

Citation for published version:

Loisios-Konstantinidis, I, Paraiso, RLM, Fotaki, N, McAllister, M, Cristofoletti, R & Dressman, J 2019, 'Application of the relationship between pharmacokinetics and pharmacodynamics in drug development and therapeutic equivalence: a PEARRL review', *Journal of Pharmacy and Pharmacology*, vol. 71, no. 4, pp. 699-723. https://doi.org/10.1111/jphp.13070

DOI: 10.1111/jphp.13070

Publication date: 2019

Document Version Peer reviewed version

Link to publication

This is the peer reviewed version of the following article: LoisiosKonstantinidis, I., Paraiso, R. L., Fotaki, N., McAllister, M., Cristofoletti, R. and Dressman, J. (2019), Application of the relationship between pharmacokinetics and pharmacodynamics in drug development and therapeutic equivalence: a PEARRL review. J Pharm Pharmacol, 71: 699-723., which has been published in final form at https://doi.org/10.1111/jphp.13070. This article may be used for non-commercial purposes in accordance with Wiley Terms and Conditions for Self-Archiving

University of Bath

General rights

Copyright and moral rights for the publications made accessible in the public portal are retained by the authors and/or other copyright owners and it is a condition of accessing publications that users recognise and abide by the legal requirements associated with these rights.

Take down policy

If you believe that this document breaches copyright please contact us providing details, and we will remove access to the work immediately and investigate your claim.

1 Application of the relationship between pharmacokinetics and pharmacodynamics in drug

- 2 *development and therapeutic equivalence: a PEARRL review*
- 3

4 Authors

5 Ioannis Loisios-Konstantinidis^a, Rafael L. M. Paraiso^a, Nikoletta Fotaki^b, Mark McAllister^c, Rodrigo Cristofoletti^d,

6 Jennifer Dressman^a

7 Author Information

- 8 ^a Institute of Pharmaceutical Technology, Goethe University, Frankfurt am Main, Germany, ^b Department of
- 9 Pharmacy and Pharmacology, Faculty of Science, University of Bath, Bath, UK, ^c Pfizer Drug Product Design,

10 Sandwich, UK and ^d Brazilian Health Surveillance Agency (ANVISA), Division of Therapeutic Equivalence, Brasilia,

11 Brazil

12 Correspondence

- 13 Jennifer Dressman, Biocenter, Institute of Pharmaceutical Technology, Johann Wolfgang Goethe University,
- 14 Max-von-Laue-Str. 9, Frankfurt am Main 60438, Germany. Email: dressman@em.uni-frankfurt.de
- 15
- 16

17 Abstract

- Objectives The objective of this review is to provide an overview of PK/PD models, focusing on drug-specific
 PK/PD models and highlighting their value-added in drug development and regulatory decision-making.
- 20 Key findings Many PK/PD models, with varying degrees of complexity and physiological understanding, have
- 21 been developed to evaluate the safety and efficacy of drug products. In special populations (e.g. pediatrics), in
- 22 cases where there is genetic polymorphism and in other instances where therapeutic outcomes are not well
- described solely by PK metrics, the implementation of PK/PD models is crucial to assure the desired clinical
 outcome. Since dissociation between the pharmacokinetic and pharmacodynamic profiles is often observed, it
- outcome. Since dissociation between the pharmacokinetic and pharmacodynamic profiles is often observed, it is proposed that physiologically-based pharmacokinetic (PBPK) and PK/PD models be given more weight by
- 26 regulatory authorities when assessing the therapeutic equivalence of drug products.
- 27 **Summary** Modeling and simulation approaches already play an important role in drug development. While slowly
- 28 moving away from "one-size fits all" PK methodologies to assess therapeutic outcomes, further work is required
- 29 to increase confidence in PK/PD models in translatability and prediction of various clinical scenarios to encourage
- 30 more widespread implementation in regulatory decision-making.
- 31
- 32
- 33 Keywords
- 34 Pharmacokinetics/ pharmacodynamics (PK/PD), modeling & simulation, drug development, regulatory science,
- 35 bioequivalence, therapeutic equivalence

Table of Contents

37	1	Introdu	ction		
38	2	The effe	ect compartment model	5	
39	2	.1 Ov	erview	5	
40	2	.2 Ap	plications and case examples	7	
41		2.2.1	d-tubocurarine and pancuronium	7	
42		2.2.2	Ibuprofen: dental pain relief	9	
43		2.2.3	Anti-nociceptive effect of morphine	14	
44	3	Modelir	ng of irreversible mechanisms of action	15	
45	3	.1 Ov	erview	15	
46	3	.2 Ap	plications and case examples	17	
47		3.2.1	Proton pump inhibitors	17	
48		3.2.2	Acetylsalicylic acid	22	
49		3.2.3	Exemestane	24	
50	4	Indirect	response and feedback control models	27	
51	4	.1 Ov	erview	27	
52		4.1.1	"Basic" and "extended basic" indirect response models	28	
53		4.1.2	Signal transduction and feedback control indirect response models	29	
54	4	.2 Ap	plications and case examples	31	
55		4.2.1	Ibuprofen: antipyretic response	31	
56		4.2.2	Rosuvastatin	34	
57		4.2.3	Escitalopram	37	
58	5	Outlook and concluding remarks			
59	6	Acknowledgements 42			
60	7	References			
61					

62 1 Introduction

63

Over the last decades pharmacokinetic/pharmacodynamics (PK/PD) models have been evolving rapidly, starting with the pioneering work in the 1960s, then moving from empirical descriptions to models based on mechanistic and physiological approaches and still evolving today in the form of state-of-the-art mathematical models describing the progression of diseases as well as entire biological systems, under the umbrella of systems pharmacology and computational biology. ^{[1],[2],[3],[4],[5],[6],[7]}

69 At the beginning of the conjunction of pharmacokinetics with pharmacodynamics, empirical models 70 which were based on the shape of the effect-concentration curve and assumed that the pharmacologic 71 response is directly related to the drug plasma concentration were introduced. Soon it was recognized 72 that this scenario is only valid when the equilibrium between the plasma and the site of action is 73 instantaneous, when the free drug concentration and the distribution to all tissues is the same (or 74 remains proportionally the same) and when the system is at steady-state. A variety of these so-called 75 steady-state empirical direct effect models have been reported in the literature: linear, power, 76 hyperbolic, sigmoid (E_{max} model), logarithmic and logistic. Even though these models have been applied in a number of situations, ^{[1],[8],[9]} they have two important limitations. First and most important, they 77 78 are time-independent (also referred to as static models). Second, they lack a mechanistic and/or 79 physiological understanding of the underlying pharmacokinetics and pharmacodynamics.^[10] For these 80 reasons, non-steady state, mechanistic and physiologically based modeling approaches were 81 introduced and these are more widely used these days in drug development.

In parallel to the developments in modeling approaches, major regulatory authorities have been moving slowly but surely from "one-size fits all" concepts to a more case-by-case, scientifically justified approach, in which the application of modeling and simulation (M&S) is playing a valuable supporting role. Physiologically-based pharmacokinetic (PBPK) and PK/PD models have already been implemented in the assessment of drug-drug interactions (DDIs) and extrapolation of results from adults to pediatric

populations. ^{[11],[12],[13],[14],[15],[16]} In addition, generic dermatologic and inhalation products have been
approved based on pharmacodynamic or clinical endpoint bioequivalence studies (BE).^{[17],[18]}

89 Most recently, pharmacokinetic metrics providing information about delivery of the drug to the body and exposure (i.e. onset and duration of action),^[19] such as partial areas under the concentration-time 90 91 curve (pAUCs) have been recommended by the US-FDA for the evaluation of several complex oral products combining immediate (IR) with extended release (ER). ^{[20],[21],[22]} However, there are still many 92 93 cases, especially for systematically acting drugs, where the value of modeling and simulation methods has not yet been widely recognized by the regulatory authorities. Such cases include the virtual 94 95 bioequivalence of oral drug products, the justification for potential extension of BCS-based biowaivers 96 to some BCS class II compounds and the reduction of the number of volunteers for bioequivalence 97 studies of highly variable drugs (HVDs). In view of the fact that single point pharmacokinetic metrics (i.e. C_{max}, AUC) used to assess bioequivalence do not always comprise an appropriate surrogate for 98 therapeutic equivalence (TE), which by definition is the ultimate goal of bioequivalence studies, ^[23] it 99 100 would seem appropriate to implement modeling and simulation approaches to assure therapeutic 101 outcomes in this arena too.

102 The aim of this review is to provide an overview of existing non-steady state PK/PD models, focusing 103 on drug-specific case examples. These are intended to serve as examples of the importance of 104 mechanistic PK/PD models in assuring desired therapeutic outcomes in clinical practice and to 105 encourage wider implementation of PK/PD in support of regulatory decision-making.

106 2

2 coThe effect compartment model

107

108 2.1 Overview

109

110 In many cases, the site of action of a drug is kinetically distinct from plasma and the equilibration 111 between the plasma and the effect site is often rather slow. In such cases, there will be a temporal 112 delay between the drug plasma (C_p) and effect site concentrations (C_e) and the effect will be a function of C_e rather than of C_p. Even though bioanalytical methods have improved greatly over the last decades, measuring the concentration at the effect site often remains a challenge, due to the lack of tissue accessibility.

In 1970, a hypothetical compartment serving as a link between the pharmacokinetic and pharmacodynamic models to address the equilibration kinetics was introduced by Segre et al.^[2] and was applied for the first time by Forester et al.^[24] to describe the time-course of effect of various cardiac glycosides.^[25] This approach, using a so-called «effect compartment» or «biophase distribution» model (Fig. 1), was further elaborated and described mathematically by Holford and Sheiner ^{[3],[26]} as follows:

122
$$\frac{dA_e}{dt} = k_{1e} \cdot A_p - k_{e0} \cdot A_e \quad (1)$$

Where A_p and A_e are the amounts of drug in the plasma (main compartment) and in the effect compartment, respectively, and k_{1e} , k_{e0} are the first-order rate constants for distribution and elimination from the hypothetical compartment, respectively.

Assuming that the effect compartment receives a negligible amount of drug and that distribution to and clearance from the biophase compartment are equal, the model can be simplified and then coupled with a pharmacodynamic model, for example a sigmoid E_{max} model:

129
$$k_{1e} \cdot V_p = k_{e0} \cdot V_e$$
 (2)

130
$$\frac{dC_e}{dt} = k_{e0} \cdot (C_p - C_e) \quad (3)$$

131
$$E(C_e(t)) = \frac{E_{max} \cdot C_e(t)^{\gamma}}{C_e(t)^{\gamma} + EC_{e50}^{\gamma}} \quad (4)$$

where C_p , V_p , C_e , V_e are the concentration and the volume in the central and effect compartment respectively; E_{max} , EC_{e50} and γ represent the maximum effect, the concentration in the effect site required to reach 50% of the maximum effect and the sigmoidicity factor, respectively. Alternatively, the hypothetical compartment could be coupled with a peripheral compartment instead of the central
compartment. However, it is not very common to use samples obtained at the effect site (e.g. using
microdialysis) or any other peripheral compartment as a pharmacokinetic surrogate.

138 A hallmark of the effect compartment model is the hysteresis observed in the effect-concentration 139 plot due to the time delay between pharmacokinetics and pharmacodynamics. In fact, this is a common attribute of non-steady-state pharmacokinetic/pharmacodynamic models.^[27] Well-known examples of 140 drugs exhibiting a biophase distribution delay related response include neuromuscular blocking agents 141 such as d-tubocurarine (see section 2.2) and pancuronium,^[28] the calcium channel blocker 142 verapamil,^[29] and the bronchodilator theophylline.^[30] Further cases that have been reported in the 143 144 literature include quinidine, disopyramide, opioids such as pethidine, morphine, fentanyl, diclofenac, organic nitrates, benzodiazepines and digoxin.^{[31],[32],[33],[34],[35],[36],[37],[38]} In the following section, the 145 models for tubocurare, pancuronium, ibuprofen and morphine are used to illustrate application of the 146 147 effect compartment model.

- 148 2.2 Applications and case examples
- 149

150 2.2.1 d-tubocurarine and pancuronium

151

152 The assumption of a direct relationship between pharmacokinetics and drug response has been 153 questioned for more than half a century, as illustrated by the case of d-tubocurarine.

Already in the early 1960s, the first attempts to simultaneously model pharmacokinetics and pharmacodynamics, based on the available plasma concentration and effect data for d-tubocurarine, were made. In 1964, Levy implemented a log-linear model to describe the time course of dtubocurarine response, assuming one-compartment pharmacokinetics following intravenous bolus administration, based on the results of Ryan et al.^[39] The log-linear model assumed that the effect of muscular relaxation is a linear function of the logarithm of the amount of d-tubocurarine present in the plasma,while elimination of the amount of d-tubocurarine in the body occurs exponentially with

time. In such cases, the pharmacologic activity declines linearly with time.^[1] In 1972, an open three-161 compartment model for the pharmacological effect of d-tubocurarine was proposed by Gibaldi et al.^[40] 162 The amount of drug in the central compartment at the time of recovery from neuromuscular block was 163 deemed by these authors to be dose-independent. This observation, combined with the very rapid 164 165 onset of action of d-tubocurarine, led the authors to the conclusion that the site of action is located in the central compartment,^[40] implying instantaneous equilibration between plasma concentration and 166 167 response. However, the data on which this model was based had been collected during the terminal 168 elimination phase, during which a pseudo-equilibrium between plasma and tissues concentration is 169 reached and the distributional delay is minimized.

By contrast, Hull et al.^[41] showed that after administration of pancuronium, a similar to d-tubocurarine 170 171 neuromuscular blocking agent, a linear relationship between the logarithm of concentration and the response is a poor predictor of the early phase response, in which a hysteresis between the 172 173 concentration in any compartment and twitch depression is observed. By adding a biophase 174 compartment, expressed similarly to equation (3), and assuming that same degree of paralysis (i.e. 175 during onset and offset of action) is associated with the same Ce, they were able to empirically relate 176 the intensity of pharmacologic effect to the concentration at the site of action at every time point using a fixed effect pharmacodynamic model.^[41] In the case of d-tubocurarine, the effect compartment 177 model, as described mathematically by Holford and Sheiner,^{[3],[26]} was successfully applied as well. 178 179 Plasma concentration and effect data after intravenous administration were analyzed from healthy 180 subjects and patients with renal failure. The model was able to fit data from both groups without statistically significant differences in the pharmacokinetic or pharmacodynamic parameters between 181 the two groups.^[42] Interestingly, the equilibration half-life (4 minutes) for pancuronium estimated in a 182 more empirical way by Hull et al.^[41] was very similar to the one for d-tubocurarine reported by Sheiner 183 et al.^[42] using an explicit pharmacokinetic/pharmacodynamic model. 184

In parallel, Stanski et al.^[43] explored the influence of various anesthetic agents on the muscle-relaxing
 effect of d-tubocurarine. Halothane induced-anesthesia, in comparison to anesthesia with morphine

187 and nitrous oxide, prolonged the equilibration half-life. An open two-compartment pharmacokinetic 188 model coupled with a hypothetical effect compartment was implemented to fit both plasma and 189 muscle paralysis data. Interestingly, changes in pharmacodynamic (k_{e0} , $t_{1/2ke0}$, EC₅₀), but not in pharmacokinetic, parameters were observed for patients under halothane anesthesia. Furthermore, it 190 191 was possible to distinguish between the effects of the agents on the EC_{50} for muscle paralysis showing 192 that halothane sensitizes the neuromuscular junction to d-tubocurarine. Provided that the diffusion of 193 tubocurarine into the extracellular fluid of the muscle and the receptor affinity is high, the rate limiting 194 step for the onset of action is the rate of muscle perfusion, which is inversely proportional to the 195 equilibration half-life $(t_{1/2ke0})$.^[43] Although the onset and the magnitude of response is dependent on muscle blood flow, the recovery from neuromuscular blockage is perfusion-independent and solely 196 related to the drug-receptor dissociation rate.^[44] The significant increase in t_{1/2ke0} under halothane-197 198 induced anesthesia is consistent with the decreased muscle blood flow, which would suggest a later 199 onset of paralysis. However, halothane also decreases the EC_{50} , which compensates for the decrease 200 in perfusion and results in a similar onset to that observed under morphine and nitrous oxide 201 anesthesia.

