC (15–20 h) vs IV (5–10 h) administration, and this may be worth a consideration in clinical settings.

Our final PK model shows consistency with previous reports. A two-compartmental PK model was best suited to explain the disposition kinetics of the drug, which is in line with previous population-based modelling reports [9]. Some previous studies have used three-compartmental models, but our initial effort showed a two-compartment model more stable with more precise parameter estimates and smaller standard errors. Furthermore, parameter estimates from our final PK model agree with previous studies reporting elimination clearance estimates between 41 and 57 L/min [9, 30]. Interestingly, the estimate of clearance from our study with SC dosing is completely in line with a previous report using an intranasal route of administration (i.e. 52.6, 44.0 and 44.5 L/min after IV, intranasal and SC dosing, respectively) [9, 13].

The estimation of bioavailability parameters in our final model demonstrated that the fat compartment receives approximately 55% of the drug from the depot, and of the remaining 45%, a further 79% reaches the systemic circulation via fast absorption. The resulting net bioavailability of dexmedetomidine via the SC route value is 89% (F = 0.89), which is slightly higher than our previous estimation using NCA. Bioavailability of dexmedetomidine after intranasal, intramuscular, oral and buccal administration has been investigated previously [6, 31, 32]. Our current findings suggest approximately similar bioavailability of dexmedetomidine after SC and intranasal administration and confirm that the bioavailability is sufficient for clinical usefulness also after SC administration. Compared to intranasal dosing, SC dosing seems more appropriate for palliative care, as the elimination is much slower.

We measured NE with dense sampling to characterise the effect on sympathetic nervous system after dexmedetomidine dosing. We assumed that for modelling purposes, NE levels at the sites of release could be used as a surrogate biomarker of sympatholysis. An indirect response model [33] could capture the sympatholytic effect of dexmedetomidine on NE release with more precise model estimates, relative standard errors and narrower confidence intervals compared with previous results [9]. Our simulations suggest doses above 4 μ g/kg nearly abolish plasma NE levels, which may have clinical implications to be considered when using dexmedetomidine at high SC infusion rates. However, it should be emphasised that much lower doses are clinically effective, indicating that the SC dosing route is a safe alternative for palliative sedation.

Dexmedetomidine exerts biphasic effects on blood pressure [34–36]. At high initial concentrations and especially after IV dosing, a fast hypertensive effect predominates due to an action on vascular α_2 -adrenoceptors [37, 38]. Sympatholysis develops somewhat later when the drug distributes into the CNS to regulate sympathetic nervous activity [7, 36, 39]. Our model estimate for EC₅₀ of dexmedetomidine for causing a rise in blood pressure was 2.45 ng/mL. It is therefore understandable that the initial hypertensive response is much more prevalent after IV than after SC dosing as the former leads to sudden high plasma concentrations of dexmedetomidine. Considering fragile patients requiring palliative care may have concomitant cardiovascular problems, SC administration should thus attenuate the haemodynamic challenges associated with dexmedetomidine administration.

The role of NE in the modulation of HR after dexmedetomidine administration has been previously documented and also utilised in PD modelling of the HR response after IV dosing [9]. Yoo et al. studied extravascular dosing via intranasal administration and used an indirect response model to estimate NE effect on HR by employing plasma NE concentrations and the NE biophase [9], but high between-subject variability was seen. A recent study of Colin et al. used dexmedetomidine plasma concentrations as a surrogate marker for HR and showed that HR model estimates followed predicted dexmedetomidine plasma concentration linearly [11]. Their results indicate that the lower boundary for HR variability after IV dexmedetomidine is 22 beats/min, but again the inter-individual variability was quite high (48%). We also first tested an indirect response model to account for the NE effect on HR, but a clear misspecification was observed in the model predictions. Changes in blood pressure are known to result in modulation of the baroreceptor reflex and changes in its activity help to maintain haemodynamic homeostasis. The effect of the baroreflex attributable to dexmedetomidine pharmacokinetics was recently shown to account for initial vasoconstriction after dexmedetomidine dosing [13].

