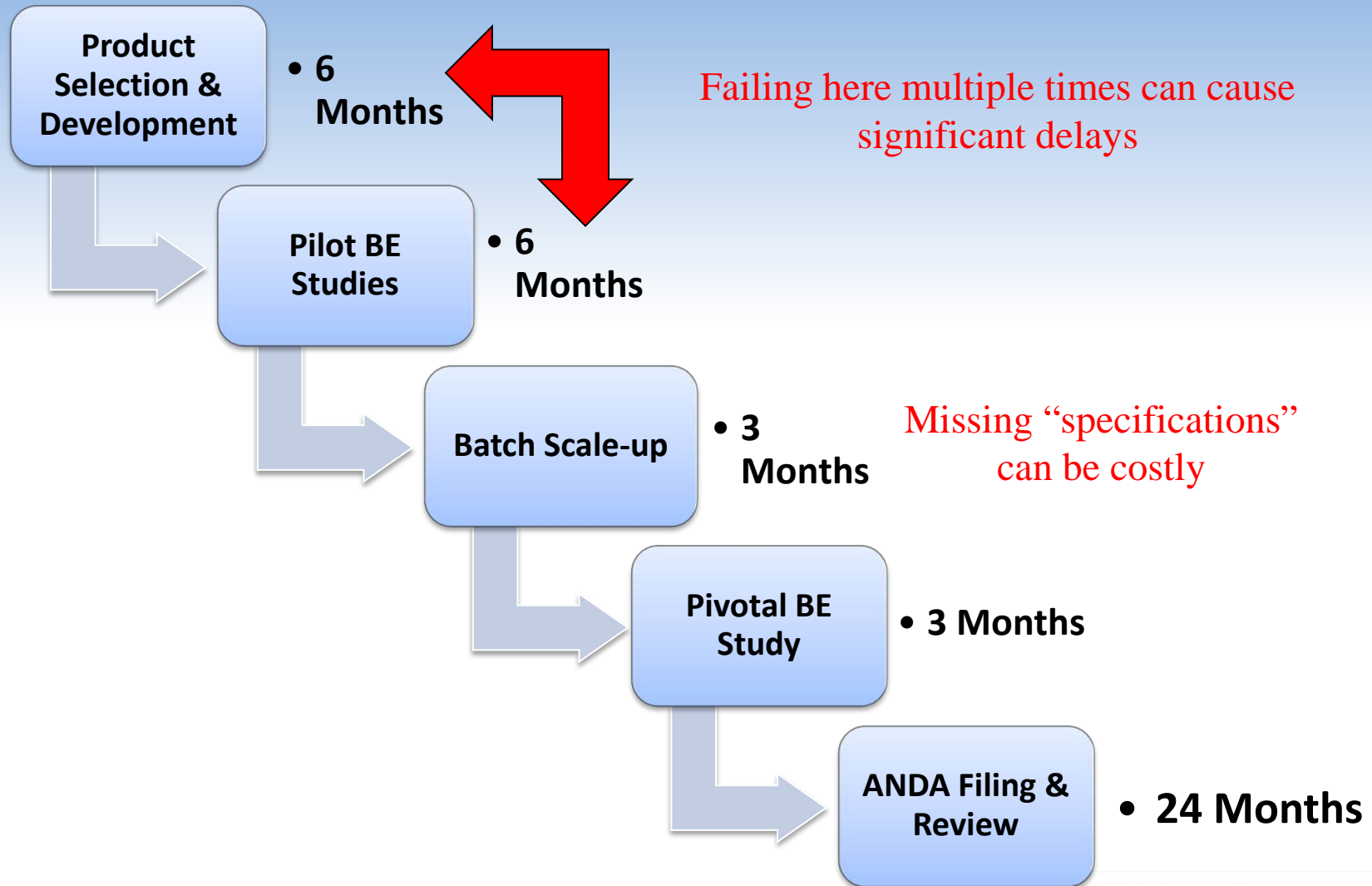


*Incorporating Mechanistic Modeling
& Simulation to Assist with
Formulation Development and
Regulatory Evaluations*

*Viera Lukacova
Simulations Plus, Inc.*

The Generic Product Development Process



Outline

- Why Modeling & Simulation?
- Overview of Mechanistic Simulation Models
 - Predicting *in vivo* absorption & PK
- Applications in Generic Product Development
 - Generating IVIVCs
 - Performing virtual bioequivalence trials and establishing dissolution specifications
 - Understanding food effects
- A successful biowaiver case study
- Conclusions

How can simulation software be used?

- Dissolution Method Development
 - Which *in vitro* method best correlates with an *in vivo* profile?
- Formulation Design
 - How do I design my formulation to achieve bioequivalence?
- Establish Dissolution Specifications
 - What is the acceptable variability in key parameters before we are no longer bioequivalent?

Flow Diagram for Simulation Studies

The AAPS Journal, Vol. 13, No. 1, March 2011 (© 2010)
DOI: 10.1208/s12248-010-9250-9

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Research Article

Utility of Physiologically Based Absorption Modeling in Implementing Quality by Design in Drug Development

Xinyuan Zhang,¹ Robert A. Lionberger,^{1,2} Barbara M. Davit,¹ and Lawrence X. Yu¹

Received 16 September 2010; accepted 14 December 2010; published online 5 January 2011

Abstract. To implement Quality by Design (QbD) in drug development, scientists need tools that link drug products properties to *in vivo* performance. Physiologically based absorption models are potentially useful tools; yet, their utility of QbD implementation has not been discussed or explored much in the literature. We simulated pharmacokinetics (PK) of carbamazepine (CBZ) after administration of four oral formulations, immediate-release (IR) suspension, IR tablet, extended-release (XR) tablet and capsule, under fasted and fed conditions and presented a general diagram of a modeling and simulation strategy integrated with pharmaceutical development. We obtained PK parameters and absorption scale factors (ASFs) by deconvolution of the PK data for IR suspension under fasted condition. The model was validated for other PK profiles of IR formulations and used to predict PK for XR formulations. We explored three key areas where a modeling and simulation approach impacts QbD. First, the model was used to help identify optimal *in vitro* dissolution conditions for XR formulations. Second, identification of critical formulations variables was illustrated by a parameter sensitivity analysis of mean particle radius for the IR tablet that showed a PK shift with decreased particle radius, C_{max} was increased and T_{max} was decreased. Finally, virtual trial simulations allowed incorporation of inter-subject variability in the model. Virtual bioequivalence studies performed for two test formulations suggested that an *in vitro* dissolution test may be a more sensitive discriminative method than *in vivo* PK studies. In summary, a well-validated predictive model is a potentially useful tool for QbD implementation in drug development.

KEY WORDS: advanced compartmental absorption and transit (ACAT) model; gastroplus™; modified release (MR); quality by design (QbD).

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Fig. 1.