In summary, the evaluation of the pharmacodynamics in concert with the pharmacodynamics of these two muscle relaxants enabled a more mechanistic description of their dose-response characteristics and a better understanding of the drug interaction with the anaesthetic. These early successes triggered further interest in combining pharmacokinetics with pharmacodynamics to achieve a more mechanistic description of the relationship between dose, dosing and clinical effects.

207 2.2.2

Ibuprofen: dental pain relief

208

209 Ibuprofen was selected as a model drug to investigate the clinical relevance of bioequivalence metrics 210 to the therapeutic effect. An analysis of 25 bioequivalence studies of Ibuprofen immediate-release oral 211 dosage forms over a dose range from 200-600 mg showed that 14 of the studies failed to prove 212 bioequivalence in C_{max}, even though AUC fell within the bioequivalence limits.^[45] The authors reported that Ibuprofen, a weakly acidic BCS class II compound, is at higher risk to fail bioequivalence because of C_{max} variations. However, in cases where the plasma concentration is related non-linearly and/or indirectly to the drug effect^{[46],[3]}, the C_{max} and t_{max} values may not be accurate metrics for the therapeutic response. For example, if the C_{max} is higher than anticipated this will not necessarily translate to toxic effects. Likewise, if the C_{max} is lower, this will not necessarily result in lack of efficacy.^[47]

Dissociation between pharmacokinetics and pharmacodynamics is common for NSAIDS. This may be because of delayed distribution to the biophase or related to an indirect response mechanism, for example when the pharmacodynamic endpoint is the inhibition of inflammation mediators.^[48] Pain relief and antipyresis after administration of ibuprofen formulations have been extensively modelled in different populations. In this section, the main studies for pain relief after third molar extraction are presented, while studies investigating the antipyretic effect are addressed in section 4.2.1.

225 Third molar extraction pain models describe the postoperative onset of inflammation, with maximum 226 pain intensity occurring in 12 hours or less. Relief from pain associated with tooth extraction exhibits 227 high reproducibility and a low placebo effect, features that are important for differentiation among various doses and thus for the identification of dose-response curves.^{[49],[50],[51],[52]} The most commonly 228 229 evaluated endpoints in dental pain models are the pain intensity difference (PID) and sum of pain 230 intensity difference (SPID), the pain relief (PAR) and total pain relief (TOTPAR), the time to re-231 medication (REMD), the time to first perceptible pain relief (TFPR) and time to first meaningful pain relief (TFMP).^{[53][54]} 232

In a double-blind, randomized, single- and multi-dose study of 254 adult patients, who had undergone third molar surgery, Hersh et al.^[50] reported a positive dose-response relationship for sum pain intensity (SPID), total pain relief (TOTPAR), time to re-medication (REMD) and overall pain relief, after administration of 200 and 400 mg of ibuprofen as a single-dose. During the multi-dose phase, no significant differences between the two dose levels were detected. The authors concluded that 238 patients could benefit from higher doses for pain treatment immediately after the extraction, but that lower doses would be satisfactory thereafter. These results suggest that the single-dose approach 239 240 adopted for bioequivalence testing might be over-discriminating for the assessment of ibuprofen 241 formulations with regard to the maintenance of dental pain relief. Indeed, McQuay et al.^[55] observed no significant differences between 200 and 400 mg of ibuprofen in a double-blind, randomized, 242 243 placebo-controlled, single-dose study comparing the analgesic effect of 200 and 400 mg of ibuprofen 244 with placebo and with 200 mg ibuprofen plus 50, 100 or 200 mg caffeine in 161 adult patients after 245 third molar removal. In a further study, a positive dose-response relationship of ibuprofen over the 246 dose range 50-400 mg with regard to sum of pain intensity difference (SPID) and total pain relief (TOTPAR) was reported by Schou et al.^[54] However, in terms of TOTPAR the doses of 200 and 400 mg 247 248 did not differ significantly.

A meta-analysis of data from 13 trials with total of 994 patients reported an absolute increase of only 9% (from 59% to 68%) in the number of patients who achieved at least 50% pain relief, when the dose of ibuprofen was doubled from 200 to 400 mg, meaning that 10 patients would need to be treated with the higher dose for just one of them to benefit. ^[56] The analysis indicates that the dose-response relationship is rather flat in the dose range 200 to 400 mg with respect dental pain relief by ibuprofen.

Li et al.^[53] applied a pharmacodynamic model to investigate the onset and offset of dental pain relief after administration of effervescent and standard tablets containing 400 mg ibuprofen. As an endpoint, a categorical pain relief score was applied and treated as a continuous variable, in agreement with Lemmens et al.^[57] The observed distributional delay of the response to ibuprofen was addressed by the addition of an effect-compartment model and the overall effect as the sum of placebo and drug was described as following:

260
$$\frac{d(C_e[t])}{dt} = k_{e0} \cdot \{C_p[t] - C_e[t]\}$$
(5)

261
$$f_d(C_e) = \frac{E_{max} \cdot C_e^{\gamma}}{C_e^{\gamma} + EC_{50}^{\gamma}} \quad (6)$$

262
$$f_p[t] = P_{max} \cdot \left(1 - e^{-k_p \cdot t}\right) \quad (7)$$

263
$$PR(t) = f_p[t] + f_d(C_e) + \varepsilon \quad (8)$$

where C_p and C_e are the drug concentrations in plasma and in the effect-site compartment, respectively; k_{e0} and k_p are the first-order rate constants for the placebo effect and equilibration, respectively; E_{max} and P_{max} are the maximum ibuprofen and placebo effect, $f_d(C_e)$ and $f_p[t]$ are the pain relief by ibuprofen and placebo, respectively; γ and EC_{50} are the sigmoidicity factor and the drug plasma concentration to achieve 50% of E_{max} , respectively; PR(t) represents the pain relief score at a given time t and ε stands for the normally distributed residual variability.

The model was able to describe the pain relief score data adequately and the effect was directly related to the effect-site concentration, which increased much faster for the effervescent than the standard tablets, with the peak effect site-concentration occurring one hour earlier than for the standard tablet (1.0 h versus 2.0 h). The sigmoidicity factor was estimated to be 2.0 ± 0.43 , confirming the relatively flat dose-response curve of ibuprofen.

275 More recently, a PBPK/PD model for Ibuprofen was developed and validated by Cristofoletti and Dressman^[58] with the SimCyp Simulator[®] version 12.2 (SimCyp Ltd.), fitting antipyretic and dental pain 276 277 relief pharmacodynamic models to pharmacokinetic and pharmacodynamic data already published in 278 the literature. The main goals of this study were a comprehensive evaluation of the clinical relevance 279 of bioequivalence criteria for ibuprofen immediate-release oral dosage forms and a risk assessment of 280 waiving in vivo bioequivalence studies of such products. To simulate the pharmacokinetic and 281 pharmacodynamic profiles, virtual populations similar to those enrolled in the clinical studies by Walson et al.^[59] and Li et al.^[60] in terms of age and gender ratio were generated, such that virtual trials 282 283 for the dental pain relief model included 100 adults per trial aging between 18-40 years and receiving 284 tablets of 100, 200, 280 or 400 mg of Ibuprofen. One-at-a-time sensitivity analysis for the gastric 285 solubility, gastric emptying time (GET), apparent permeability coefficient (P_{app}) and small intestine pH 286 was conducted and the effect of applying different dissolution rates in the simulations on the resulting pharmacokinetic and pharmacodynamic profiles was also investigated.^[58] The authors found that the 287 dose-response curve for dental pain relief is shallow and as a result relatively insensitive to changes in 288 289 plasma concentrations within the range 12-23 mg/L (applying an EC₅₀ of 10.2 mg/L). Comparing the 290 pharmacodynamic response after the simulated administration of 280 versus 400 mg lbuprofen tablets 291 to adults undergoing third molar extraction, no significant differences in the response occurred. 292 Interestingly, although (under the assumption that the 400 mg tablet is the reference product and the 293 280 mg tablet is the test product in a virtual bioequivalence scenario) the test product would not be 294 bioequivalent to the reference product in terms of pharmacokinetics (Cmax ratio (C_{max-T}/C_{max-R}) of 0.7), 295 the 280 mg tablet would be still considered therapeutically equivalent to the 400 mg tablet for dental 296 pain relief in adult patients.

297 Cristofoletti and Dressman combined in vitro in vivo extrapolation with PBPK/PD model to simulate the 298 effect of different dissolution rates from products containing ibuprofen free acid (IBU-H) and salts (IBU 299 salts) and to investigate whether these would a) reflect reported differences in pharmacokinetics as 300 well as whether b) differences in pharmacokinetics would translate into difference in the ability of ibuprofen to relieve dental pain in adults.^[61] The model was able to adequately predict the observed 301 302 pharmacokinetic profiles. The pain relief model by Li et al.^[60] was adopted to simulate ibuprofen 303 response. As expected from the faster dissolution of the products containing salt forms of ibuprofen, 304 the 90% confidence intervals (CI) for C_{max} did not meet the average bioequivalence (ABE) acceptance 305 criteria. However, pain relief scores elicited by ibuprofen free acid and salts were identical. 306 Interestingly, the simulated peak effect-site concentrations for both IBU-H and IBU salts 400 mg were 307 found to be higher than the estimated $EC_{80} \approx 20 \text{ mg/L}$, indicating that the extent of pain relief would be 308 insensitive to pharmacokinetic changes at this dose level. Importantly, the duration over which the 309 effect-site concentrations are maintained above EC₈₀ should be also taken into account. The authors concluded that the bioequivalence criteria for C_{max} might be over-discriminatory and not clinically 310

311 relevant for assessing therapeutic equivalence of ibuprofen products in terms of overall dental pain relief. 312

313 As illustrated by the example of ibuprofen, therapeutic equivalence is not always captured 314 appropriately by simple plasma concentration measurements due to the insensitivity of the 315 pharmacodynamic response to the pharmacokinetics in the dose range typically applied. From this 316 case example, it is evident that the interaction of the drug pharmacokinetics with the pharmacologic 317 response should be taken into account to set clinically relevant specifications ("safe spaces") for drug 318 products. Modeling and simulation techniques would be a powerful tool in this direction, facilitating a 319 regulatory transition from the current "one size fits all" bioequivalence paradigm to a scenario based 320 on the clinically-based, specific PK/PD characteristics of the drug product and thus able to provide a 321 more accurate assessment of therapeutic equivalence.

- 322 2.2.3 Anti-nociceptive effect of morphine
- 323

324 For drugs, which exhibit high biological target affinity and/or reach their site of action by active 325 transport mechanisms, distribution to the biophase may or may not impose a rate-limiting step. Over 326 the past few years, several specific transporters that may influence the distribution of drugs to their site of action in the central nervous system (CNS) have been identified.^{[62],[63],[64],[65]} However, the 327 328 number of pharmacokinetic/pharmacodynamic (PK/PD) studies exploring the functional role of these 329 transporters in the distribution to the effect site are few. One interesting example is the anti-330 nociceptive effect of morphine, for which mechanism-based models of the biophase distribution 331 within the central nervous system were established using intracerebral micro-dialysis.

Letrent et al.^[66] investigated the effect of GF120918, a potent and selective P-glycoprotein (P-gp) 332 333 inhibitor, on the pharmacokinetics and pharmacodynamics of morphine in rats, which were 334 randomized into GF120918 pretreated, vehicle and control groups. The concentrations of both 335 morphine and its metabolite, morphine-3-glucoronide (M3G), in serum were quantified and the anti-336 nociception was expressed as the percentage of maximum possible response (% MPR). A two-

337 compartment pharmacokinetic model, together with an effect compartment coupled to a sigmoidal E_{max} model was employed to simultaneously fit the pharmacokinetic and pharmacodynamic data. 338 339 Among the pharmacokinetic (AUC, Cl, MRT, V_{ss}) and pharmacodynamic (k_{e0} , EC₅₀, γ) parameters 340 evaluated, only the equilibration rate constant (ke0) and the %MPR were significantly altered by pre-341 treatment with GF120918, indicating a faster onset and more intense action, respectively (p=0.0023). 342 The increased pharmacodynamic response could not be attributed to pharmacokinetic changes or to 343 the elevated M3G concentrations. Since M3G does not possess any anti-nociceptive 344 properties,^{[67],[68],[69]} the authors suggested that the inhibition of P-gp by GF1920918 might diminish the 345 efflux of morphine from brain capillary endothelial cells, leading to more rapid distribution and higher concentrations of morphine at its site of action. These data were supported by Xie et al.^[70], who 346 347 demonstrated, using trans-cortical micro-dialysis, that morphine concentrations in the brain were 348 increased (1.7-fold) after administration to mdr-1a genetic deficient rats, whereas the metabolite M3G 349 was unaffected.

Evaluation of the kinetics of biophase distribution within the central nervous system by intracerebral microdialysis, which has already been successfully applied to the characterization of the distributional behavior in several cases ^{[71],[70],[72],[73]}, is a promising tool for the development of more sophisticated, mechanism-based models, enabling as yet unexplained aspects of the pharmacodynamics of the central nervous system acting drugs to be illuminated.

355

356 3 Modeling of irreversible mechanisms of action

357

358 3.1 Overview

359

In this section, we describe some examples of drugs that act in the human body through irreversible inhibition at the site of action. In general, pharmacodynamic (PD) effects are initiated by the interaction of drugs with targets such as receptors, enzymes, ion channels, cell membranes etc. Such interactions may be reversible, with a balance between association and dissociation of the drug with the target, or irreversible when a drug bonds covalently to the target or the dissociation rate is extremely slow for the relevant time span. As a result of these interactions, a cascade of events is triggered, leading to the pharmacological effect, which can either stimulate (agonist) or inhibit (antagonist) a physiological process.^{[74],[75]}

368 In many cases, drugs that irreversibly inhibit a physiological process are transformed, as a first step, 369 into reactive metabolites, which then bind covalently to their target, resulting in its inactivation. In 370 order for the pre-existing situation to be reestablished, it is necessary to resynthesize the target. In 371 such cases, the duration of action is likely to be independent of the pharmacokinetic half-life of 372 elimination of the drug and instead depends essentially on the *de novo* synthesis of the target. The 373 irreversible inactivation of endogenous enzymes or receptors caused by drugs e.g. the antiplatelet effect of aspirin after binding cyclo-oxygenase-1,^{[76],[77]} the 5 α -reductase inhibitors,^{[78],[79]} and the 374 proton pump inhibition by proton pump inhibitors (PPI),^{[80],[81],[82]} are often described using such 375 376 turnover models. Further examples are drugs that trigger apoptosis in human cells, bactericidal antibiotics,^[83] reduction of viral load due to the treatment with antivirals,^[84] cell death processes 377 induced by anticancer drugs^[85] and cytotoxic drugs which cause myelosupression.^[86] 378

In general, the turnover models that have been presented in the literature are based on the following
 differential equation:^[87]

381
$$\frac{dR}{dt} = k_{in} - k_{out} \cdot R - f(C) \cdot R \qquad R(0) = R_0$$
(9)

where *R* denotes the response produced by the drug, R_0 is its initial response value, k_{in} is a zero-order rate constant for the response, k_{out} is a first-order elimination rate constant and the function of the drug concentration f(C) can be interpreted as a bimolecular interaction of the drug or its active metabolite with the target. This is the general equation representing the turnover rate of the response, however, more complex scenarios are also possible, requiring more mechanistic models to be developed as will be discussed later.