We hypothesised, based on this finding [13] and our results, that changes in central neural reflexes that drive the HR effect and depend on baroreflex activity are essential for accurate prediction of the HR time course. The HR model was significantly improved after including an effect component dependent on blood pressure levels, which mediates the effects of the central neural activity. Simulations with our final HR model indicate that the dexmedetomidine has a more pronounced effect on HR after intravascular dosing, and SC dosing seems to be also safer in this regard. Our results agree with a previous finding demonstrating that extravascular dosing may avoid strong acute haemodynamic changes seen after intravascular use.

Our study has several limitations. Although we aimed to implement mechanistic modelling solutions, many assumptions regarding underlying physiological processes had to be made during the development of our models for simplification. These assumptions override the deeply complex and multi-layered nature of the effects of dexmedetomidine on haemodynamic parameters and subjective effects. Additionally, we could not achieve a stable epinephrine model because of over-parameterisation issues because of the large amount of concentrations below the limit of quantification. However, our current models show reasonable performance with low standard errors and narrow confidence intervals. This indicates that although data from only ten subjects were studied, our dense sampling could provide enough information for model development. Finally, the data used in the modelling were collected from healthy volunteers, thus our findings should be translated to clinical practice only with appropriate caution.

5 Conclusions

Our semi-mechanistic models were able to capture the PK and PD profiles of both intravenously and subcutaneously administered dexmedetomidine, describe the inhibition of NE release, and evaluate the haemodynamic and subjective drug effects. The PK model provides a mechanistic exploration of the absorption process and disposition after SC dexmedetomidine with excellent precision. Endogenous mediators were utilised in an indirect response model for inhibition of NE release to control the diverse effects of the drug. Norepinephrine effects on HR, SAP and DAP were linked via a biophase compartment, which reflects the effects of endogenous mediators and their access to the heart and blood vessels. Our current model provides accurate and clinically plausible predictions, which can provide additional information for drug therapy used in palliative care.

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Compliance with Ethical Standards

Conflict of interest MWA, PU and TIS received a speaker honorarium from Orion Pharma Ltd (Espoo, Finland). MS had contract research relationships with Orion Corporation Ltd (Espoo, Finland), Novartis Pharma (Espoo, Finland), Desentum Ltd (Helsinki, Finland), Gabather AB (Gothenburg, Sweden), AbbVie AG (Germany) and AC Immune SA (Lausanne, Switzerland). In addition, MS has received fees for participation on Data and Safety Committees in Faron Pharmaceuticals (Turku, Finland) and Herantis Pharma (Helsinki, Finland). MS has received a speaker's fee from Pharma Industry Finland (Helsinki, Finland) and he has stock options for Santhera Pharmaceuticals (Pratteln, Switzerland). **Open Access** This article is licensed under a Creative Commons Attribution-NonCommercial 4.0 International License, which permits any non-commercial use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by-nc/4.0/.