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Flow Diagram for Simulation Studies

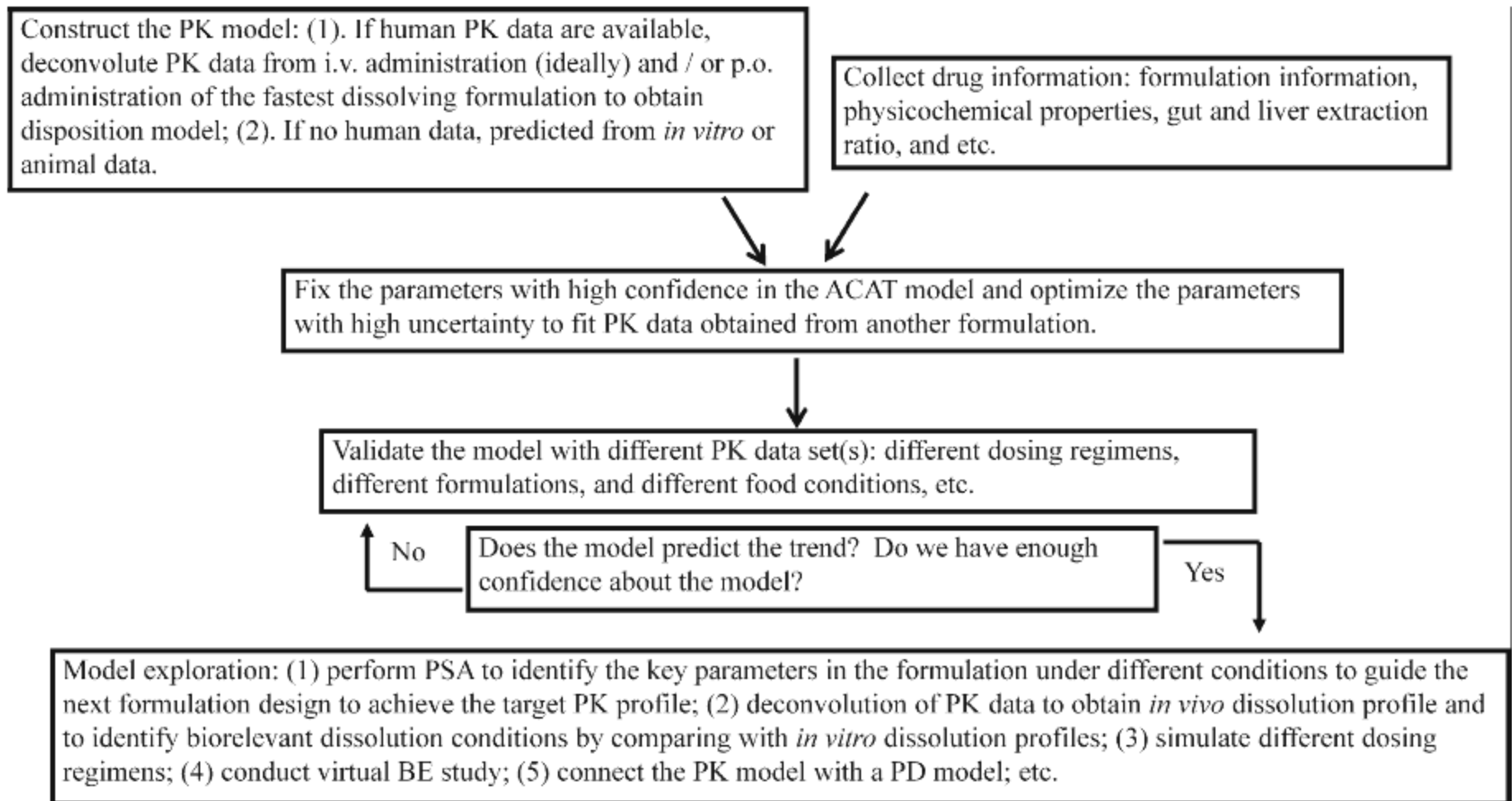
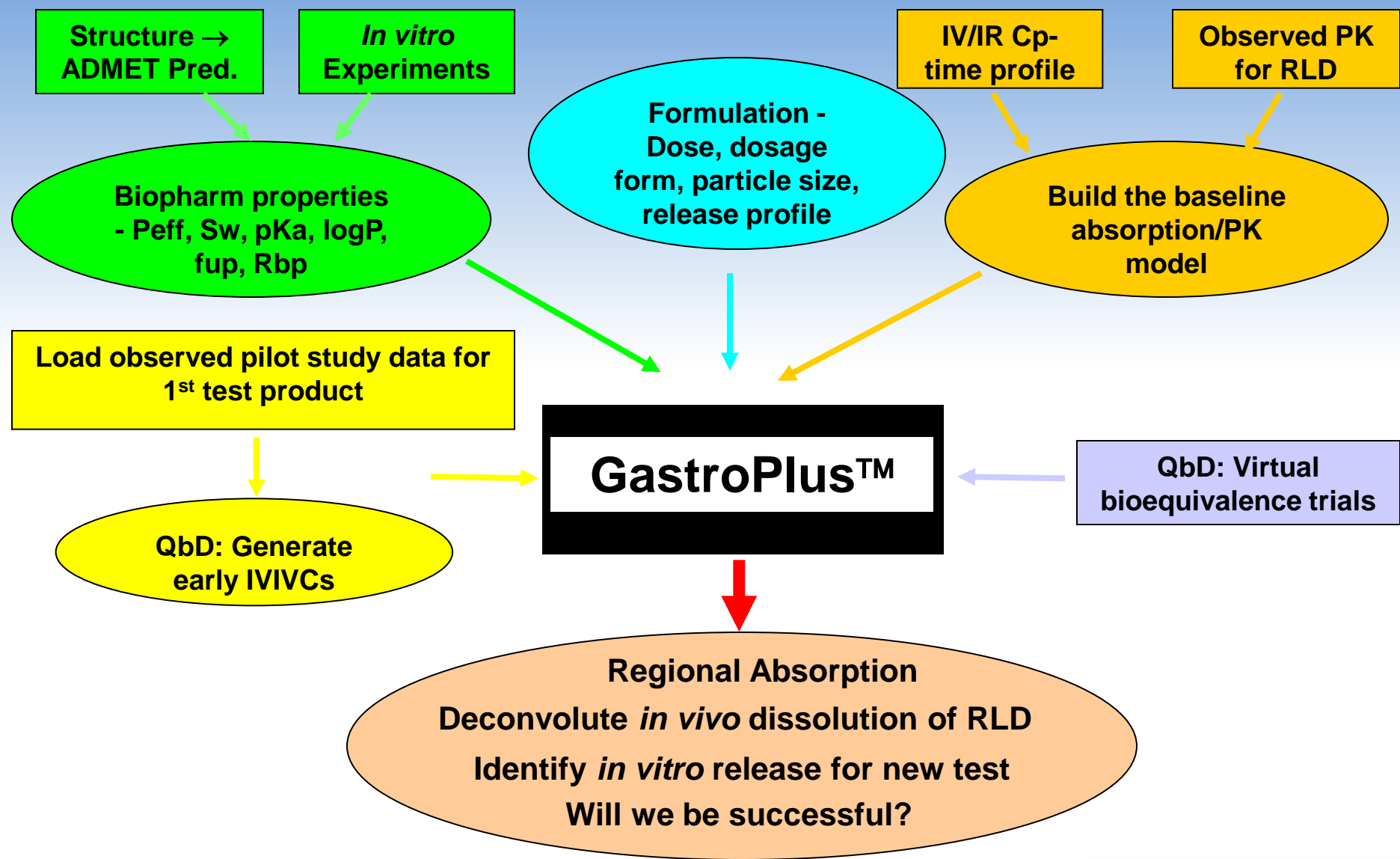