Figure 2 depicts a turnover model that can be applied to the interaction between the drugs with receptors, enzymes or ion channels. In the case of interaction with endogenous enzymes, the k_{in} and k_{out} parameters represent apparent rates of response formation and dissipation respectively and f(C)represents the effect as a function of drug concentration.

393

394

4 3.2 Applications and case examples

- 395
- **396** 3.2.1 Proton pump inhibitors
- 397

Proton pump inhibitors (PPIs) were chosen as the drug model for this topic since their inhibition of the proton pump (H⁺, K⁺-ATPase) enzyme present in the parietal cells of the stomach is irreversible. To understand the mechanism of inhibition by the PPIs, models describing the turnover of H⁺, K⁺-ATPase have been described.

402 The PPIs are, in and of themselves, inactive drugs that require an acid environment for their activation. 403 These weakly basic substances reach the general circulation after absorption from the gastrointestinal 404 tract and then become concentrated in the acid compartment of the parietal cells present in the gastric 405 mucosa. Following their activation by conversion to the sulphonamide form in the acidic intracellular 406 environment of the parietal cells, a covalent bond occurs between the activated PPI and cysteine 407 residues present in H⁺, K⁺-ATPase. This enzyme is responsible for the final step in the secretory gastric 408 acid process.^{[81],[88],[89]} As a consequence of the binding, the enzyme is inactivated and this results in suppression of acid secretion into the gastric lumen.^{[90],[80]} PPIs inhibit both basal and stimulated gastric 409 410 acid secretion, regardless of the nature of stimulation of the parietal cells. In order for the acid secretion to be re-established, *de novo* synthesis of H⁺, K⁺-ATPase is necessary.^{[90],[91],[92]} 411

Even though the elimination half-life of PPIs is only 1-2 hours, the pharmacodynamic half-life of the inhibitory effect on H⁺, K⁺-ATPase is about 48 hours, rendering a rapid elimination (PK) but long duration of response (PD) to members of this class.^{[92],[93],[94]} By comparison, the pharmacodynamics of

415 drugs that reversibly bind to the proton pump to decrease acidic secretion in the stomach, such as 416 cimetidine and other H₂ receptor antagonists, can be described with a direct response PD model.^[95] To construct a mechanistic PK/PD model for PPIs, several factors have to be considered: the 417 accumulation of PPI in the parietal cell, the amount of active enzymes present in the canaliculus of 418 419 parietal cell, the rate of de novo synthesis of new proton pump enzymes, the metabolism and 420 inactivation of PPIs, the extent of covalent PPI binding to the proton pump in the parietal cell and the stability of this binding.^[96] Because of this complexity, several different models have been proposed to 421 422 describe the relationship between PK and PD for this class of drugs. There are empirical models that 423 simply consider the turnover of the proton pump and those that are more mechanistic, taking into 424 account the relevant physiology and PPI characteristics. In this section we will focus on PK/PD models 425 that have been used to describe the difference between the elimination half-life (PK) of PPIs and the temporal inhibition of acid secretion (PD) that results from binding of the PPI with H⁺, K⁺-ATPase. 426

Katashima and co-workers^[95] were the first to publish a mechanistic PK/PD model for PPIs. In the first study, a model relating the unbound plasma concentration (C_f) of lanzoprazole and omeprazole to the inhibitory effect on stomach acid secretion was developed. This model, illustrated in Figure 3, utilizes the apparent turnover process of H⁺, K⁺-ATPase to describe the relationship between plasma concentration and the inhibitory effect of the PPIs on gastric acid secretion.^[97]

432

According to this PK/PD model, the inactive form of the PPI is present in the plasma, and only after reaching the acid environment of the parietal cells is it transformed into the active form. This form then reacts with active H⁺, K⁺-ATPase according to a second order reaction with the rate constant, *K*, to establish a covalent bond between the activated PPI and H⁺, K⁺-ATPase, resulting in inactivation of the enzyme.

438

439 The total amount of proton pump (E_t) remains at a constant level (k_s/k_I) because H⁺, K⁺-ATPase is 440 synthesized, on the one hand, at a rate described by the rate constant, K_s , but also eliminated, on the

other hand, at a rate described by the first order rate constant k_1 . The inactive proton pump recovers at a rate described by the first order rate constant k_2 . Under these circumstances, the apparent turnover rate constant, k, is represented by $k_1 + k_2$. The time courses of variation in the amount of active H⁺, K⁺-ATPase (*E*) and the inactive fraction (*E_c*) are expressed by the following equations:

445
$$\frac{dE}{dt} = -K \cdot C_f \cdot E - k \cdot E + k_2 \cdot E_c + K_s \quad (10)$$

446
$$\frac{dE_c}{dt} = K \cdot C_f \cdot E - (k_1 - k_2) \cdot E_c \quad (11)$$

An *in vivo* pharmacokinetic and pharmacodynamic study in rats was conducted over a dose range of 0.006 - 3 mg/kg (IV) with omeprazole and lanzoprazole. Using the data from intravenous administration in rats, the estimated half-life of the proton pump was 27 times longer than the elimination half-life for omeprazole and 66 times longer for lansoprazole. Using the PK/PD model described above, good agreement between predicted and observed data was achieved for both drugs.

After their success with the PK/PD model in describing the data from rats, Katashima and co-workers^[81] 453 454 extended the model to human studies with pantoprazole (PPZ), lansoprazole (LPZ) and omeprazole 455 (OPZ). The PK/PD analysis of these PPIs in humans was conducted using data obtained after oral 456 administration of OPZ (40mg), LPZ (30mg) and PPZ (40mg). Again, good agreement between the 457 predicted and observed values for the parameters was achieved. The estimated half-life of elimination 458 for omeprazole was 0.854 h, for lansoprazole 1.66 h and for pantoprazole 1.52 h, while the apparent 459 recovery half-life of the inhibitory effect on gastric acid secretion was 27.5 h for omeprazole, 12.9 h 460 for lanzaprole and 49.9 h for pantaprazole. These results confirmed the divergence between plasma 461 concentration (PK) and the inhibitory effect on gastric acid secretion (PD) of these there PPIs.

462

The mechanistic PK/PD model was extended by Puchalski and co-workers for lansoprazole.^[82] Their model was set up to describe the intra-gastric pH time profile over a 24 hour period, enabling the circadian rhythm of acid secretion and food effects on intra-gastric pH to be taken into account. Using this model, the estimated value for lansoprazole half-life of elimination was 3.2h, somewhat longer

than in the Katashima model (1.66 h), while in the clinical study the pH had not returned to the baseline 467 level after 24h. As this proposed model took into account several factors that can interfere in the PPI 468 absorption and activation, it should be particularly useful in the design of clinical studies, the prediction 469 470 of the optimal dosing regimen and the investigation of PPI effects in different patient populations.^[82] 471 The inhibitory effect of PPIs on gastric acid secretion has also been described by Abelo and coworkers^[80] using a simpler, empirical turnover model type I, as introduced by Dayneka et al.^[98] (see 472 473 section 4.1.1). In the basic turnover model shown in Eq. 12 and applied to omeprazole in Figure 4, it is 474 assumed that the drug inhibits or stimulates the production of an effect, which can be characterized 475 by the zero order k_{in} turnover and the elimination first order k_{out} rate constants as appropriate. The 476 rate of change of the response (R) provoked in the absence of the drug is described with the following 477 equation:

- 478
- 479

 $\frac{dR}{dt} = k_{in} - k_{out} \cdot R \quad (12)$

480

481 According to Eq. 12 the acid secretion (*AS*) is directly proportional to the concentration of the active
482 proton pump enzyme (*E*). Equation 13 can be used to correct for the placebo effect on acid secretion:

483
$$R = \frac{AS(Drug,t)}{AS(Placebo,t)} = \frac{E(Drug,t)}{E(Placebo,t)} \quad (13)$$

484

Omeprazole irreversibly removes the enzyme from the system at a rate proportional to the amount of enzyme and the inhibitor concentration. Irreversible removal of the enzyme results in a decrease in the response according to equation 14:

488
$$\frac{dR}{dt} = k_{in} - \left(k_{out} + k_{ome} \cdot C_p\right) \cdot R \quad (14)$$

489

490 For a given concentration of omeprazole, the value for R at steady state (R_{ss}) will be:

491
$$R_{ss} = \frac{k_{in}}{k_{out} + k_{ome} \cdot c_{pss}} \quad (15)$$

492 This relationship states that with increasing omeprazole concentration, R_{ss} approaches zero.

Data from studies in dogs were used to predict the PK and PD parameters for omeprazole for this species, leading to a prediction for the half-life of elimination of 1.3 h and for the effective half-life for inhibition of acid secretion ($t_{1/2 \text{ Kout}}$) of 51h. Using allometric scaling, the predicted half-life for humans was 1.5 h and the effective half-life for inhibition of acid secretion ($t_{1/2 \text{ Kout}}$) was 71.7 h. The discrepancy between predicted (71.7 h) and observed (48) $t_{1/2 \text{ Kout}}$ in humans was attributed to differences in basal acid secretion between dogs and humans. ^[99]

499

Ferron and co-workers ^[100] also used the basic turnover irreversible PK/PD approach, in this case to describe the inhibition of gastric acid secretion by pantoprazole in rats and humans. The model was able to adequately describe the time course of gastric acid secretion in rats at all doses studied. The next step it was to apply it to gastric secretion data obtained after single or multiple oral or intravenous administration of pantoprazole in humans. The estimated half-life for pantoprazole was 0.5 h in rats and 0.8 h in humans, in agreement with the observed data in both species.

506

507 Both the mechanistic and empirical models described in this section were able to predict the 508 discrepancy between the half-life elimination (PK) of PPIs and the time-course of inhibition of acid 509 secretion (PD). The models were also successful in describing further characteristics of PPIs, namely that the effect in acid secretion inhibition of PPIs is linked to the extent of exposure (AUC), and that 510 511 the onset of action is governed by the maximum concentration (C_{max}). Thus, PK/PD modelling provides 512 a powerful tool for analysing/predicting effects achieved with other dosing regimens. To circumvent 513 the use of invasive methods in clinical studies for monitoring the gastric pH and inhibition of gastric 514 acid secretion, it would be necessary to build PK/PD models that can also predict the extent of acid 515 inhibition in terms of the pH value and the duration over which the pH is kept above a clinically relevant threshold value (usually pH 4) by the PPI. 516

In conclusion, modelling and simulation clearly shows why PPIs, despite having a short plasma half-life,
are able to have a long duration of effect. Such models enable better decisions to be made about
dosing intervals and also help to identify the time-frames over which drug/drug interactions with PPIs
may persist.

522 3.2.2 Acetylsalicylic acid

523

524 Similarly to the PPIs, aspirin (ASA) has a long duration of action, even though it has a short elimination half-life (t_{1/2} 18-30 min).^{[101],[102]} ASA inhibits platelet-derived thromboxane (TXB2), with approximately 525 60% inhibition still observed four days after discontinuation of ASA.^{[101],[102]} This pronounced 526 527 dissociation between the elimination half-life (PK) and the time-frame of drug action (PD) occurs 528 because ASA binds covalently to TXB2 causing irreversible inhibition of this enzyme. The TXB2 activity 529 can only be re-established by synthesis of new platelets, which is a process that occurs over a period of approximately 10-14 days.^[101] Because platelets are not nucleated, they are unable to synthesize 530 531 new COX-1, and for this reason platelet function will only normalize after the platelets that have been 532 acetylated by ASA are removed from the systemic circulation and replaced by new platelets derived from megakaryocytes.^[103] 533

534

The first model describing cyclooxygenase activity in platelets and the blood vessel endothelium after oral administration of aspirin was developed by Yamamoto and co-workers.^[77] These authors used irreversible inhibition, with renewal by enzymatic turnover, to explain the long duration of the antiplatelet effect of aspirin in humans. In this study thromboxane B₂ concentrations and the percentage of prostacyclin production in the blood vessels were used as biomarkers.^[77]

540

It has been suggested that non-selective COX-1 inhibitors, e.g. ibuprofen, could limit the cardioprotective effect of aspirin.^[104] For this reason Hong and co-workers^[76] developed a PK/PD model that was based on the turnover of the COX-1 enzyme, in which the irreversible inhibition by aspirin and the reversible binding by ibuprofen were both incorporated. The rate changes of free

enzyme concentration available for aspirin binding (*E*) and the ibuprofen-enzyme complex (*EI*) were
described by the following equations:

547
$$\frac{dE}{dt} = k_{in} - k_{out} \cdot E - K \cdot C_{asa} \cdot E - k_{on} \cdot C_{ibu} \cdot E + k_{off} \cdot EI \quad (16)$$

548
$$\frac{dEI}{dt} = k_{on} \cdot C_{ibu} \cdot E - k_{off} \cdot EI - k_{out} \cdot EI \quad (17)$$

where k_{in} is the zero-order production effect rate constant, k_{out} is the first order elimination rate constant, K is the second-order rate constant for the irreversible enzyme inactivation by aspirin, and k_{on} and k_{off} are the association and dissociation rate constants for binding of ibuprofen on the enzyme. C_{asa} and C_{ibu} represent the aspirin and ibuprofen concentrations in the plasma, assuming that both drugs follow a one compartment PK model with first order rate constants for absorption and elimination.

The mechanistic PK/PD model was able to reflect the anti-platelet effect of aspirin administered either alone or concomitantly with ibuprofen. As well as simulating the PK and PD time courses, significant inhibition of the antiplatelet effects of aspirin in the presence of a typical ibuprofen regimen was also demonstrated.

The most mechanistic PK/PD model describing the effects of aspirin on COX-1 activity to date was proposed by Giareta and co-workers.^[105] This model uses a population of megakaryocytes (MK) and peripheral platelets present in the blood circulation to describe aspirin's antiplatelet activity, as shown in Figure 5.

For the construction of the PK/PD model for aspirin, the inactivation of COX-1 by low dose aspirin and the recovery of COX-1 after stopping treatment were taken into consideration. Other physiological processes, e.g. the description of the megacariopoiese process responsible for the maturation and generation of new platelets, were also accounted for. The basic characteristics of the megacariopoiese process are shown in Figure 5. The schematic description of the resulting PK/PD model is shown in Figure 6. It consists of three linear compartments to describe the PK behavior of aspirin and two non-

linear compartments to describe the mechanism of inactivation of COX-1 (PD) in MK cells and in the
platelets generated from them. A full mathematical description of the model has been published by
Giaretta and co-workers.^[105]

572

The PK and PD parameters of the model were inferred from the literature and calibrated by measurements of TXB2, which represents the COX-1 activity in peripheral platelets, in 17 healthy subjects and 24 patients with essential thrombocythemia (ET).^[105] The model was able to reproduce both the mean TXB2 inhibition time in healthy patients and the reduced inhibition of TXB2 seen in patients with ET. Thus, this mechanistic PK/PD model may helpful to customize aspirin regimens under conditions of altered megakaryopoiesis.

579

In addition to the dissociation between PK (short half-life of elimination) and PD (long response period) demonstrated by the models described above, the dose-response relationship for platelet inhibition by aspirin is flat. Feldman and co-workers^[101] demonstrated that even with a 10-fold increase in dose of aspirin, only a two-fold increase in response (inhibition of TXB2) was observed. Since doses of 81 and 325 mg of ASA are not significantly different with regard to this clinical response, applying a low dose of aspirin to prevent platelet aggregation is justified.^[101]

In summary, mechanistic models of the pharmacodynamic action of aspirin on platelets appear to be useful for customizing the prevention of thrombus formation and for designing clinical trials in special patient populations e.g. the elderly, pregnant women, children, obese patients, etc. Indeed, regulatory authorities are increasingly relying on and encouraging the use of modeling and simulation to forecast changes in PK and PD in rare diseases and in special populations of patients in whom it is challenging to perform clinical trials.