References

- 1. Bodnar J. A review of agents for palliative sedation/continuous deep sedation: pharmacology and practical applications. J Pain Palliat Care Pharmacother. 2017;31:16–37.
- Garetto F, Cancelli F, Rossi R, Maltoni M. Palliative sedation for the terminally ill patient. CNS Drugs. 2018;32:951–61.
- Weerink MAS, Struys MMRF, Hannivoort LN, Barends CRM, Absalom AR, Colin P. Clinical pharmacokinetics and pharmacodynamics of dexmedetomidine. Clin Pharmacokinet. 2017;26:335–46.
- Nelson LE, Lu J, Guo T, Saper CB, Franks NP, Maze M. The alpha2-adrenoceptor agonist dexmedetomidine converges on an endogenous sleep-promoting pathway to exert its sedative effects. Anesthesiology. 2003;98:428–36.
- Mahmoud M, Mason KP. Dexmedetomidine: review, update, and future considerations of paediatric perioperative and periprocedural applications and limitations. Br J Anaesth. 2015;115:171–82.
- Uusalo P, Al-Ramahi D, Tilli I, Aantaa RA, Scheinin M, Saari TI. Subcutaneously administered dexmedetomidine is efficiently absorbed and is associated with attenuated cardiovascular effects in healthy volunteers. Eur J Clin Pharmacol. 2018;74:1047–54.
- Ebert TJ, Hall JE, Barney JA, Uhrich TD, Colinco MD. The effects of increasing plasma concentrations of dexmedetomidine in humans. Anesthesiology. 2000;93:382–94.
- Lodenius Å, Ebberyd A, Hårdemark Cedborg A, Hagel E, Mkrtchian S, Christensson E, et al. Sedation with dexmedetomidine or propofol impairs hypoxic control of breathing in healthy male volunteers: a nonblinded, randomized crossover study. Anesthesiology. 2016;125:700–15.
- Yoo H, Iirola T, Vilo S, Manner T, Aantaa R, Lahtinen M, et al. Mechanism-based population pharmacokinetic and pharmacodynamic modeling of intravenous and intranasal dexmedetomidine in healthy subjects. Eur J Clin Pharmacol. 2015;71:1197–207.
- Colin PJ, Hannivoort LN, Eleveld DJ, Reyntjens KMEM, Absalom AR, Vereecke HEM, et al. Dexmedetomidine pharmacokineticpharmacodynamic modelling in healthy volunteers: 1. Influence of arousal on bispectral index and sedation. Br J Anaesth. 2017;119:200–10.
- Colin PJ, Hannivoort LN, Eleveld DJ, Reyntjens KMEM, Absalom AR, Vereecke HEM, et al. Dexmedetomidine pharmacodynamics in healthy volunteers: 2. Haemodynamic profile. Br J Anaesth. 2017;119:211–20.
- Li A, Yuen VM, Goulay-Dufaÿ S, Sheng Y, Standing JF, Kwok PCL, et al. Pharmacokinetic and pharmacodynamic study of intranasal and intravenous dexmedetomidine. Br J Anaesth. 2018;120:960–8.

- 13. Talke P, Anderson BJ. Pharmacokinetics and pharmacodynamics of dexmedetomidine-induced vasoconstriction in healthy volunteers. Br J Clin Pharmacol. 2018;84:1364–72.
- Pérez-Guillé M-G, Toledo-López A, Rivera-Espinosa L, Alemon-Medina R, Murata C, Lares-Asseff I, et al. Population pharmacokinetics and pharmacodynamics of dexmedetomidine in children undergoing ambulatory surgery. Anesth Analg. 2018;127(3):716–23.
- Greenberg RG, Wu H, Laughon M, Capparelli E, Rowe S, Zimmerman KO, et al. Population pharmacokinetics of dexmedetomidine in infants. J Clin Pharmacol. 2017;57:1174–82.
- Liu H-C, Lian Q-Q, Wu F-F, Wang C-Y, Sun W, Zheng L-D, et al. Population pharmacokinetics of dexmedetomidine after short intravenous infusion in Chinese children. Eur J Drug Metab Pharmacokinet. 2017;42:201–11.
- Iirola T, Ihmsen H, Laitio R, Kentala E, Aantaa R, Kurvinen JP, et al. Population pharmacokinetics of dexmedetomidine during long-term sedation in intensive care patients. Br J Anaesth. 2012;108:460–8.
- Välitalo PA, Ahtola-Sätilä T, Wighton A, Sarapohja T, Pohjanjousi P, Garratt C. Population pharmacokinetics of dexmedetomidine in critically ill patients. Clin Drug Investig. 2013;33:579–87.
- Ji QC, Zhou JY, Gonzales RJ, Gage EM, El-Shourbagy TA. Simultaneous quantitation of dexmedetomidine and glucuronide metabolites (G-Dex-1 and G-Dex-2) in human plasma utilizing liquid chromatography with tandem mass spectrometric detection. Rapid Commun Mass Spectrom. 2004;18:1753–60.
- Beal SL, Sheiner LB, Boeckmann AJ, Bauer RJ. NONMEM 7.4.3 users guides (1989-2018). Hanover, MD: ICON Development Solutions; 2018.
- Lindbom L, Ribbing J, Jonsson EN. Perl-speaks-NONMEM (PsN): a Perl module for NONMEM related programming. Comput Methods Progr Biomed. 2004;75:85–94.
- R Core Team. The R Project for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. 2018. https://www.R-project.org/. Accessed 10 May 2020.
- 23. Frechen S, Junge L, Saari TI, Suleiman AA, Rokitta D, Neuvonen PJ, et al. A semiphysiological population pharmacokinetic model for dynamic inhibition of liver and gut wall cytochrome P450 3A by voriconazole. Clin Pharmacokinet. 2013;52:763–81.
- Ashraf MW, Peltoniemi MA, Olkkola KT, Neuvonen PJ, Saari TI. Semimechanistic population pharmacokinetic model to predict the drug-drug interaction between S-ketamine and ticlopidine in healthy human volunteers. CPT Pharmacomet Syst Pharmacol. 2018;7:687–97.
- Nguyen THT, Mouksassi M-S, Holford N, Al-Huniti N, Freedman I, Hooker AC, et al. Model evaluation of continuous data pharmacometric models: metrics and graphics. CPT Pharmacomet Syst Pharmacol. 2017;6:87–109.
- 26. Dosne A-G, Bergstrand M, Harling K, Karlsson MO. Improving the estimation of parameter uncertainty distributions in nonlinear