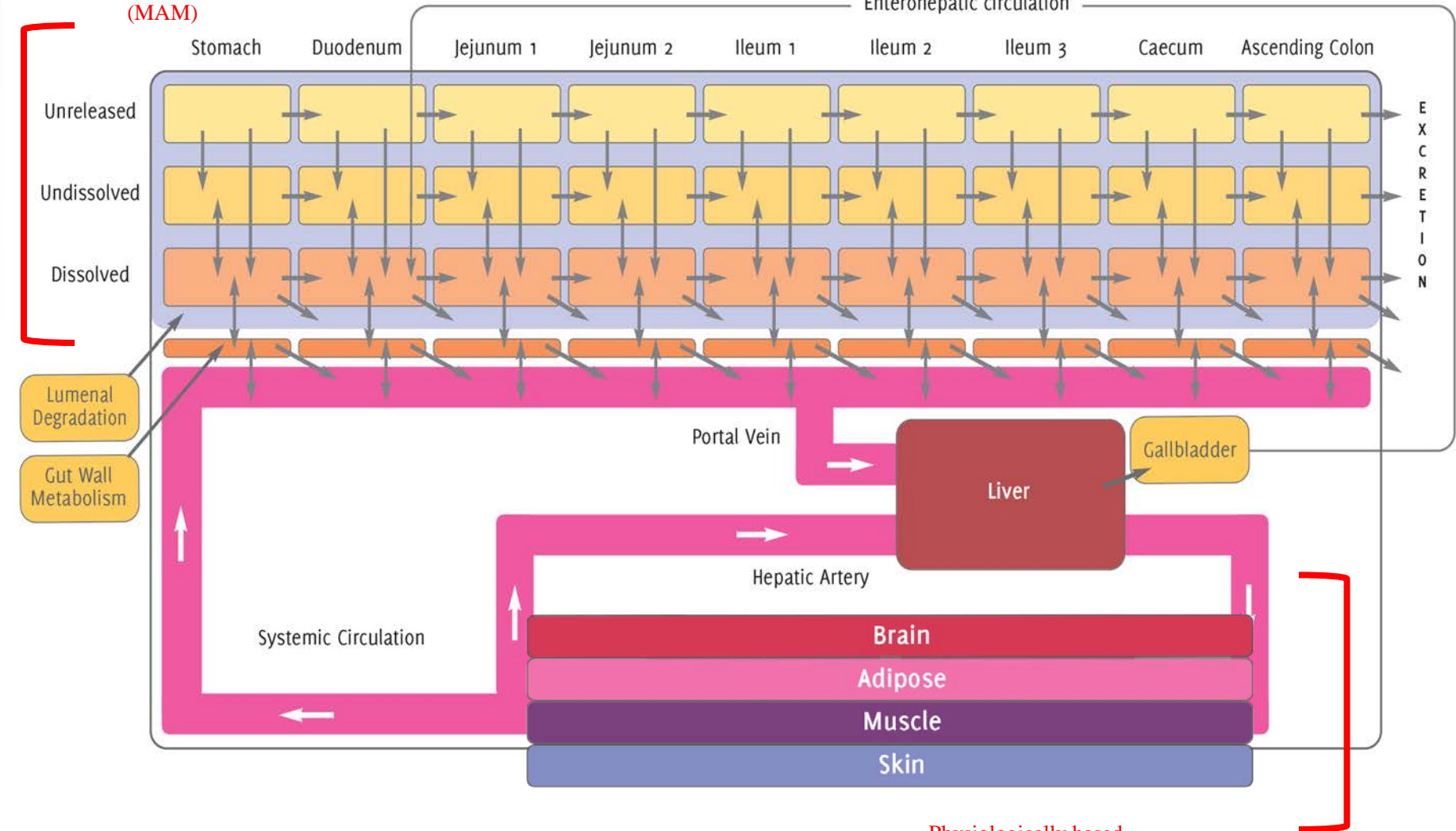
Fig. 1. The flow diagram shows a general process of using a physiologically based absorption model in QbD-based drug development

The Big Picture



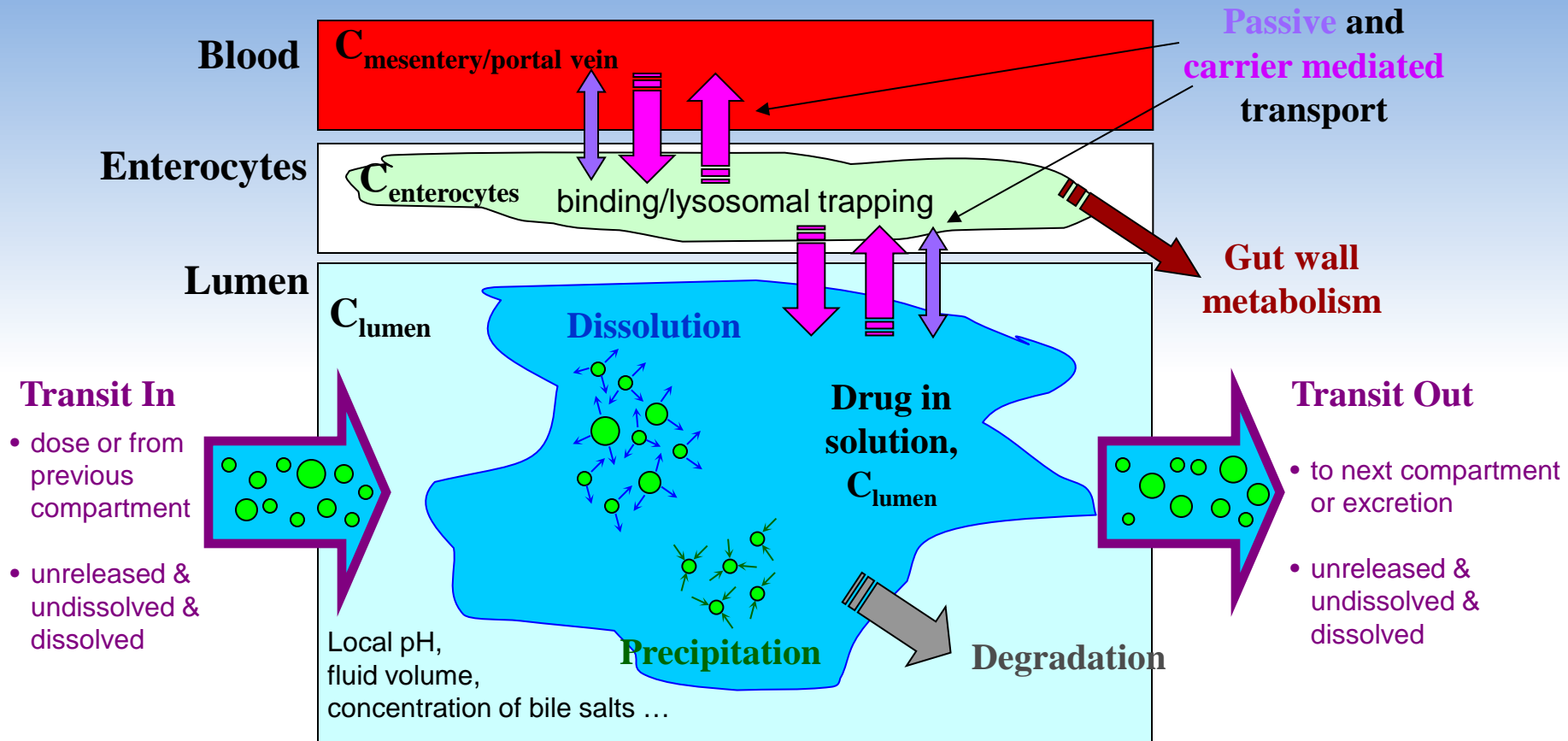
Advanced Compartmental Absorption and Transit Model (ACAT™)