592 3.2.3 Exemestane

Exemestane, an irreversible aromatase type I (Ar type I) inhibitor for the treatment of advanced breast
cancer of postmenopausal women, provides a further, interesting example of irreversible binding and
biological target inactivation.

597

598 In an open, three-period, randomized, crossover study of twelve healthy post-menopausal women 599 Valle et al. investigated the effects of formulation (suspension versus tablet) and administration of 600 food (i.e. fasted versus fed) on the pharmacokinetics and pharmacodynamics of exemestane. As had 601 already been demonstrated by previous clinical trials, oral administration of exemestane (25 mg/day) 602 inactivates peripheral aromatase, leading to a 85-95% decrease in basal plasma estrone, estradiol and 603 estrone sulphate (EIS) concentrations in post-menopausal women with advanced breast cancer. ^{[106],[107],[108]} First, population pharmacokinetic models, consisting of a mono- or bi- exponential 604 605 absorption and three compartment distribution function, with empirical Bayesian estimates for each 606 individual were developed. Absorption lag times were determined for both absorption models. An 607 inhibitory (type I) indirect response pharmacodynamic model (see more details in section 4.1), in which 608 synthesis and elimination of EIS (which is indirectly related to aromatase activity) are governed by zero-609 and first-order rate constants, respectively, was implemented to describe the dissociation between 610 plasma concentrations and the observed effect:

611
$$\frac{dC_{EIS}}{dt} = k_s - k_o \cdot C_{EIS} \quad (18)$$

612
$$\frac{dC_{EIS}}{dt} = k_s \cdot \left(\frac{C^{\gamma}}{C^{\gamma} + IC_{50}^{\gamma}}\right) - k_o \cdot C_{EIS} \quad C_{EIS}(0) = C_{EIS 0} \quad (19)$$

613 where C_{EIS} is the plasma concentration of estrone sulphate, k_s is the zero order rate constant for 614 synthesis and k_o is the first-order rate constant for elimination, C^{γ} is the exemestane plasma 615 concentration, IC_{50} represents the exemestane plasma concentration at which 50% of inhibition is 616 achieved and γ is the Hill-coefficient. This semi-empirical, non-linear mixed-effect modeling approach 617 fitted the data adequately. 618 A more mechanistic model, incorporating the irreversible aromatase inactivation by exemestane, was 619 also applied. In this model the aromatase concentration, Ar, is assumed to be the system variable 620 controlling the rate of synthesis of EIS. The production and elimination rate of aromatase is in turn governed by a zero-order (k_{se}) and first-order (k_{oe}) rate constant, respectively. The irreversible 621 622 inhibition of aromatase by exemestane is characterized by an increase in the elimination of aromatase 623 and represented by a second-order rate constant k_i . Assuming that the concentration of EIS precursor 624 is constant and the concentration of aromatase is known, the model is fully identifiable. The rate of 625 concentration changes of EIS and Ar are defined by the equations:

626
$$\frac{dC_{EIS}}{dt} = k_s \cdot Ar - k_o \cdot C_{EIS} \quad C_{EIS}(0) = C_{EIS 0} \quad (20)$$

627
$$\frac{dAr}{dt} = k_{se} - k_{oe} \cdot Ar - k_i \cdot C_{EIS} \cdot Ar \quad Ar(0) = Ar_0 \quad (21)$$

628

629 where Ar_0 is the baseline concentration of aromatase.

630

631 The adoption of a more physiological relevant mechanism of action in the model was expected to 632 provide better results. Nevertheless, the goodness of fit was not significantly improved over the type 633 I indirect response model. Despite being semi-empirical, the type I indirect-response model was able 634 to predict the drug effect in different scenarios (i.e. doses, dosage regimens), providing an external 635 validation. In a sense, the initial, indirect response type I model could be considered as a "collapsed" 636 form of the mechanism-based model, under the assumptions that Hill-coefficient is equal to one (γ =1) 637 and that the aromatase dynamics equation is solved at equilibrium and then substituted in the EIS equation. These assumptions appear to be justified in the case of exemestane, since the 638 639 pharmacodynamic parameters do not change significantly in the data range studied and a value of Hill-640 coefficient 1.75 (γ =1.75) has been reported. Hence, a relatively flat dose-response is implied.

An almost 4-fold increase in the absorption rate of exemestane when administered as a suspension as compared to a tablet was detected, while food intake decreased the absorption rate. Interestingly, these differences were mitigated in terms of pharmacodynamic response such that the maximum effect and time to maximum effect were not significantly different among treatment groups. The authors concluded that even large differences in pharmacokinetics arising from formulation or administration with food were not translated to a meaningful difference in pharmacodynamics.

648

The example of exemestane is interesting for two main reasons: a) it illustrates that a mechanismbased model of irreversible pharmacodynamics can be transformed, depending on data availability or fast equilibration, to a simplified, "collapsed" model, without influencing the outcome appreciably, and b) observed differences in absorption patterns and food effects are not always clinically relevant, especially when there is a long delay between plasma levels and the elicited drug response. Again, these findings support the consideration of pharmacodynamics as well as pharmacokinetics when determining whether two drug products or two dosing scenarios are therapeutically equivalent.

656

657 4 Indirect response and feedback control models

658

659 4.1 Overview

660

661 Most pharmacological targets are subject to homeostatic mechanisms, characterized by continuous 662 degradation on the one hand and re-synthesis of one or more biomarkers (e.g. enzymes, antibodies, 663 circulating proteins or inflammation factors) to compensate for elimination on the other hand, which 664 balance each other to maintain a stable steady-state. This is often referred to as the turnover process. 665 Some drugs elicit their action by perturbing the steady-state, resulting in a temporary or a more 666 permanent change in the marker value. Such mechanisms of actions, which do not affect the response 667 itself but rather influence the turnover process, are inherently indirect and the models describing their 668 effect-time course are usually referred to as turnover or indirect response models. These models 669 typically exhibit a delay between the drug concentration-time and response-time profiles. The 670 amplitude of the response and the extent of the time delay are dependent on the turnover rates 671 (synthesis and degradation) of the pharmacological target as well as the magnitude of the effect.

672 4.1.1 "Basic" and "extended basic" indirect response models

Nagashima et al.^[109] were the first to implement an indirect response model, which was used to explain the anticoagulant effect of warfarin on the activity of the prothrombin complex. In 1993, Dayneka et al.^[110] introduced four basic mathematical models describing the indirect pharmacological processes, according to which the production and loss of the response, R, are governed by zero- and first-order rate constants, k_{in} and k_{out}, respectively. The drug can inhibit or stimulate the synthesis and/or the elimination process as follows:

680 Model I (inhibition of k_{in}):

673

681
$$\frac{dR}{dt} = k_{in} \cdot \left(1 - \frac{I_{max} \cdot C}{C + IC_{50}}\right) - k_{out} \cdot R, \quad R(0) = R_0 \quad (22)$$

682 Model II (inhibition of k_{out}):

683
$$\frac{dR}{dt} = k_{in} - k_{out} \cdot \left(1 - \frac{I_{max} \cdot C}{C + IC_{50}}\right) \cdot R, \ R(0) = R_0 \quad (23)$$

684 Model III (stimulation of k_{in}):

685
$$\frac{dR}{dt} = k_{in} \cdot \left(1 + \frac{E_{max} \cdot C}{C + EC_{50}}\right) - k_{out} \cdot R, \quad R(0) = R_0 \quad (24)$$

686 Model IV (stimulation of k_{out}):

687
$$\frac{dR}{dt} = k_{in} - k_{out} \cdot \left(1 + \frac{E_{max}C}{C + EC_{50}}\right) \cdot R, \ R(0) = R_0 \quad (25)$$

where k_{in} , k_{out} are the zero order production and first order elimination rate constants, C is the drug plasma concentration, and EC_{50} and IC_{50} represent the drug plasma concentrations achieving 50% of the maximum stimulating, E_{max} , and inhibitory, I_{max} , effects, respectively.

691 These four basic models, which are illustrated in Figure 7, have been applied extensively and some examples have been summarized by Jusko and Ko.^[4] The inhibition of basophil trafficking by 692 693 methylprednisolone and the furosemide-mediated inhibition of water reabsorption from the tubules 694 and collecting duct were assessed by Model I and II, respectively, while the stimulation of the cyclic 695 adenosine monophosphate (cAMP)-induced bronchodilation by the β-adrenergic receptor agonist 696 terbutaline was described by Model III. In a further example, it was shown that the increase in cAMP 697 by terbutaline activates the cellular membrane sodium-potassium pump, resulting in an increase of 698 efflux of potassium ions from the plasma into cells, an effect that can be described with Model IV.

699 These basic turnover models can be modified and/or extended to account for more complex physiological processes such as time-dependent production (k_{in}(t)),^[111] the rate of loss of cells 700 701 according to their lifespan^{[112],[113],[114]} and capacity limited processes such as nonlinear synthesis and degradation functions.^[115] Further, many physiological processes such as secretion of hormones and 702 703 gastric acid, gene expression, cardiac output and blood pressure are known to be subject to circadian rhythms, which might influence the pharmacokinetics and pharmacodynamics of various 704 drugs.^{[116],[117],[118]} Symmetric circadian rhythms have been described by trigonometric functions, such 705 706 as the cosine model introduced by Lew et al.,^[119] whereas asymmetric circadian rhythms have been 707 modelled with the addition of exponential, dual cosine or harmonic functions.^{[120],[111]} The detailed mathematical formalism around these functions has been summarized by Krzyzanski.^[121] 708

709

4.1.2 Signal transduction and feedback control indirect response models711

712 When a sequence of events takes place between receptor binding or activation and the observable effect, this is referred to as signal transduction and can involve signaling cascades, activation or 713 714 inhibition of secondary messengers, gene up- or down-regulation and mRNA transcription to functional proteins. By definition, every transduction process has two inherent attributes: the 715 716 transformation of the original signal and the introduction of a time-delay.^{[122],[123]} Depending on the 717 experimental time-scale, the time delay might or might not be discernable and in the latter case the 718 response is described by a transduction model with no delay, for example in the operational model of agonism introduced by Black and Leff.^[124] This model has been applied to describe the 719 720 pharmacokinetic/pharmacodynamic relationships of A1 adenosine, µ-opioid and 5-HT1A receptor agonists.^{[125],[126],[127],[128],[129]} However, in other cases the time delay produced by the transduction 721 722 process is significant and the mathematical models need to be adjusted accordingly. The most common 723 approach is the so-called transit compartment model (Fig. 8), which has been applied to the modeling of the genomic effects of corticosteroids, in this case known as the 5th generation model for 724 725 corticosteroids, as well as myelosuppression and hematologic toxicity in cancer chemotherapy.^{[130],[131],[132],[133]} 726

727 Most physiological processes are subject to feedback control and belong to the so-called autoregulation systems. The pharmacokinetic/pharmacodynamic (PK/PD) models that do not address 728 729 these auto-regulatory mechanisms fail to provide a complete insight of the drug-exposure relationship and it has been shown that this can lead to underestimation of the drug's potency.^[123] The feedback 730 731 control indirect response (FC IDR) models (see Figure 9) usually incorporate terms proportional to the 732 error signal itself, the integral and the derivative of the error signal in linear and, less commonly, in 733 nonlinear combinations. There are also FC IDR models which include an additional state, the "moderator" state, which feeds back to alter the synthesis or turnover of the response.^[134] Numerous 734 735 applications of PK/PD models incorporating feedback regulation mechanisms have been published in the literature.^{[132],[135],[136]} The example of (S)-citalopram, a widely used selective serotonin receptor 736 737 inhibitor (SSRI), is presented in detail in section 4.3.

738 739

4.2 Applications and case examples

740 741

4.2.1 Ibuprofen: antipyretic response

742 As mentioned in section 2.2.2, the antipyretic effect of ibuprofen resulting from the inhibition of 743 prostaglandin synthesis has been investigated in numerous clinical studies and an indirect response 744 model has been applied to fit the reported pharmacodynamic data. In a single-dose, placebocontrolled, double-blind and parallel-group trial by Walson et al.,^[137] the safety, efficacy, tolerability 745 746 and dose-effect relationships of ibuprofen products, formulated as a suspension at doses of 5 mg/kg 747 and 10 mg/kg to treat febrile children, were compared to liquid formulations of acetaminophen. The 748 patients (N=127) were split into groups according to their initial temperature and on whether 749 antibiotics were being administered concurrently. A positive dose-response relationship between 750 ibuprofen suspension 5 mg/kg and 10 mg/kg in the higher temperature (102.6-104°F), non-antibiotic 751 group was demonstrated, whereas in the lower temperature group (101-102.5°F) both doses were 752 equally effective. However, the authors pointed out that the plasma levels necessary for maximum 753 effective antipyresis of ibuprofen (approximately 10 mg/L) are achievable at doses even less than 5 754 mg/kg, implying a ceiling effect in the antipyretic response at doses of 5 mg/kg or higher.

Similar results in 178 children were observed by Wilson et al.^[138] In a single-dose, placebo-controlled study, during which age and initial temperature were considered as co-variates, both the 5 and 10 mg/kg doses were significantly superior to placebo, but not different from each other in terms of maximum reduction in temperature. However, it was concluded, based on the temperature at 6 hours after administration, the change of temperature from the baseline value and the percentage of efficacy, that the 10 mg/kg dose was more effective. The effect of the age and the initial temperature value on the magnitude of the pharmacological action was also emphasized.

In a double-blind, randomized, single-dose study of 5 and 10 mg/kg ibuprofen to treat febrile children
 (N=153) Brown et al.^[139] noted a dissociation between t_{max} and time of maximum temperature
 decrease and found no correlation between the extent of temperature change and plasma levels at

 $t_{R,max}$ or 6 hours post-administration. Further, there was no evidence that pretreatment with 765 766 antibiotics, race or gender influenced the antipyretic effect. By contrast, age and initial temperature 767 were shown to be co-variates. Interestingly, after compartmental pharmacokinetic analysis, only the pharmacodynamic, but not the pharmacokinetic parameters related to absorption (Cmax, tmax) and 768 769 elimination (k_{el}, t_{1/2}), were affected by the age of the child. In a subsequent paper, Brown et al. ^[140] 770 implemented an effect-compartment model coupled with a sigmoid E_{max} pharmacodynamic model to 771 describe the antipyretic effect of ibuprofen in children and further elaborated the model by adding a linear and/or sinusoidal cyclic function for the decrease in temperature as co-variates to fit their own 772 773 as well as previously reported data ^[138]. Values of the estimated sigmoidicity factor (γ) were 3.97 ± 0.58 774 and 4.27 ± 0.63 for ibuprofen 5 mg/kg and 10 mg/kg, respectively, implying that the dose-response 775 relationship for antipyresis in children might be steeper than for dental pain relief in adults.

776 Troconiz et al.^[47] reported a temporal disconnection between t_{max} after administration to febrile 777 children of 7 mg/kg ibuprofen as a suspension or as effervescent granules dosed at 200 or 400 mg (0.5 778 for the suspension and 1.9 hours for the effervescent granules) and time of maximum decrease in body 779 temperature (3 hours in both cases), suggesting that the formulation and its pharmacokinetic behavior 780 has little impact on the antipyretic effect of ibuprofen. The antipyretic response of non-steroidal antiinflammatory drugs (NSAIDs) has been attributed to their ability to inhibit the synthetic pathway of 781 782 prostaglandins, particularly of prostaglandin E₂ (PGE₂), via an indirect mechanism.^[141] The following 783 equation was derived to describe the pharmacodynamics of antipyresis by this mechanism:

784
$$\frac{dT}{dt} = k_{syn} \cdot \left(1 - E_{\max} \cdot \frac{C^{\gamma}}{C^{\gamma} + EC_{50}^{\gamma}}\right) - k_{out} \cdot T \qquad (26)$$

where dT/dt represents the rate of body temperature change with time, k_{syn} and k_{out} are the zeroorder and first-order rate constants for synthesis and degradation of the inflammation mediator (i.e. PGE₂), respectively, *T* is the body temperature, E_{max} is the maximum antipyretic effect, EC_{50} is the 788 drug plasma concentration (*C*) required to achieve half of the maximum effect and γ is the sigmoidicity 789 factor.