mixed effects models using sampling importance resampling. J Pharmacokinet Pharmacodyn. 2016;43:583–96.

- Acharya C, Hooker AC, Türkyılmaz GY, Jönsson S, Karlsson MO. A diagnostic tool for population models using non-compartmental analysis: the ncappc package for R. Comput Methods Progr Biomed. 2016;127:83–93.
- Linares OA, Jacquez JA, Zech LA, Smith MJ, Sanfield JA, Morrow LA, et al. Norepinephrine metabolism in humans: kinetic analysis and model. J Clin Investig. 1987;80:1332–411.
- European Medicines Agency: EMEA/H/C/002268 Dexdor: EPAR - Product Information. 2020. https://www.ema.europa.eu/ en/documents/product-information/dexdor-epar-product-infor mation_en.pdf. Accessed 10 May 2020.
- Flexman AM, Wong H, Riggs KW, Shih T, Garcia PA, Vacas S, et al. Enzyme-inducing anticonvulsants increase plasma clearance of dexmedetomidine: a pharmacokinetic and pharmacodynamic study. Anesthesiology. 2014;120:1118–25.
- Iirola T, Vilo S, Manner T, Aantaa R, Lahtinen M, Scheinin M, et al. Bioavailability of dexmedetomidine after intranasal administration. Eur J Clin Pharmacol. 2011;67:825–31.
- 32. Uusalo P, Guillaume S, Siren S, Manner T, Vilo S, Scheinin M, et al. Pharmacokinetics and sedative effects of intranasal dexmedetomidine in ambulatory pediatric patients. Anesth Analg. 2020;130:949–57.
- Dayneka NL, Garg V, Jusko WJ. Comparison of four basic models of indirect pharmacodynamic responses. J Pharmacokinet Biopharm. 1993;21:457–78.
- Bloor BC, Ward DS, Belleville JP, Maze M. Effects of intravenous dexmedetomidine in humans. II. Hemodynamic changes. Anesthesiology. 1992;77:1134–42.
- Dyck JB, Maze M, Haack C, Vuorilehto L, Shafer SL. The pharmacokinetics and hemodynamic effects of intravenous and intramuscular dexmedetomidine hydrochloride in adult human volunteers. Anesthesiology. 1993;78:813–20.
- 36. Kallio A, Scheinin M, Koulu M, Ponkilainen R, Ruskoaho H, Viinamäki O, et al. Effects of dexmedetomidine, a selective alpha 2-adrenoceptor agonist, on hemodynamic control mechanisms. Clin Pharmacol Ther. 1989;46:33–42.
- Link RE, Desai K, Hein L, Stevens ME, Chruscinski A, Bernstein D, et al. Cardiovascular regulation in mice lacking alpha2adrenergic receptor subtypes b and c. Science. 1996;273:803–5.
- Talke P, Stapelfeldt C, Lobo E, Brown R, Scheinin M, Snapir A. Effect of alpha2B-adrenoceptor polymorphism on peripheral vasoconstriction in healthy volunteers. Anesthesiology. 2005;102:536–42.
- Talke P, Lobo E, Brown R. Systemically administeredalpha2-agonist-induced peripheral vasoconstriction in humans. Anesthesiology. 2003;99:65–70.