Mechanistic Absorption Modeling (MAM)



Physiologically based Pharmacokinetics (PBPK)

Processes Involved in Oral Absorption

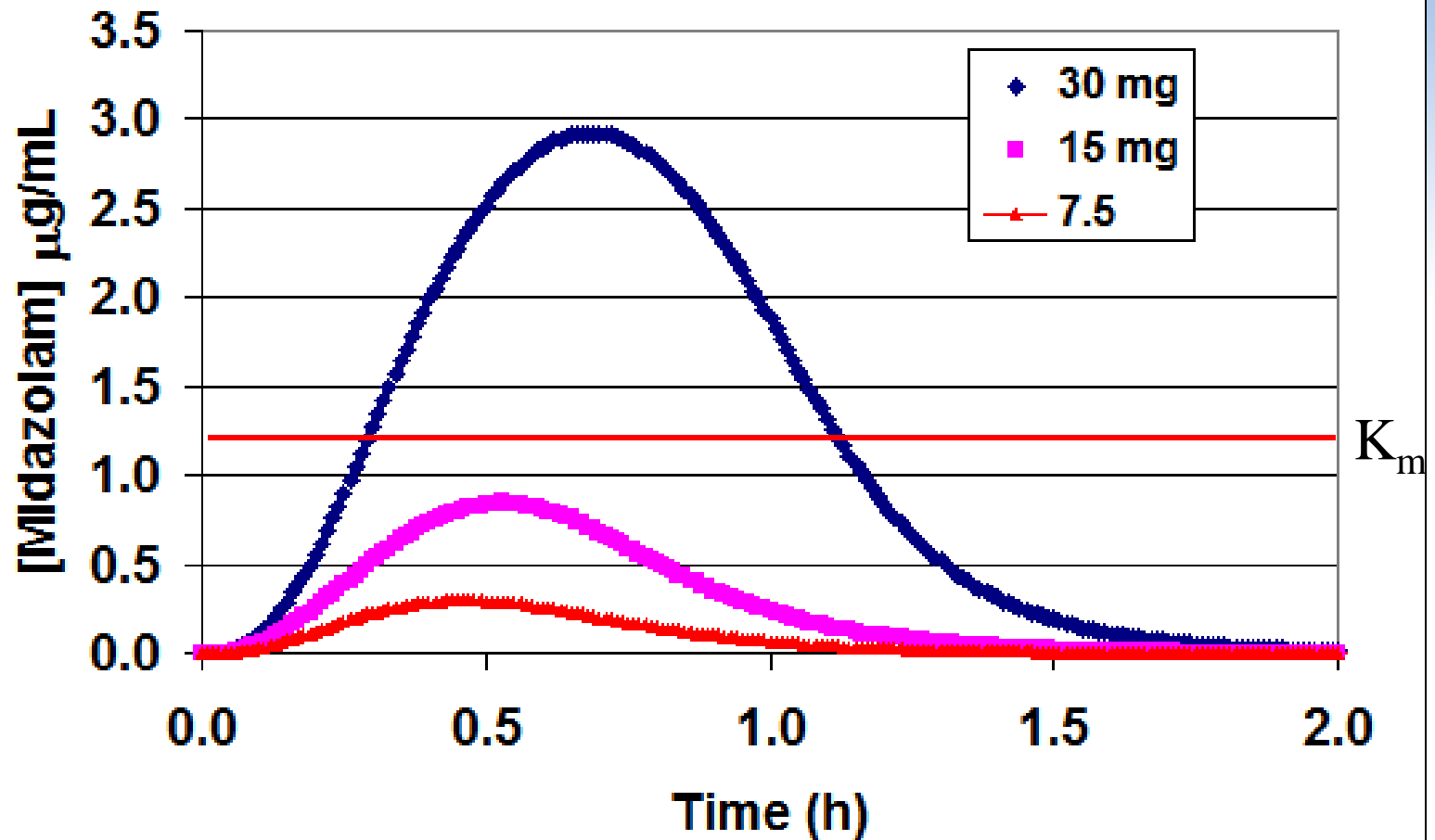


These phenomena:

- are happening simultaneously
- are repeated in each of the compartments of the gastrointestinal tract

[Jejunal Enterocyte]

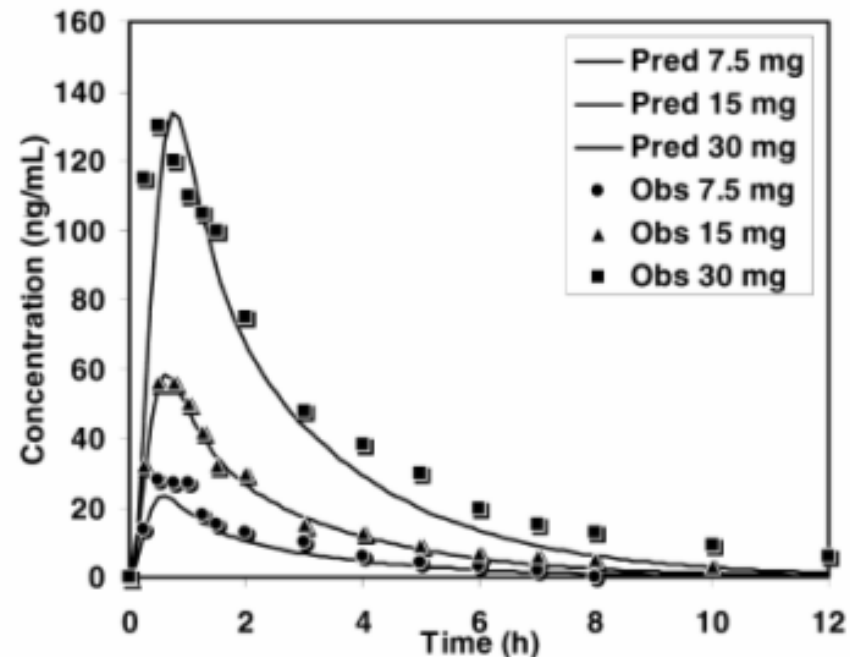
Note: Midazolam $K_m=1.2 \mu\text{g/mL}$



Nonlinear Dose Dependence of Midazolam Metabolism in Gut and Liver

Dose	Experimental		GastroPlus Compartmental Simulated				
	C _{max}	AUC	C _{max}	AUC	Fa%	FDP%	Fb%
7.5	0.028	69	0.021	65	99	45	24
15	0.056	154	0.052	158	99	55	29
30	0.13	453	0.120	369	99	64	34

GastroPlus simulations of nonlinear dose dependence for midazolam using *in vitro* K_m and V_{max} and *iv* PK. (Agoram et al., 2001)