The proposed pharmacokinetic-pharmacodynamic model fitted the antipyretic profiles well. The estimated EC_{50} and k_{out} parameters were in agreement with those previously reported by Garg and Jusko (6.18 versus 10.2 mg/L for EC_{50} and 1.17 versus 0.89 h⁻¹ for k_{out}), who had also applied an indirect response model.^[142] The sigmoidicity factor was calculated to be 2.71 ± 0.18, suggesting a relatively flat dose-response curve. In contrast to previous studies, however, age and initial temperature did not elicit covariate effects. ^{[138],[143]}

796 Based solely on the differences in C_{max} and t_{max} between the suspension and the effervescent granule formulations, a delayed onset of drug action would be expected for the effervescent granules. 797 798 Nevertheless, the maximum antipyretic effect was similar and occurred at the same time for both 799 formulations. Importantly, an almost identical mean effect time course of 200 and 400mg of Ibuprofen 800 effervescent granules in febrile children was observed, implying that at least for this formulation there 801 was no significant clinical benefit with a dose increase (Fig. 10). Therefore, the authors concluded that 802 the formulation-dependent pharmacokinetic differences are mitigated by the response mechanism, 803 leading to similar pharmacodynamic responses for both formulations at both doses in febrile children.

804 Using a verified PBPK/PD model Cristofoletti and Dressman simulated the antipyretic response with 805 virtual trials of 2, 5, 7 or 10 mg/kg dosing of Ibuprofen suspension to 100 febrile children per trial in the age range of 2-11 years.^[58] In terms of maximum decrease in temperature from the baseline value, 806 807 the 5, 7 and 10 mg/kg doses were proven to be significantly superior to 2 mg/kg but not statistically 808 different from one another. A rather flat dose-response curve (with EC₅0≈6.18 mg/L) was confirmed 809 for the antipyretic effect in children. Under the assumption that the 7 and 10 mg/kg dose represent 810 the test and reference products, respectively, the test product would be bioinequivalent to the 811 reference in terms of C_{max} and AUC ratios (C_{max,T}/C_{max,R} and AUC_{max,T}/AUC_{max,R} around 0.7), but still therapeutically equivalent in children. This conclusion is supported by the data from Troconiz et al.^[47], 812

whose clinical trial demonstrated superimposable antipyretic profiles between ibuprofen suspension
7 mg/kg and effervescent granules 400 mg (normalized by children mean body weight as 11.8 mg/kg)
after administration to febrile children.

816 4.2.2 Rosuvastatin

817

Of the currently available 3-hydroxy-3-methylglutaryl coenzyme A reductase (HMG-CoA reductase) inhibitors, rosuvastatin is one of the most effective at lowering the low density lipoprotein (LDL) cholesterol. Mevalonic acid synthesis, which takes place in the liver, is catalyzed by HMG-CoA reductase and is the first irreversible stage of the cholesterol biosynthetic pathway.^{[144],[145],[146]}

822 A pharmacokinetic/pharmacodynamic model was developed to predict the response of rosuvastatin to different dosage regimens and identify differences in response between morning (at 07:00 a.m.) 823 and evening (at 06:00 p.m.) administration. For this purpose, Aoyama et al.^[147] used a two-824 825 compartment pharmacokinetic model with first order absorption and elimination from the central 826 compartment, which was then linked to a modified inhibitory indirect response pharmacodynamic 827 model describing the plasma concentrations of mevalonic acid (MVA). The model was further extended 828 by incorporating a time-dependent periodic function in the zero-order synthesis rate constant of mevalonic acid to account for the circadian rhythm, as introduced by Krzyzanski et al.^{[148],[149]} The model 829 830 is presented in Figure 11 and described by the following equations:

831
$$\frac{dR}{dt} = k_{in} \cdot \left(1 - \frac{C_p^{\gamma}}{C_p^{\gamma} + IC_{p50}^{\gamma}}\right) - k_{out} \cdot R \quad (27)$$

where R is the response, k_{in} is the time-dependent zero order rate constant for the increase in plasma MVA concentration, k_{out} is the first order rate constant for the decrease in plasma MVA concentration, C_p represents the plasma concentration of rosuvastatin, IC_{p50} is the plasma concentration at which k_{in} is reduced 50% and γ is the sigmoidicity factor. The time-dependent k_{in} to account for the circadian rhythm is defined as follows

837
$$k_{in} = k_m + k_{amp} \cdot \cos(2 \cdot \pi (t - tz)/24) \quad (28)$$

838 where k_m and k_{amp} represent the mean MVA synthesis and its amplitude rate constants, respectively, 839 and tz is the acrophase time, during which MVA is synthesized at the maximum rate. The following 840 function to describe the circadian rhythm of k_m was proposed by Krzyzanski et al.^[148]:

841
$$k_m = k_{out} \cdot IC - \frac{k_{amp} \cdot k_{out}^2}{k_{out}^2 + (2\pi/24)^2} \cdot \left[\cos\left(\frac{2 \cdot \pi \cdot (tz)}{24}\right) - \left(\frac{2 \cdot \pi}{24 \cdot k_{out}}\right) \cdot \sin\left(\frac{2 \cdot \pi \cdot (tz)}{24}\right)\right]$$
(29)

842 where IC is the initial plasma MVA concentration measured at 6 a.m., set to 4.32 ng/ml.

843 Application of the time course of rosuvastatin and mevalonic acid plasma concentration to the model enabled an adequate prediction of the clinical data reported by Martin et al.^[150] A higher reduction 844 845 ratio of 7.7% in the area under the plasma MVA concentration-time curves over 24 hours at steady 846 state (AUEC₀₋₂₄) was observed after administration in the evening. Furthermore, sensitivity analysis on 847 the pharmacokinetic parameters showed that changes in the pharmacokinetics have a greater effect 848 on the AUEC₀₋₂₄ reduction ratio after morning than after evening administration. This was attributed 849 to the circadian rhythm, with the acrophase time estimated to be 15.5 hours. The authors concluded that evening administration of rosuvastatin might be useful in clinical practice.^[147] The main limitation 850 851 of the model is that it is based only on the mean plasma pharmacokinetic and pharmacodynamic data. 852 Therefore, it does not address the concentration at the effect site, which is the liver and not the 853 plasma, or the inter-subject variability. Most importantly, the use of only one mean PK/PD data set 854 raises questions about the identifiability of the estimated parameters and caution should be exercised 855 in drawing conclusions about the validity of this model.

Since the liver is the effect site for the statins, uptake into the liver is an important factor in their efficacy. Multiple transporters of the family of the organic anion transporting polypeptide (OATP) family are abundant in the liver, facilitating the active hepatic uptake of endogenous substances and xenobiotics, including statins, from sinusoidal blood.^{[151],[152],[153],[154],[155]} Rosuvastatin is a substrate of the organic anion transporting polypeptide 1B1, 1B2, 1B3, 1A2 and the sodium-dependent

taurocholate co-transporting polypeptide.^{[151],[156]} The expression of OATP1B1 on the sinusoidal 861 membrane of human hepatocytes is encoded by the gene SLCO1B1, which is subjected to single-862 nucleotide polymorphisms (SNPs). As already demonstrated for paravastatin, pitavastatin and 863 864 simvastatin, such polymorphisms are associated with reduced OATP1B1 in vitro activity and markedly increased plasma concentrations.^{[157],[158],[159],[160],[161]} Pasanen et al.^[158] investigated the effect of 865 866 SLCO1B1 polymorphism on the pharmacokinetics of atorvastatin and rosuvastatin, after oral 867 administration in 32 healthy volunteers, with the following genotypes: SLCO1B1 c.521CC (n=4), 868 SLCO1B1 c.521CT (n=12), SLCO1B1 c.521TT (wild type, n=16). Significant increases in the AUC_{0-48 h} and 869 C_{max} (65% and 79%, respectively) in *SLCO1B1* c.521CC subjects compared to the reference genotype, 870 SLCO1B1 c.521TT, were observed. By contrast, increases in the AUC_{0-48 h} (144% increase), but not the 871 C_{max}, were reported after administration of atorvastatin. This study implies that the reduced OATP1B1-872 mediated hepatic uptake of rosuvastatin due to SLCO1B1 polymorphism results in an increased risk of 873 a reduced cholesterol-lowering effect as well as adverse effects such as myopathy and/or 874 rhabdomyolysis.

Based on the model of Aoyama et al., [147] a full PBPK/PD model was built in the SimCyp Simulator® by 875 Rose et al.^[162] to investigate the impact of polymorphic hepatic uptake (OATP1A1, OATP1B4) and efflux 876 877 transposers (BcRP, MRP2) on the disposition, pharmacologic and toxic effects of rosuvastatin. First, 878 plasma concentrations were linked to the cholesterol-lowering effect of rosuvastatin, according to the 879 plasma AUC of MVA. The simulations performed with the PBPK/PD model showed a large increase in 880 the mean plasma AUC infinity (AUC∞) of rosuvastatin by 63% and 111% for the SLCO1B1 c.521CT and 881 SLCO1B1 c.521CC, respectively, compared to the wild type (SLCO1B1 c.521TT). Similarly, a significant 882 increase in MVA plasma AUC of 30% and 35% for the same genotypes was observed. However, the 883 hepatic unbound intracellular water concentration (Cu_{IW}) of rosuvastatin, which was predicted by a 884 permeability limited liver model, was considered to be a more relevant driver of its pharmacodynamic 885 effect. Interestingly, only a slight decrease in Cu_{IW} based AUC_{∞} of 5.7% and 9.6%, with a parallel decrease in MVA plasma AUC of 3.1% and 5.8% were reported for the heterozygote and homozygote, 886

respectively. The latter findings are in agreement with a number of studies showing that OATP1B1 c.521T>C SNP has either no or only a slight effect on the cholesterol-lowering response to statins,^{[163],[164],[165]} and that when plasma concentrations were used as the input, the results were misleading.

With regard to toxic effects, the effect of genetic polymorphism on rosuvastatin-mediated myopathy was investigated by prediction of muscle concentrations using a perfusion-limited model. A strong correlation between plasma concentrations and the risk of muscle-related adverse effects was observed. Thus, in contrast to the results for the cholesterol-lowering effect of rosuvastatin, the plasma concentration appears to be a good surrogate for the concentration at the muscle when assessing the risk of statin-induced muscle toxicity in individuals with polymorphic hepatic uptake transporter activity. This result was also in agreement with an already published study.^[166]

898 High inter-individual variability among the different genotypes, limited availability of accurate in vitro 899 data and/or published clinical studies at different dose levels as well as incomplete understanding of 900 the impact of transporters on pharmacokinetics and/or pharmacodynamics, are some of the 901 limitations which restrict the robustness of the models for rosuvastatin and their confidence in 902 simulating different clinical scenarios. Despite these limitations, rosuvastatin serves as a useful case 903 example to demonstrate the potential of linking PBPK with PD model to enhance physiological 904 understanding and improve the ability to assess the impact of transporters on the pharmacologic 905 and/or toxic response. Of particular importance was the finding that, in some instances, parameters 906 other than the plasma concentration are appropriate indicators of the therapeutic and/or toxic effect. 907 This example illustrates that implementation of (PB)PK/PD models (even on an exploratory basis) can 908 provide valuable information during clinical drug development and significantly contribute to the 909 clinical ramifications of genetic polymorphism and facilitate an optimal dosing regimen.

910 4.2.3 Escitalopram

911

Selective serotonin reuptake inhibitors (SSRIs), such as escitalopram, block the neuronal reuptake of serotonin (5-HT), resulting in increased neurotransmitter concentration at the terminal and somatodendritic areas. However, the auto-receptors 5-HT_{1A} and 5-HT_{1B}, which regulate the 5-HT release from neurons by negative feedback control, are also situated at the terminal and somato-dendritic neuronal parts, respectively (Fig. 12).^[167] Intracerebral microdialysis can be used to measure the extracellular concentration of 5-HT and thus its concentration at the site of action.^{[168],[169]}

Bundgaard et al.^[170] developed an indirect response PK/PD model for escitalopram, including a 918 919 moderator state (tolerance model) to account for the auto-inhibitory feedback. For this purpose, 920 different doses of escitalopram were administered intravenously at a constant infusion rate over 60 921 minutes in four groups (vehicle, 2.5, 5 and 10 mg/kg) of six male Sprague-Dawley rats and the response 922 was expressed as the change in extracellular 5-HT concentration. A two-compartment 923 pharmacokinetic model with first order elimination from the main compartment was used to fit the 924 individual mean unbound plasma concentration-time profiles for each dose group and the predicted 925 profiles were used as the input to drive the pharmacodynamic model. A type II basic indirect response 926 model was implemented to describe the inhibition of 5-HT reuptake. In this model, the increase in the 927 response, R, over the baseline value R₀, feeds back to the moderator compartment and stimulates the production of the moderator, M. As a simplifying approximation, the rates in and out of M are 928 929 described by a first-order rate constant k_{tol} . An increase in M induces a negative feedback on the 930 generation of the response and thus enables the baseline value to be reestablished. The model is 931 illustrated in Figure 13 and described by the following equations:

932
$$\frac{dR}{dt} = \frac{k_{in}}{M} - k_{out} \cdot R \cdot I(C_p) \quad (30)$$

933
$$\frac{dM}{dt} = k_{tol} \cdot R - k_{tol} \cdot M \quad (31)$$

934
$$I(C_p) = 1 - \frac{I_{max} \cdot C_p^n}{IC_{50}^n + C_p^n} \quad (32)$$

where R, M and C_p represent the response, the moderator and the escitalopram unbound plasma concentration respectively, I_{max} , IC_{50} and n are the maximum inhibitory effect, the potency and sigmoidicity factor respectively, and k_{in} , k_{out} and k_{tol} represent the turnover rate, fractional turnover rate and feedback rate constants, respectively (see Fig.13). By setting equations 30 and 31 equal to zero, the initial baseline conditions are obtained:

$$k_{in} = k_{out} \cdot R_0^2 \quad (33)$$

941
$$R_0 = M_0 = \sqrt{\frac{k_{in}}{k_{out}}}$$
 (34)

942 The feedback control model fitted the response-time data well. Between unbound plasma 943 concentration and 5-HT response, a distinct time-delay was observed for all doses, leading to a 944 counter-clockwise hysteresis loop. The development of tolerance was confirmed by the fact that the 945 terminal phases of the hysteresis loops were not superimposable as a function of dose: the higher dose 946 groups exhibited a lower response at the same concentration. Based on one-way analysis of variance 947 (ANOVA) and post hoc analysis, maximal increases in 5-HT extracellular levels reached 337%, 424% and 948 456% of the baseline and the levels remained elevated for 135, 175 and 235 minutes at the 2.5, 5 and 949 10 mg/kg doses, respectively. Despite the significant differences in plasma concentrations, the basal 950 response value was recovered within 360 min following the administration of all tested doses. In fact, 951 neither the duration nor the magnitude of the response increased when the dose was increased from 952 5 to 10 mg/kg. These findings are in agreement with previous studies in rats, in which increasing the dose of escitalopram exhibited a ceiling effect in the extracellular levels of 5-HT in the frontal cortex, 953 as measured by microdialysis.^{[171],[172]} 954

The results from this study established the high potency (IC₅₀= 4.4 μ g/L) of escitalopram, with almost complete (I_{max}= 0.9) inhibition of reuptake. A fast neuronal 5-HT reuptake with a half-life of less than 5 minutes ($t_{1/2k_{out}}$) was reported, whereas the half-life for the development of tolerance, $t_{1/2k_{tol}}$ was estimated at 10 hours. The importance of incorporating a moderator state to account for the 959 physiological homeostatic autoregulation mechanisms was demonstrated by comparison of the pharmacodynamic parameters of this more mechanistic model with the conventional effect-960 compartment model. The effect-compartment model predicted higher EC_{50} values at increased doses, 961 962 which was inconsistent with the physiological response. In addition, Zhang and D'Argenio^[123] used the 963 same data sets to compare the performance of the basic model II inhibitory model with and without 964 the addition of proportional and proportional-plus-integral feedback gain. When the feedback was 965 omitted, the drug's potency was underestimated, while the model with the proportional-plus-integral 966 feedback gain performed the best (lowest Akaike information criterion value).