Observed nonlinear dose dependence for valacyclovir

600 Weller et al.

CLINICAL PHARMACOLOGY & THERAPEUTICS
DECEMBER 1993

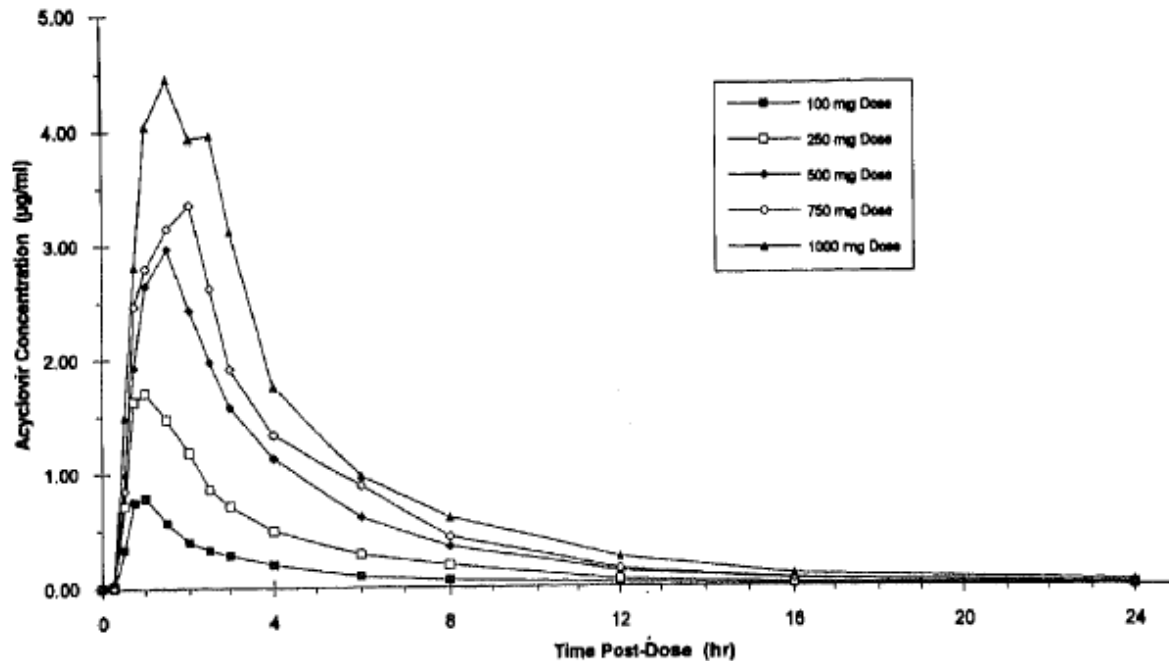
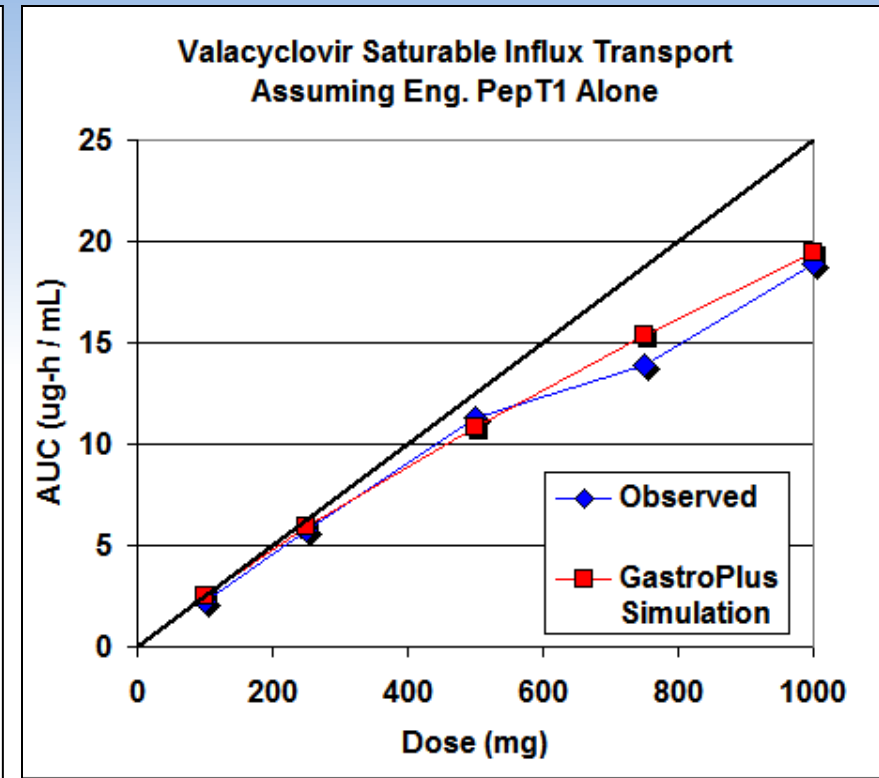
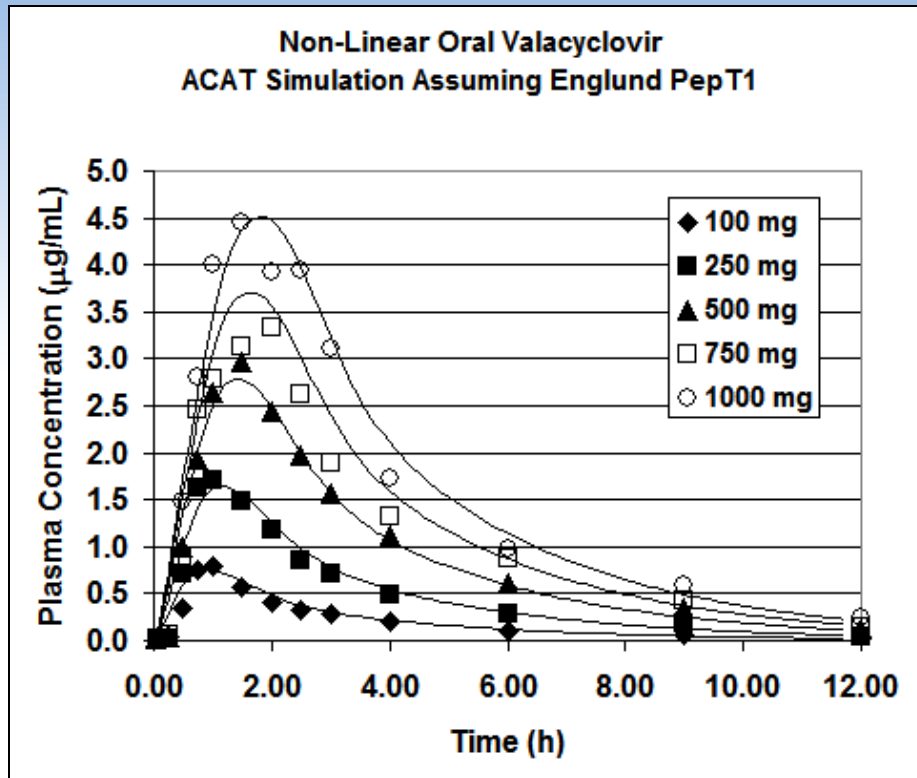


Fig. 1. Mean plasma acyclovir concentrations after single-dose administration of valacyclovir to normal volunteers. A single cohort of subjects received valacyclovir at each dose level ($n = 7$ except for the 100 mg dose, for which $n = 8$).

Weller, S. Clin. Pharm. Ther. 54(6):595 (1993)

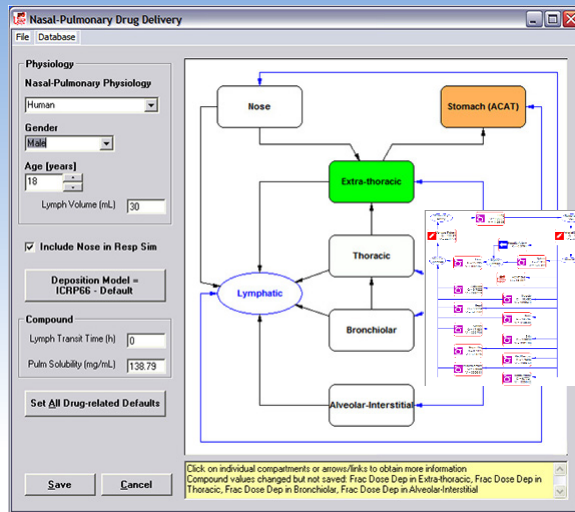
GastroPlus simulation of nonlinear dose dependence for influx transport of valacyclovir



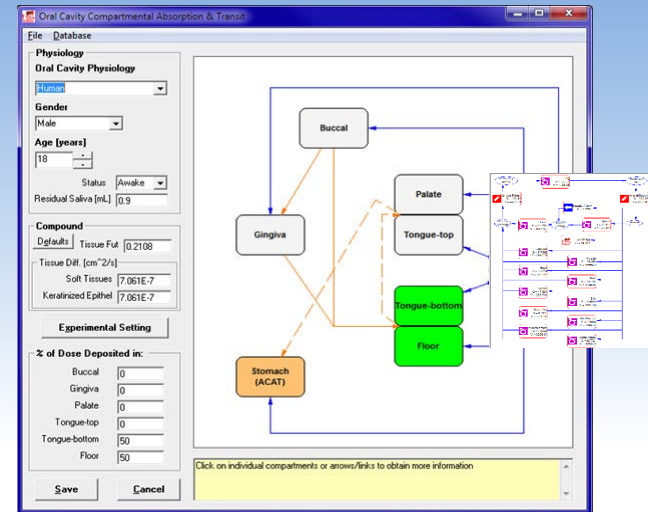
Bolger MB, et al. AAPS Journal 11(2):353 (2009)
GastroPlus results were first reported in Feb. 2003
at AAPS Drug Transport Workshop, Peachtree City, GA

QbD: Beyond mechanistic oral absorption

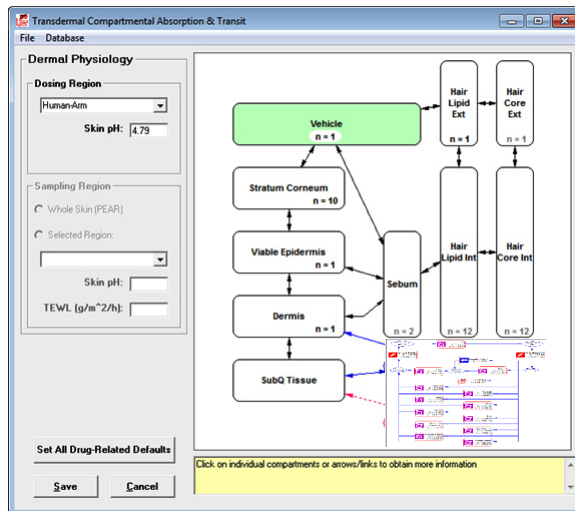
Pulmonary (PCAT™)



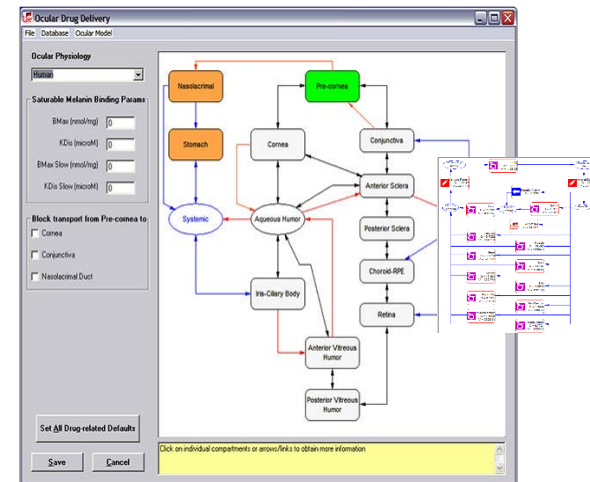
Oral Cavity (OCCAT™)



Dermal (TCAT™)



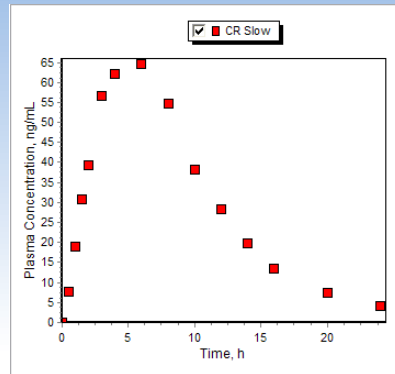
Ocular (OCAT™)



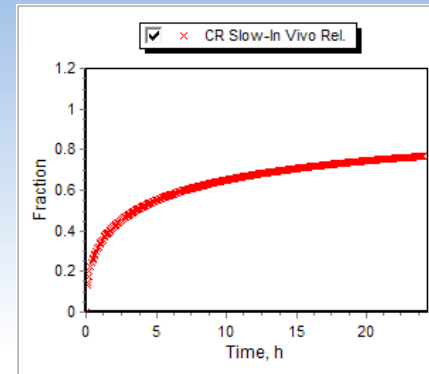
*Developing a mechanistic
in vitro-in vivo correlation
(IVIVC)*

Deconvolution

(with GastroPlus™ Mechanistic Absorption method)