967 These findings not only highlight the usefulness of implementing feedback control mechanisms in 968 pharmacodynamic models, but also the importance of assessing the PK/PD at multiple doses. It is 969 evident that when the autoregulation of the pharmacodynamic response is not taken into account, the 970 evaluation of in vivo potency can lead to an underestimation of drug's potency and application of 971 unnecessarily high doses. Additionally, feedback control models may be useful for the comparison of 972 the pharmacodynamic behavior among SSRIs, to improve understanding of their antidepressant 973 effects and as a guide to set effective plasma concentrations in clinical practice.

- 5 974
- 975

Outlook and concluding remarks

976 This review describes the large variety of pharmacokinetic/pharmacodynamic modeling approaches 977 available to predict dose-concentration-effect relationships and to simulate various clinical scenarios. 978 Models incorporating a physiological understanding of the underlying mechanism(s) of action of the 979 drug and progression of disease can serve as powerful tools for exploring and predicting clinical drug 980 product performance. Provided such models are adequately validated, they can also be implemented 981 with confidence to drive model-informed decisions during drug development as well as at the 982 regulatory level.

983 An even more complete understanding of a drug's therapeutic value would be possible if dose-984 concentration-adverse reactions relationships were to be simultaneously established through toxicokinetic/toxicodynamic models, so that not only efficacy, but also safety can be evaluated. This is
important, since dose-response curves may differ significantly between the therapeutic and adverse
effects in different patient populations as well as among different indications of the same drug.

988 A current limitation of mechanistic models is that their complexity often leads to issues of identifiability 989 and reproducibility of parameters. The commercially available physiologically based pharmacokinetic 990 models are often implemented with mostly (or only) literature data. In these models the number of 991 parameters is often far greater than would be required for application of classical compartmental 992 models and it may be difficult to acquire reliable values for some parameters. The advent of more 993 sophisticated analytical techniques such as microdialysis will promote a better understanding of the 994 time profile of drug concentration at the effect site. In the meantime, to ensure maximum quality and 995 to facilitate the interpretation of PK/PD models, transparency in the parameter values applied in the 996 model, as well as in the underlying assumptions and the derived equations, together with 997 harmonization based on good coding practice (GCP), is essential.

998 Once there is enough confidence in the translatability, estimation and prediction of preclinical and 999 clinical PK/PD and systems pharmacology models, a move towards linking them with biorelevant in 1000 vitro tools to guarantee therapeutic equivalence will be another key step forward in the drive to link 1001 the laboratory to the patient, which seems not only promising, but also imminent. Bridging the gap 1002 between in vitro, in vivo and in silico methods by applying the Quality by Design (QbD) and the 1003 Biopharmaceutics Risk Assessment Roadmap (BioRAM),^{[173],[174]} will allow pharmaceutical scientists to 1004 correctly assess the relative impact of formulation, dose and dosing interval during development of 1005 new drugs.

For the formulation scientist, modeling and simulation used in this way will assist in the selection of the most appropriate dosage form and to set formulation targets, knowing to what extent the formulation can be expected to steer the *in vivo* performance of the drug product. For the clinician, the approach helps to identify the dosing strategy which optimizes the efficacy/safety ratio.

For the analyst, modeling and simulation can provide guidance in setting clinically relevant dissolution specifications, taking into account not only which formulation factors steer the drug plasma concentration (critical quality attributes) but also how any differences in these will translate in the clinical outcome. In this context, robust PK/PD modeling approaches will play an essential role in model-informed drug development.

1015 Finally, from a regulatory decision-making point of view, a seamless description of the relationship 1016 between the pharmacokinetic and pharmacodynamic characteristics of a drug together with a 1017 knowledge of how, and to what extent, formulation and formulation performance can influence the 1018 PK and PD, provides an excellent, clinically relevant basis for an integrated approach to assessing 1019 applications for drug approval. Currently, pharmacodynamics considerations are taken into account 1020 in the approval of labeling of new drug products, for example, whether taking the drug before vs. after 1021 a meal will influence efficacy. There is also a thrust towards virtual bioequivalence, for example using 1022 PBPK modeling to determine whether a change in the dissolution characteristics will impact the plasma 1023 profile significantly. A logical further step would be to combine these two approaches to optimize the 1024 approval process. Foreseen is a scenario in which the release testing in the laboratory reflects the 1025 release in the target patient population(s), the data are combined with verified PBPK models tailored 1026 to the target population(s) and then translated with PK/PD modeling into a prediction of the clinical 1027 outcome. This scenario would not only provide sponsors as well as the regulatory authority with more 1028 flexibility in the approval procedure, without sacrificing efficacy or safety, but also be a way forward 1029 to move effectively towards a more personalized medicine concept.

- 1030 6 Acknowledgements
- 1031

1032 This work was supported by the European Union's Horizon 2020 Research and Innovation

1033 Programme under grant agreement No 674909 (PEARRL)

1034 1035	7	References
1036 1037	1.	Levy G. Relationship between rate of elimination of tubocurarine and rate of decline of its pharmacological activity. <i>Br J Anaesth</i> 1964; 36(11): 694–695.
1038 1039	2.	Segre G. Kinetics of interaction between drugs and biological systems. <i>Farmaco Sci</i> 1968; 23(10): 907–18.
1040 1041	3.	Holford NHG, Sheiner LB. Understanding the Dose-Effect Relationship: Clinical Application of Pharmacokinetic-Pharmacodynamic Models. <i>Clin Pharmacokinet</i> 1981; 6(6): 429–453.
1042 1043	4.	Jusko WJ, Ko HC. Physiologic indirect response models characterize diverse types of pharmacodynamic effects. <i>Clin Pharmacol Ther</i> 1994; 56(4): 406–419.
1044 1045	5.	Rowland M <i>et al.</i> Physiologically based pharmacokinetics is impacting drug development and regulatory decision making. <i>CPT Pharmacometrics Syst Pharmacol</i> 2015; 4(6): 313–315.
1046 1047	6.	Jusko WJ. Moving from Basic Toward Systems Pharmacodynamic Models. <i>J Pharm Sci</i> 2013; 102(9): 2930–2940.
1048 1049 1050	7.	Androulakis IP. Systems engineering meets quantitative systems pharmacology: from low- level targets to engaging the host defenses. <i>Wiley Interdiscip Rev Syst Biol Med</i> 2015; 7(3): 101–112.
1051 1052	8.	Galeazzi RL <i>et al.</i> Relationship between the pharmacokinetics and pharmacodynamics of procainamide. <i>Clin Pharmacol Ther</i> 1976; 20(3): 278–89.
1053 1054	9.	Frazier EP <i>et al</i> . Effects of gender, age and hypertension on β-adrenergic receptor function in rat urinary bladder. <i>Naunyn Schmiedebergs Arch Pharmacol</i> 2006; 373(4): 300–309.
1055 1056	10.	Wright DFB <i>et al.</i> Understanding the time course of pharmacological effect: A PKPD approach. <i>Br J Clin Pharmacol</i> 2011; 71(6): 815–823.
1057 1058 1059 1060	11.	U.S. Department of Health and Human Services <i>et al.</i> Physiologically Based Pharmacokinetic Analyses — Format and Content Guidance for Industry (Draft guidance). 2016. Available at: http://www.fda.gov/Drugs/GuidanceComplianceRegulatoryInformation/Guidances/default.ht m. Accessed May 7, 2018.
1061 1062 1063 1064	12.	Medicines Agency E. Guideline on the qualification and reporting of physiologically based pharmacokinetic (PBPK) modelling and simulation (Draft). 2016. Available at: http://www.ema.europa.eu/docs/en_GB/document_library/Scientific_guideline/2016/07/WC 500211315.pdf. Accessed May 7, 2018.
1065 1066 1067 1068 1069	13.	U.S. Department of Health and Human Services <i>et al.</i> Clinical Drug Interaction Studies — Study Design, Data Analysis, and Clinical Implications Guidance for Industry (Draft Guidance). 2017. Available at: http://www.fda.gov/Drugs/GuidanceComplianceRegulatoryInformation/Guidances/default.ht m. Accessed May 7, 2018.
1070 1071 1072	14.	Medicines Agency E. Guideline on the investigation of drug interactions. 2012. Available at: http://www.ema.europa.eu/docs/en_GB/document_library/Scientific_guideline/2012/07/WC 500129606.pdf. Accessed May 7, 2018.
1073	15.	EMA. Committee for Medicinal Products for Human use (CHMP): Guidelin on the role of

- 1074pharmacokinetics in the development of medicinal products in the paediatric population1075DRAFT AGREED BY EFFICACY WORKING PARTY GUIDELINE ON GUIDELINE ON THE ROLE OF1076PHARMACOKINE. In: EMEA/CHMP/EWP/147013/2004 C, 2006, eds., 2006. Available at:1077http://www.emea.europa.eu. Accessed May 7, 2018.
- Center for Drug Evaluation and Research (CDER), Center for Biologics Evaluation and Research (CBER), Food and Drug Administration. Guidance for Industry: General Clinical Pharmacology Considerations for Pediatric Studies for Drugs and Biological Products . 2014. Available at: http://www.fda.gov/Drugs/GuidanceComplianceRegulatoryInformation/Guidances/default.ht m. Accessed May 7, 2018.
- 1083 17. U.S Department of Health and Human Services F and DAC for DE and R (CDER). Topical
 1084 dermatological corticosteriods: in vivo bioequivalence. FDA 1995. In: , 1995. Available at:
 1085 https://www.fda.gov/downloads/Drugs/GuidanceComplianceRegulatoryInformation/Guidanc
 1086 es/UCM070234.pdf. Accessed May 7, 2018.
- 1087 18. Lionberger RA. FDA Critical Path Initiatives: Opportunities for Generic Drug Development.
 2008.
- 108919.Chen ML *et al.* Challenges and opportunities in establishing scientific and regulatory standards1090for determining therapeutic equivalence of modified-release products: Workshop summary1091report. Clin Ther 2010; 32(10): 1704–1712.
- 1092 20. U.S. Department of Health and Human Services, Food and Drug Administration, Center
 1093 forDrug Evaluation and Research. Individual Product Bioequivalence Recommendation—
 1094 Methylphenidate hydrochloride (Draft guidance). In: *Guidance for Industry: Bioequivalence* 1095 *Recommendations for Specific Products. May 2007.*
- 1096 http://www.fda.gov/downloads/Drugs/GuidanceCompliance
- 1097 *RegulatoryInformation/Guidances/ucm072872.pdf.*, 2017. Available at:
- 1098 https://www.fda.gov/downloads/Drugs/GuidanceComplianceRegulatoryInformation/Guidanc 1099 es/UCM581432.pdf. Accessed May 7, 2018.
- 110021.U.S. Department of Health and Human Services, Food and Drug Administration, Center1101forDrug Evaluation and Research. Individ- ual Product Bioequivalence Recommendation—1102Budesonide (Draft guidance). In: Guidance for Industry: Bioequivalence Recommendations for1103Specific Products. May 2007. http://www.fda.gov/downloads/Drugs/GuidanceCompliance1104RegulatoryInformation/Guidances/ucm072872.pdf., 2014. Available at:1105https://www.fda.gov/downloads/Drugs/GuidanceCompliance1106es/UCM426317.pdf. Accessed May 7, 2018.
- 110722.U.S. Department of Health and Human Services, Food and Drug Administration, Center1108forDrug Evaluation and Research. Individ- ual Product Bioequivalence Recommendation—1109Zolpidem (Final guidance). In: Guidance for Industry: Bioequivalence Recommendations for1110Specific Products. May 2007. http://www.fda.gov/downloads/Drugs/GuidanceCompliance1111RegulatoryInformation/Guidances/ucm072872.pdf., 2011. Available at:
- 1112https://www.fda.gov/downloads/Drugs/GuidanceComplianceRegulatoryInformation/Guidanc1113es/UCM175029.pdf. Accessed May 7, 2018.
- 111423.CFR Code of Federal Regulations Title 21. Available at:1115https://www.accessdata.fda.gov/scripts/cdrh/cfdocs/cfcfr/CFRSearch.cfm?fr=314.3. Accessed1116August 16, 2018.
- 111724.Forester W *et al.* The onset and magnitude of the contractile response to commonly used1118digitalis glycosides in normal subjects. *Circulation* 1974; 49(3): 517–21.

1119 1120	25.	Shapiro W <i>et al.</i> Relationship of plasma digitoxin and digoxin to cardiac response following intravenous digitalization in man. <i>Circulation</i> 1970; 42(6): 1065–72.
1121 1122	26.	Holford NHG, Sheiner LB. Kinetics of pharmacologic response. <i>Pharmacol Ther</i> 1982; 16(2): 143–166.
1123 1124	27.	Louizos C <i>et al.</i> Understanding the hysteresis loop conundrum in pharmacokinetic/pharmacodynamic relationships. <i>J Pharm Pharm Sci</i> 2014; 17(1): 34–91.
1125 1126	28.	Evans MA <i>et al</i> . Pharmacokinetic and pharmacodynamic modelling with pancuronium. <i>Eur J Clin Pharmacol</i> 1984; 26(2): 243–50.
1127 1128	29.	Schwartz JB <i>et al.</i> Pharmacodynamic modeling of verapamil effects under steady-state and nonsteady-state conditions. <i>J Pharmacol Exp Ther</i> 1989; 251(3): 1032–8.
1129 1130	30.	Whiting B <i>et al</i> . Modelling theophylline response in individual patients with chronic bronchitis. <i>Br J Clin Pharmacol</i> 1981; 12(4): 481–7.
1131 1132 1133	31.	Holford NH <i>et al.</i> The effect of quinidine and its metabolites on the electrocardiogram and systolic time intervals: concentrationeffect relationships. <i>Br J Clin Pharmacol</i> 1981; 11(2): 187–95.
1134 1135	32.	Gabrielsson JL <i>et al.</i> Analysis of pethidine disposition in the pregnant rat by means of a physiological flow model. <i>J Pharmacokinet Biopharm</i> 1986; 14(4): 381–395.
1136 1137 1138	33.	Björkman S <i>et al.</i> Comparative physiological pharmacokinetics of fentanyl and alfentanil in rats and humans based on parametric single-tissue models. <i>J Pharmacokinet Biopharm</i> 1994; 22(5): 381–410.
1139 1140 1141	34.	Lemmens HJM <i>et al.</i> Pharmacokinetic-pharmacodynamic modeling in drug development: Application to the investigational opioid trefentanil. <i>Clin Pharmacol Ther</i> 1994; 56(3): 261– 271.
1142 1143	35.	Torres-López JE <i>et al.</i> Pharmacokinetic-pharmacodynamic modeling of the antinociceptive effect of diclofenac in the rat. <i>J Pharmacol Exp Ther</i> 1997; 282(2): 685–90.
1144 1145	36.	Morrison RA <i>et al.</i> Isosorbide dinitrate kinetics and dynamics after intravenous, sublingual, and percutaneous dosing in angina. <i>Clin Pharmacol Ther</i> 1983; 33(6): 747–756.
1146 1147	37.	Mould DR <i>et al.</i> Simultaneous modeling of the pharmacokinetics and pharmacodynamics of midazolam and diazepam. <i>Clin Pharmacol Ther</i> 1995; 58(1): 35–43.
1148 1149	38.	Kelman AW, Whiting B. Modeling of drug response in individual subjects. <i>J Pharmacokinet Biopharm</i> 1980; 8(2): 115–30.
1150 1151	39.	Ryan AR. Tubocurarine administration based upon its disappearance and accumulation curves in anaesthetized man. <i>BJA Br J Anaesth</i> 1964; 36(5): 287–294.
1152 1153	40.	Gibaldi M <i>et al.</i> Kinetics of the elimination and neuromuscular blocking effect of d- tubocurarine in man. <i>Anesthesiology</i> 1972; 36(3): 213–218.
1154 1155	41.	Hull CJ <i>et al.</i> A pharmacodynamic model for pancuronium. <i>Br J Anaesth</i> 1978; 50(11): 1113– 23.
1156 1157	42.	Sheiner LB <i>et al.</i> Simultaneous modeling of pharmacokinetics and pharmacodynamics: application to d-tubocurarine. <i>Clin Pharmacol Ther</i> 1979; 25(3): 358–71.