Deconvolution



in vivo dissolution
vs. time along the
gut– **NOT F%!**

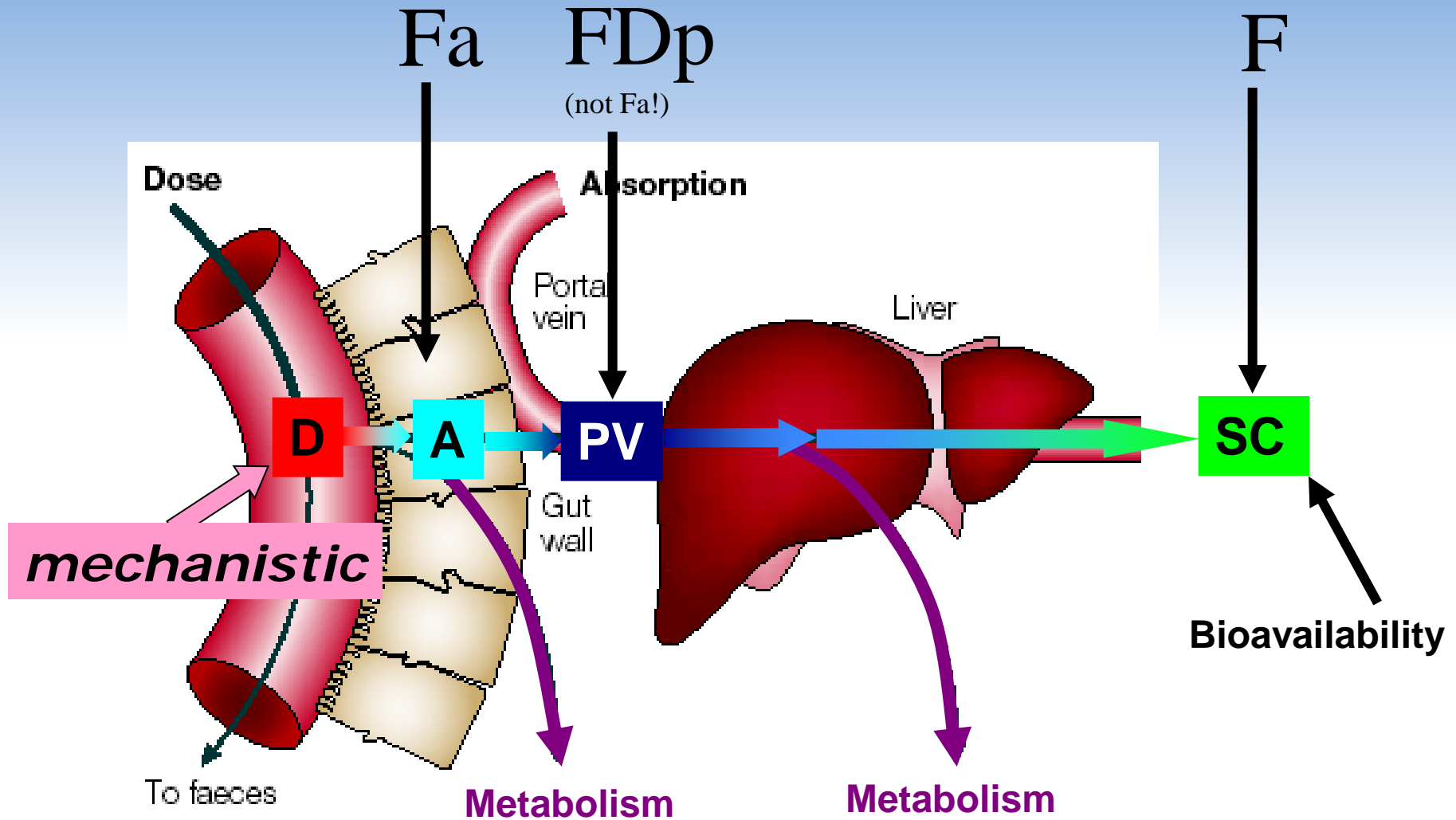
- Inputs (in addition to the data required for the traditional methods):
 - Physiological parameters
 - Drug properties (solubility, Peff, logP, pKa, etc.)

- Outputs:

A model that combines all available *in silico*, *in vitro* and *in vivo* information and provides:

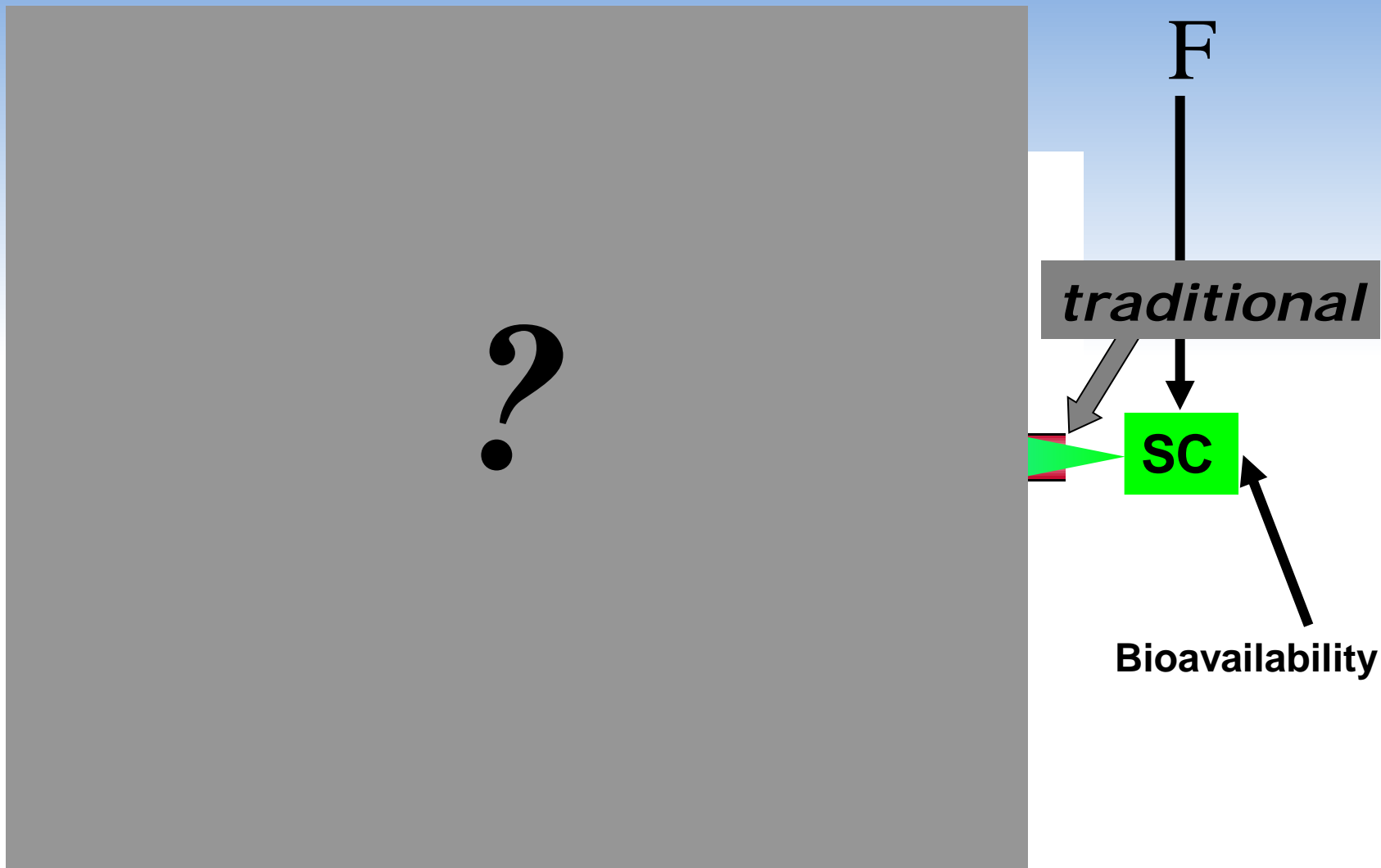
- *in vivo* dissolution, absorption and bioavailability vs. time profiles
- Description of site dependent absorption
- Description of tissue contributions to first pass extraction

Difference between traditional and mechanistic deconvolution?



* Modified from van de Waterbeemd, H, and Gifford, E. ADMET In Silico Modelling: Towards Prediction Paradise? Nat. Rev. Drug Disc. 2003, 2:192-204

Difference between traditional and mechanistic deconvolution?



* Modified from van de Waterbeemd, H, and Gifford, E. *ADMET In Silico Modelling: Towards Prediction Paradise?* Nat. Rev. Drug Disc. 2003, 2:192-204

Comparison of IVIVC Methods: Predicting PK of new products

RESEARCH PAPER

Use of *In Vitro*–*In Vivo* Correlation to Predict the Pharmacokinetics of Several Products Containing a BCS Class I Drug in Extended Release Matrices

Tahseen Mirza • Srikant A. Bykadi • Christopher D. Ellison • Yongsheng Yang • Barbara M. Davit • Mansoor A. Khan

Received: 28 February 2012 / Accepted: 9 August 2012
© Springer Science+Business Media, LLC 2012

ABSTRACT

Purpose To determine if an IVIVC model can predict PK profiles of varying formulations of a BCS Class I drug that is a salt of a weak base.