43. Stanski DR et al. Pharmacokinetics and Pharmacodynamics of d-Tubocurarine during Nitrous 1158 1159 Oxide–Narcotic and Halothane Anesthesia in Man. Anesthesiology 1979; 51(3): 235–241. Goat VA et al. The effect of blood flow upon the activity of gallamine triethiodide. Br J 1160 44. 1161 Anaesth 1976; 48(2): 69-73. 45. 1162 Blume H, Mutschler M. Bioäquivalenz, Qualitätsbewertung wirkstoffgleicher 1163 Fertigarzneimittel, Teil I/II, Isosorbiddinitrat 6. Ergänzungslieferung, Govi-Verlag 1164 Pharmazeutischer Verlag, Frankfurt/Main-Eschborn. 1996. 1165 46. Holford NH, Sheiner LB. Pharmacokinetic and pharmacodynamic modeling in vivo. Crit Rev 1166 Bioeng 1981; 5(4): 273-322. 1167 47. Trocóniz IF et al. Pharmacokinetic-Pharmacodynamic Modelling of the Antipyretic Effect of 1168 Two Oral Formulations of Ibuprofen. Clin Pharmacokinet 2000; 38(6): 505–518. 1169 48. Lon H-K et al. Pharmacokinetic/pharmacodynamic modeling in inflammation. Crit Rev Biomed 1170 Eng 2012; 40(4): 295–312. 1171 49. Jain AK et al. Analgesic efficacy of low-dose ibuprofen in dental extraction pain. 1172 *Pharmacotherapy* 6(6): 318–22. 50. 1173 Hersh E V. et al. Single dose and multidose analgesic study of ibuprofen and meclofenamate 1174 sodium after third molar surgery. Oral Surgery, Oral Med Oral Pathol 1993; 76(6): 680–687. 1175 51. Seymour RA et al. Post-operative dental pain and analgesic efficacy. Part II. Analgesic usage 1176 and efficacy after dental surgery. Br J Oral Surg 1983; 21(4): 298–303. 1177 52. Laska EM et al. The correlation between blood levels of ibuprofen and clinical analgesic 1178 response. Clin Pharmacol Ther 1986; 40(1): 1-7. 53. Li H et al. Modeling the Onset and Offset of Dental Pain Relief by Ibuprofen. J Clin Pharmacol 1179 1180 2012; 52(1): 89–101. 1181 54. Schou S et al. Analgesic dose-response relationship of ibuprofen 50, 100, 200, and 400 mg 1182 after surgical removal of third molars: a single-dose, randomized, placebo-controlled, and 1183 double-blind study of 304 patients. J Clin Pharmacol 1998; 38(5): 447-54. 1184 55. Mcquay HJ et al. Ibuprofen compared with ibuprofen plus caffeine after third molar surgery. P 1185 1996; 66: 247-251. 1186 56. McQuay HJ, Moore RA. Dose-response in direct comparisons of different doses of aspirin, 1187 ibuprofen and paracetamol (acetaminophen) in analgesic studies. Br J Clin Pharmacol 2007; 1188 63(3): 271-278. 1189 57. Lemmens H et al. Pharmacokinetics-pharmacodynamics (PK/PD) of Ibuprofen in Dental Pain. J 1190 Clin Pharmacol 1996; 36(9): 856. 1191 58. Cristofoletti R, Dressman JB. Use of Physiologically Based Pharmacokinetic Models Coupled 1192 with Pharmacodynamic Models to Assess the Clinical Relevance of Current Bioequivalence 1193 Criteria for Generic Drug Products Containing Ibuprofen. J Pharm Sci 2014; 103(10): 3263-1194 3275. 1195 59. Walson PD, Galletta G, Braden NJ AL. Ibuprofen, acetaminophen and placebo treatment of 1196 febrile children. *Clin Pharmacol Ther* 1989; 46(1): 9–17. 1197 60. Li H et al. Modeling the Onset and Offset of Dental Pain Relief by Ibuprofen. J Clin Pharmacol

1198 2012; 52(1): 89–101.

- 119961.Cristofoletti R, Dressman JB. Bridging the Gap Between In Vitro Dissolution and the Time1200Course of Ibuprofen-Mediating Pain Relief. J Pharm Sci 2016; 105(12): 3658–3667.
- 1201 62. Jonker JW, Schinkel AH. Pharmacological and Physiological Functions of the Polyspecific
 1202 Organic Cation Transporters: OCT1, 2, and 3 (SLC22A1-3). *J Pharmacol Exp Ther* 2003; 308(1):
 1203 2–9.
- de Lange ECM, Danhof M. Considerations in the Use of Cerebrospinal Fluid Pharmacokinetics
 to Predict Brain Target Concentrations in the Clinical Setting. *Clin Pharmacokinet* 2002; 41(10):
 691–703.
- 1207 64. Lee G *et al.* Drug transporters in the central nervous system: brain barriers and brain
 1208 parenchyma considerations. *Pharmacol Rev* 2001; 53(4): 569–96.
- 1209 65. De Boer AG *et al.* The role of drug transporters at the blood-brain barrier. *Annu Rev*1210 *Pharmacol Toxicol* 2003; 43: 629–56.
- 1211 66. Letrent SP *et al.* Effect of GF120918, a Potent P-glycoprotein Inhibitor, on Morphine
 1212 Pharmacokinetics and Pharmacodynamics in the Rat. *Pharm Res* 1998; 15(4): 599–605.
- 121367.Suzuki N *et al.* Intrathecal morphine-3-glucuronide does not antagonize spinal antinociception1214by morphine or morphine-6-glucuronide in rats. *Eur J Pharmacol* 1993; 249(2): 247–50.
- 1215 68. Ouellet DM, Pollack GM. Effect of prior morphine-3-glucuronide exposure on morphine
 1216 disposition and antinociception. *Biochem Pharmacol* 1997; 53(10): 1451–7.
- 121769.Hewett K *et al.* Lack of effect of morphine-3-glucuronide on the spinal antinociceptive actions1218of morphine in the rat: an electrophysiological study. *Pain* 1993; 53(1): 59–63.
- 1219 70. Xie R *et al.* The role of P-glycoprotein in blood-brain barrier transport of morphine:
 1220 transcortical microdialysis studies in mdr1a (-/-) and mdr1a (+/+) mice. *Br J Pharmacol* 1999;
 1221 128(3): 563–568.
- 1222 71. de Lange EC. *et al.* Methodological considerations of intracerebral microdialysis in
 1223 pharmacokinetic studies on drug transport across the blood–brain barrier. *Brain Res Rev*1224 1997; 25(1): 27–49.
- 1225 72. Hammarlund-Udenaes M. The use of microdialysis in CNS drug delivery studies:
 1226 Pharmacokinetic perspectives and results with analgesics and antiepileptics. *Adv Drug Deliv*1227 *Rev* 2000; 45(2–3): 283–294.
- 1228 73. Hammarlund-Udenaes M *et al.* Drug equilibration across the blood-brain barrier-1229 pharmacokinetic considerations based on the microdialysis method. *Pharm Res* 1997; 14(2):
 1230 128–34.
- 123174.Mager DE *et al.* Diversity of mechanism-based pharmacodynamic models. *Drug Metab Dispos*12322003; 31(5): 510–518.
- 1233 75. Danhof M *et al.* Mechanism-Based Pharmacokinetic-Pharmacodynamic Modeling: Biophase
 1234 Distribution, Receptor Theory, and Dynamical Systems Analysis. *Annu Rev Pharmacol Toxicol* 1235 2007; 47(1): 357–400.
- 123676.Hong Y et al. Population pharmacodynamic modelling of aspirin- and ibuprofen-induced1237inhibition of platelet aggregation in healthy subjects. Clin Pharmacokinet 2008; 47(2): 129–

- 1238 137.
- 1239 77. Yamamoto K *et al.* Pharmacodynamics analysis of antiplatelet effect of aspirin in the literature
 1240 Modeling based on inhibition of cyclooxygenase in the platelet and the vessel wall
 1241 endothelium. *Jpn J Hosp Pharm* 1996; 22: 133–141.
- 1242 78. Gisleskog PO *et al.* A model for the turnover of dihydrotestosterone in the presence of the
 1243 irreversible 5 alpha-reductase inhibitors GI198745 and finasteride. *Clin Pharmacol Ther* 1998;
 1244 64(6): 636–647.
- 1245 79. Katashima M *et al.* Pharmacokinetic and pharmacodynamic study of a new nonsteroidal 5
 1246 alpha-reductase inhibitor, 4-[3-[3-[Bis(4-isobutylphenyl)methylamino]benzoyl]-1H-indol-1-yl]1247 butyr ic acid, in rats. *J Pharmacol Exp Ther* 1998; 284(3): 914–920.
- 124880.Abelo A *et al.* A turnover model of irreversible inhibition of gastric acid secretion by1249omeprazole in the dog. J Pharmacol Exp Ther 2000; 295(2): 662–669.
- 125081.Katashima M *et al.* Comparative pharmacokinetic/pharmacodynamic analysis of proton pump1251inhibitors omeprazole, lansoprazole and pantoprazole, in humans. *Eur J Drug Metab*1252*Pharmacokinet* 1998; 23(1): 19–26.
- 125382.Puchalski TA *et al.* Pharmacodynamic modeling of lansoprazole using an indirect irreversible1254response model. J Clin Pharmacol 2001; 41(3): 251–258.
- 1255 83. Nielsen El *et al.* Pharmacokinetic/pharmacodynamic (PK/PD) indices of antibiotics predicted
 1256 by a semimechanistic PKPD model: a step toward model-based dose optimization. *Antimicrob* 1257 *Agents Chemother* 2011; 55(10): 4619–4630.
- 125884.Snoeck E *et al.* A comprehensive hepatitis C viral kinetic model explaining cure. *Clin Pharmacol*1259*Ther* 2010; 87(6): 706–713.
- 1260 85. Simeoni M *et al.* Predictive pharmacokinetic-pharmacodynamic modeling of tumor growth
 1261 kinetics in xenograft models after administration of anticancer agents. *Cancer Res* 2004; 64(3):
 1262 1094–1101.
- 126386.Friberg LE *et al.* Semiphysiological model for the time course of leukocytes after varying1264schedules of 5-fluorouracil in rats. J Pharmacol Exp Ther 2000; 295(2): 734–740.
- 126587.Russu A, Poggesi I. Turnover model with irreversible inactivation. In: Mager DE, Kimko HHC,1266eds. Systems Pharmacology and Pharmacodynamics. Springer Nature, 2016: 217.
- 1267 88. Nagaya H *et al.* Possible mechanism for the inhibition of gastric (H+ + K+)-adenosine
 1268 triphosphatase by the proton pump inhibitor AG-1749. *J Pharmacol Exp Ther* 1989; 248(2):
 1269 799–805.
- 127089.Shin JM *et al.* The site of action of pantoprazole in the gastric H+/K(+)-ATPase. *Biochim*1271*Biophys Acta* 1993; 1148(2): 223–233.
- 127290.Fitton A, Wiseman L. Pantoprazole. A review of its pharmacological properties and1273therapeutic use in acid-related disorders. *Drugs* 1996; 51(3): 460–482.
- 1274 91. Im WB *et al.* Irreversible inactivation of rat gastric (H+-K+)-ATPase in vivo by omeprazole.
 1275 *Biochem Biophys Res Commun* 1985; 126(1): 78–82.
- 1276 92. Sachs G *et al.* Gastric acid secretion: activation and inhibition. *Yale J Biol Med* 1994; 67(3–4):
 1277 81–95.

1278 93. 1279	Gedda K <i>et al</i> . Turnover of the gastric H+,K(+)-adenosine triphosphatase alpha subunit and its effect on inhibition of rat gastric acid secretion. <i>Gastroenterology</i> 1995; 109(4): 1134–41.
1280 94. 1281	Metz DC <i>et al.</i> Proton pump activation in stimulated parietal cells is regulated by gastric acid secretory capacity: A human study. <i>J Clin Pharmacol</i> 2002; 42(5): 512–519.
1282 95. 1283	Katashima M <i>et al.</i> Comparative pharmacokinetic/pharmacodynamic study of proton pump inhibitors, omeprazole and lansoprazole in rats. <i>Drug Metab Dispos</i> 1995; 23(7): 718 LP-723.
1284 96. 1285	Shin JM, Sachs G. Differences in binding properties of two proton pump inhibitors on the gastric H +,K +-ATPase in vivo. <i>Biochem Pharmacol</i> 2004; 68(11): 2117–2127.
1286 97. 1287 1288	Sugiura M <i>et al.</i> Prediction of Therapeutic Doses Based on the Pharmacokinetic/Pharmacodynamic Model of Omeprazole, a Proton Pump Inhibitor. <i>Drug Metab Pharmacokinet</i> 1992; 7(6): 813–820.
1289 98. 1290	Dayneka NL <i>et al.</i> Comparison of four basic models of indirect pharmacodynamic responses. <i>J</i> <i>Pharmacokinet Biopharm</i> 1993; 21(4): 457–478.
1291 99. 1292	Polentarutti B <i>et al.</i> Modification of gastric pH in the fasted dog. <i>J Pharm Pharmacol</i> 2010; 62(4): 462–9.
1293 100. 1294	Ferron GM <i>et al.</i> Pharmacodynamic Modeling of Pantoprazole's Irreversible Effect on Gastric Acid Secretion in Humans and Rats. <i>J Clin Pharmacol</i> 2001; 41: 149–156.
1295 101. 1296 1297	Feldman M <i>et al.</i> A comparison of every-third-day versus daily low-dose aspirin therapy on serum thromboxane concentrations in healthy men and women. <i>Clin Appl Thromb Hemost</i> 2001; 7(1): 53–7.
1298 102. 1299 1300	Nagelschmitz J <i>et al.</i> Pharmacokinetics and pharmacodynamics of acetylsalicylic acid after intravenous and oral administration to healthy volunteers. <i>Clin Pharmacol Adv Appl</i> 2014; 6(1): 51–59.
1301 103. 1302	Patrignani P <i>et al.</i> Selective cumulative inhibition of platelet thromboxane production by low- dose aspirin in healthy subjects. <i>J Clin Invest</i> 1982; 69(6): 1366–1372.
1303 104. 1304	Renda G <i>et al.</i> Celecoxib, ibuprofen, and the antiplatelet effect of aspirin in patients with osteoarthritis and ischemic heart disease. <i>Clin Pharmacol Ther</i> 2006; 80(3): 264–274.
1305 105. 1306	Giaretta A <i>et al.</i> In Silico Modeling of the Antiplatelet Pharmacodynamics of Low-dose Aspirin in Health and Disease. <i>Clin Pharmacol Ther</i> 2017; 102(5): 823–831.
1307 106. 1308	Paridaens R <i>et al.</i> Safety, activity and estrogen inhibition by exemestane in postmenopausal women with advanced breast cancer: a phase I study. <i>Anticancer Drugs</i> 1998; 9(8): 675–83.
1309 107. 1310 1311	Johannessen DC <i>et al.</i> Endocrine and clinical effects of exemestane (PNU 155971), a novel steroidal aromatase inhibitor, in postmenopausal breast cancer patients: a phase I study. <i>Clin Cancer Res</i> 1997; 3(7): 1101–8.
1312 108. 1313 1314	Geisler J <i>et al.</i> In vivo inhibition of aromatization by exemestane, a novel irreversible aromatase inhibitor, in postmenopausal breast cancer patients. <i>Clin Cancer Res</i> 1998; 4(9): 2089–93.
1315 109. 1316	Nagashima R <i>et al</i> . Kinetics of pharmacologic effects in man: The anticoagulant action of warfarin. <i>Clin Pharmacol Ther</i> 1969; 10(1): 22–35.
1317 110.	Dayneka NL <i>et al.</i> Comparison of Four Basic Models of Indirect Pharmacodynamic Responses.

1318 J Pharmacokinet Biopharm 1993; 21(22). 1319 111. Chakraborty A et al. Mathematical modeling of circadian cortisol concentrations using indirect 1320 response models: comparison of several methods. J Pharmacokinet Biopharm 1999; 27(1): 23-43. 1321 1322 112. Krzyzanski W et al. Basic Pharmacodynamic Models for Agents That Alter Production of Natural Cells. J Pharmacokinet Pharmacodyn 1999; 27(5): 467-489. 1323 1324 Budha NR et al. Comparative Performance of Cell Life Span and Cell Transit Models for 113. 1325 Describing Erythropoietic Drug Effects. AAPS J 2011; 13(4): 650-661. 1326 114. Samtani MN et al. Pharmacokinetic and Pharmacodynamic Modeling of Pegylated 1327 Thrombopoietin Mimetic Peptide (PEG-TPOm) After Single Intravenous Dose Administration 1328 in Healthy Subjects. J Clin Pharmacol 2009; 49(3): 336–350. 1329 Yao Z et al. Assessment of Basic Indirect Pharmacodynamic Response Models with 115. 1330 Physiological Limits. J Pharmacokinet Pharmacodyn 2006; 33(2): 167–193. 1331 116. Labrecque G, Bélanger PM. Biological rhythms in the absorption, distribution, metabolism and 1332 excretion of drugs. Pharmacol Ther 1991; 52(1): 95-107. 1333 117. Sällström B et al. A Pharmacodynamic Turnover Model Capturing Asymmetric Circadian 1334 Baselines of Body Temperature, Heart Rate and Blood Pressure in Rats: Challenges in Terms of 1335 Tolerance and Animal-handling Effects. J Pharmacokinet Pharmacodyn 2005; 32(5-6): 835-1336 859. 1337 118. Sukumaran S et al. Circadian rhythms in gene expression: Relationship to physiology, disease, 1338 drug disposition and drug action. Adv Drug Deliv Rev 2010; 62(9–10): 904–917. 1339 119. Lew KH et al. Gender-based effects on methylprednisolone pharmacokinetics and 1340 pharmacodynamics. Clin Pharmacol Ther 1993; 54(4): 402–14. 1341 120. Rohatagi S et al. Dynamic modeling of cortisol reduction after inhaled administration of 1342 fluticasone propionate. J Clin Pharmacol 1996; 36(10): 938-41. 1343 Krzyzanski W. Direct, Indirect, and Signal Transduction Response Modeling. In: Mager DE, 121. 1344 Kimko HHC, eds. Systems Pharmacology and Pharmacodynamics. Springer, 2016: 177–210. 1345 122. Mager DE, Jusko WJ. Pharmacodynamic modeling of time-dependent transduction systems. 1346 *Clin Pharmacol Ther* 2001; 70(3): 210–216. 1347 123. Zhang Y, D'Argenio DZ. Feedback Control Indirect Response Models. In: Mager DE, Kimko 1348 HHC, eds. Systems Pharmacology and Pharmacodynamics., 2016: 229–254. 1349 124. Black JW, Leff P. Operational models of pharmacological agonism. Proc R Soc London Ser B, 1350 *Biol Sci* 1983; 220(1219): 141–62. 1351 125. Van Der Graaf PH et al. Mechanism-based pharmacokinetic-pharmacodynamic modeling of 1352 the effects of N6-cyclopentyladenosine analogs on heart rate in rat: estimation of in vivo 1353 operational affinity and efficacy at adenosine A1 receptors. J Pharmacol Exp Ther 1997; 1354 283(2): 809-16. 1355 126. Greene SJ et al. Partial adenosine A1 receptor agonism: a potential new therapeutic strategy 1356 for heart failure. Heart Fail Rev 2016; 21(1): 95–102. 1357 127. Cox EH et al. Pharmacokinetic-pharmacodynamic modelling of the EEG effect of alfentanil in

- 1358 rats. J Pharmacol Toxicol Methods 1997; 38(2): 99–108.
- 1359 128. Cox EH *et al.* Pharmacokinetic-pharmacodynamic modeling of the electroencephalogram
 1360 effect of synthetic opioids in the rat: correlation with the interaction at the mu-opioid
 1361 receptor. *J Pharmacol Exp Ther* 1998; 284(3): 1095–103.
- 129. Zuideveld KP *et al.* Pharmacokinetic-pharmacodynamic modelling of the hypothermic and
 corticosterone effects of the 5-HT1A receptor agonist flesinoxan. *Eur J Pharmacol* 2002;
 445(1–2): 43–54.
- 130. Ramakrishnan R *et al.* Fifth-Generation Model for Corticosteroid Pharmacodynamics:
 Application to Steady-State Receptor Down-Regulation and Enzyme Induction Patterns during
 Seven-Day Continuous Infusion of Methylprednisolone in Rats. *J Pharmacokinet Pharmacodyn* 2002; 29(1): 1–24.
- 1369 131. Sandström M *et al.* Model Describing the Relationship Between Pharmacokinetics and
 1370 Hematologic Toxicity of the Epirubicin-Docetaxel Regimen in Breast Cancer Patients. *J Clin* 1371 Oncol 2005; 23(3): 413–421.
- 1372132.Friberg LE *et al.* Model of Chemotherapy-Induced Myelosuppression With Parameter1373Consistency Across Drugs. *J Clin Oncol* 2002; 20(24): 4713–4721.
- 1374133.Friberg LE *et al.* Semiphysiological Model for the Time Course of Leukocytes after Varying1375Schedules of 5-Fluorouracil in Rats. J Pharmacol Exp Ther 2000; 295(2): 734–40.
- 1376134.Gabrielsson J, Peletier LA. A Flexible Nonlinear Feedback System That Captures Diverse1377Patterns of Adaptation and Rebound. AAPS J 2008; 10(1): 70–83.
- 1378 135. Wakelkamp M *et al.* Pharmacodynamic modeling of furosemide tolerance after multiple
 1379 intravenous administration. *Clin Pharmacol Ther* 1996; 60(1): 75–88.
- 1380136.Ahlström C *et al.* Feedback modeling of non-esterified fatty acids in obese Zucker rats after1381nicotinic acid infusions. J Pharmacokinet Pharmacodyn 2013; 40(6): 623–638.
- 1382137.Walson PD, Galletta G, Braden NJ AL. Ibuprofen, acetaminophen, and placebo treatment of1383febrile children. Clin Pharmacol Ther 1989; 46(1): 9–17.
- 1384138.Wilson JT *et al.* Single-dose, placebo-controlled comparative study of ibuprofen and1385acetaminophen antipyresis in children. J Pediatr 1991; 119(5): 803–11.
- 1386139.Brown RD *et al.* Single-dose pharmacokinetics of ibuprofen and acetaminophen in febrile1387children. J Clin Pharmacol 1992; 32(3): 231–41.
- 1388 140. Brown RD *et al.* Integrated Pharmacokinetic-Pharmacodynamic Model for Acetaminophen,
 1389 Ibuprofen, and Placebo Antipyresis in Children. *J Pharmacokinet Biopharm* 1998; 26(5).
- 1390 141. Mackowiak PA. Concepts of Fever. Arch Intern Med 1998; 158(17): 1870–1881.
- 1391 142. Garg V, Jusko WJ. Pharmacodynamic modeling of nonsteroidal anti-inflammatory drugs:
 1392 antipyretic effect of ibuprofen. *Clin Pharmacol Ther* 1994; 55(1): 87–88.
- 1393143.Kauffman RE, Nelson M V. Effect of age on ibuprofen pharmacokinetics and antipyretic1394response. J Pediatr 1992; 121(6): 969–73.
- 1395144.Olsson AG *et al.* Effect of rosuvastatin on low-density lipoprotein cholesterol in patients with1396hypercholesterolemia. Am J Cardiol 2001; 88(5): 504–8.

1397 145. Davidson MH. Rosuvastatin: a highly efficacious statin for the treatment of dyslipidaemia. 1398 *Expert Opin Investig Drugs* 2002; 11(1): 125–141. 1399 146. Schachter M. Chemical, pharmacokinetic and pharmacodynamic properties of statins: An 1400 update. Fundam Clin Pharmacol 2005; 19(1): 117-125. 1401 147. Aoyama T et al. Pharmacokinetic/pharmacodynamic modeling and simulation of rosuvastatin 1402 using an extension of the indirect response model by incorporating a circadian rhythm. Biol 1403 *Pharm Bull* 2010; 33(6): 1082–7. 1404 148. Krzyzanski W et al. Algorithm for application of Fourier analysis for biorhythmic baselines of 1405 pharmacodynamic indirect response models. Chronobiol Int 2000; 17(1): 77–93. 1406 149. Krzyzanski W, Jusko WJ. Indirect Pharmacodynamic Models for Responses with 1407 Multicompartmental Distribution or Polyexponential Disposition. J Pharmacokinet 1408 Pharmacodyn 2001; 28(1). 1409 150. Martin PD et al. Pharmacodynamic effects and pharmacokinetics of a new HMG-CoA 1410 reductase inhibitor, rosuvastatin, after morning or evening administration in healthy 1411 volunteers. Br J Clin Pharmacol 2002; 54(5): 472-7. 1412 151. HO R, KIM R. Transporters and drug therapy: Implications for drug disposition and disease. 1413 *Clin Pharmacol Ther* 2005; 78(3): 260–277. 1414 152. Ho RH et al. Drug and Bile Acid Transporters in Rosuvastatin Hepatic Uptake: Function, 1415 Expression, and Pharmacogenetics. Gastroenterology 2006; 130(6): 1793–1806. 1416 153. Hirano M et al. Contribution of OATP2 (OATP1B1) and OATP8 (OATP1B3) to the Hepatic 1417 Uptake of Pitavastatin in Humans. Kameyama Y et al. Functional characterization of SLCO1B1 (OATP-C) variants, SLCO1B1*5, 1418 154. 1419 SLCO1B1*15 and SLCO1B1*15+C1007G, by using transient expression systems of HeLa and 1420 HEK293 cells. Pharmacogenet Genomics 2005; 15(7): 513-22. 1421 Hsiang B et al. A novel human hepatic organic anion transporting polypeptide (OATP2). 155. 1422 Identification of a liver-specific human organic anion transporting polypeptide and 1423 identification of rat and human hydroxymethylglutaryl-CoA reductase inhibitor transporters. J 1424 Biol Chem 1999; 274(52): 37161-8. 1425 156. Kitamura S et al. Involvement of Multiple Transporters in the Hepatobiliary Transport of 1426 Rosuvastatin. Drug Metab Dispos 2008; 36(10): 2014–2023. 1427 157. Nishizato Y et al. Polymorphisms of OATP-C (SLC21A6) and OAT3 (SLC22A8) genes: 1428 Consequences for pravastatin pharmacokinetics. Clin Pharmacol Ther 2003; 73(6): 554–565. 1429 Pasanen MK et al. Different Effects of SLCO1B1 Polymorphism on the Pharmacokinetics of 158. 1430 Atorvastatin and Rosuvastatin. Clin Pharmacol Ther 2007; 82(6): 726–733. 1431 159. Pasanen MK et al. SLCO1B1 polymorphism markedly affects the pharmacokinetics of simvastatin acid. Pharmacogenet Genomics 2006; 16(12): 873-879. 1432 1433 160. Niemi M et al. SLCO1B1 polymorphism and sex affect the pharmacokinetics of pravastatin but 1434 not fluvastatin. Clin Pharmacol Ther 2006; 80(4): 356–366. 1435 Niemi M et al. High plasma pravastatin concentrations are associated with single nucleotide 161. 1436 polymorphisms and haplotypes of organic anion transporting polypeptide-C (OATP-C,

- 1437 SLCO1B1). *Pharmacogenetics* 2004; 14(7): 429–40.
- 1438 162. Rose RH *et al.* Application of a Physiologically Based Pharmacokinetic Model to Predict
 1439 OATP1B1-Related Variability in Pharmacodynamics of Rosuvastatin. *CPT pharmacometrics*1440 Syst Pharmacol 2014; 3(April): e124.
- 1441 163. Tachibana-Iimori R *et al.* Effect of genetic polymorphism of OATP-C (SLCO1B1) on lipid1442 lowering response to HMG-CoA reductase inhibitors. *Drug Metab Pharmacokinet* 2004; 19(5):
 1443 375–80.
- 1444 164. Pasanen MK *et al.* Polymorphism of the hepatic influx transporter organic anion transporting
 1445 polypeptide 1B1 is associated with increased cholesterol synthesis rate. *Pharmacogenet*1446 *Genomics* 2008; 18(10): 921–926.
- 1447 165. Niemi M *et al.* Organic Anion Transporting Polypeptide 1B1: a Genetically Polymorphic
 1448 Transporter of Major Importance for Hepatic Drug Uptake. *Pharmacol Rev* 2011; 63(1): 157–
 1449 181.
- 1450166.Niemi M. Transporter Pharmacogenetics and Statin Toxicity. Clin Pharmacol Ther 2010; 87(1):1451130–133.
- 1452 167. Piñeyro G, Blier P. Autoregulation of serotonin neurons: role in antidepressant drug action.
 1453 *Pharmacol Rev* 1999; 51(3): 533–91.
- 1454168.Bourne JA. Intracerebral microdialysis: 30 years as a tool for the neuroscientist. Clin Exp1455Pharmacol Physiol 30(1-2): 16-24.
- 1456169.Westerink BH., Timmerman W. Do neurotransmitters sampled by brain microdialysis reflect1457functional release? Anal Chim Acta 1999; 379(3): 263–274.
- 1458 170. Bundgaard C *et al.* Mechanistic model of acute autoinhibitory feedback action after
 1459 administration of SSRIs in rats: Application to escitalopram-induced effects on brain serotonin
 1460 levels. 2006.
- 1461 171. Ceglia I *et al.* Effects of chronic treatment with escitalopram or citalopram on extracellular 51462 HT in the prefrontal cortex of rats: role of 5-HT1A receptors. *Br J Pharmacol* 2004; 142(3):
 1463 469–78.
- 1464 172. Mørk A *et al.* The R-enantiomer of citalopram counteracts escitalopram-induced increase in
 1465 extracellular 5-HT in the frontal cortex of freely moving rats. *Neuropharmacology* 2003; 45(2):
 1466 167–73.
- 1467173.Selen A *et al.* The biopharmaceutics risk assessment roadmap for optimizing clinical drug1468product performance. J Pharm Sci 2014; 103(11): 3377–3397.
- 1469 174. Dickinson PA *et al.* Clinical Relevance of Dissolution Testing in Quality by Design. *AAPS J* 2008;
 1470 10(2): 380–390.