Method An IVIVC model (Level A) was created by correlating deconvoluted *in vivo* absorption data obtained from oral administration of 50 mg, 100 mg, and 200 mg fast and slow extended release formulations with *in vitro* percent dissolved using residual regression analysis. The model was then used to predict the *in vivo* profile of five test products that varied in formulation characteristics.

Results The model passed internal validation for predicted C_{max} and AUC. For external validation, *in vitro* data of five different test formulations was utilized. The model passed external validation for two test formulations that were different but belonging to the same release mechanism as that of the reference formulation. Three formulations failed external validation because they belonged to either a mixed or different release mechanism. The model and results were further confirmed using GastroPlus™ simulation software.

Conclusions These observations indicate that an IVIVC model for a BCS class I drug may be applicable to varying formulations if the principle of the drug release is similar.

KEY WORDS BCS Class I drug • convolution • deconvolution • dissolution • IVIVC

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ABBREVIATIONS

AUC area under the curve
BCS biopharmaceutics classification system
 C_{max} maximum drug concentration observed in the blood plasma profile
FRA fraction of drug absorbed into the body
FRD fraction of drug dissolved during *in vitro* experimentation
IVIVC *in vitro*–*in vivo* correlation
 k_e constant of elimination
MAPE mean absolute percentage error
rpm revolutions per minute
SUPAC-MR scale up post approval changes modified release
 V_d volume of distribution
%PE_{AUC} percent error of AUC prediction
%PE _{C_{max}} percent error of C_{max} prediction

INTRODUCTION

In vitro–*in vivo* correlation (IVIVC) has been defined by the United States Pharmacopeia (USP) Subcommittee on Biopharmaceutics as: “the establishment of a rational relationship between a biological property, or parameter derived from a biological property produced by a dosage form, and a physicochemical property or characteristic of the same dosage form” (1). The Food and Drug Administration defines IVIVC as “A predictive mathematical model describing the relationship between an *in vitro* property of an extended release dosage form (usually the

Table I Formulations Used for *In Vitro* and *In Vivo* Testing

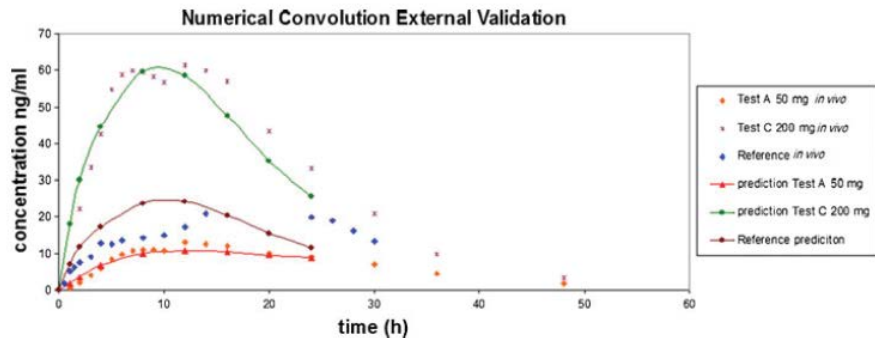
Product	Strength of dosage for <i>in vitro</i> testing	Strength of dosage for <i>in vivo</i> testing
Reference extended release	25 mg, 100 mg ^a , 200 mg	50 mg, 100 mg ^a , 200 mg
Reference fast release	100 mg	100 mg
Test A	50 mg	50 mg
Test B	200 mg	200 mg
Test C	200 mg	200 mg
Test D	200 mg	200 mg
Test E	200 mg	200 mg

^a Only used for external validation

Table III Physicochemical Properties of Drug Compound Used in Creating the GastroPlus IVIVC Model

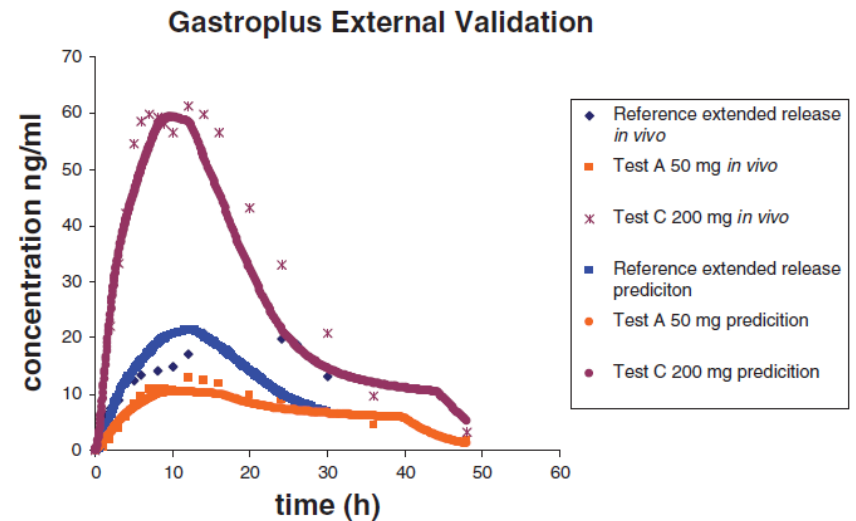
Parameters	Values
Log P	1.9
Molecular weight	261.36 g
Ph off or reference solubility fully saturated solution	5.48
Concentration of fully saturated solution	16.9 mg/ml
Mean precipitation time	5 s
Diffusion coefficient ($cm^2/s \times 10^5$)	0.74081
Drug particle density	1.2 g/ml
Particle size (diameter)	50 μ m
Human jejunal permeability (P_{eff}) ($cm/s \times 10^4$)	1.34

“External” Validation: Predicting PK of new products



Numerical Deconvolution

- Internal validation of the IVIVC showed similar prediction accuracy
 - Internal validation = applying the same products used to build the IVIVC to test it
- GastroPlus showed “greater prediction accuracy” for the new products
 - External validation = predicting PK of new products with the IVIVC



GastroPlus

IVIVC for BCS Class II (F = 66%)

AAPS PharmSciTech (© 2012)
DOI: 10.1208/s12249-012-9814-3

Research Article

Developing *In Vitro*–*In Vivo* Correlation of Risperidone Immediate Release Tablet

Yardi Saibi,^{1,3} Hitoshi Sato,¹ and Hidehisa Tachiki²

Received 6 March 2012; accepted 30 May 2012

Abstract. The present study was aimed to predict the absorption profile of a risperidone immediate release tablet (IR) and to develop the level A *in vitro*–*in vivo* correlation (IVIVC) of the drug using the gastrointestinal simulation based on the advanced compartmental absorption and transit model implemented in GastroPlus™. Plasma concentration data, physicochemical, and pharmacokinetic properties of the drug were used in building its absorption profile in the gastrointestinal tract. Since the fraction absorbed of risperidone in simulation was more than 90% with low water solubility, the drug met the criteria of class II of the Biopharmaceutics Classification System. The IVIVC was developed based on the model built using the plasma data and the *in vitro* dissolution data in several dissolution media based on the Japanese Guideline for Bioequivalence Studies of Generic Products. The gastrointestinal absorption profile of risperidone was successfully predicted. A level A IVIVC was also successfully developed in all

IVIVC for Risperidone IR Tablet

Table IV. Percent Prediction Error (PE) for Cmax and AUC of Reference Tablet

Observed values : Cmax=9.648 (ng/ml), AUC=57.83 (ng h/mL)

Dissolution Media	Cmax (ng/ml)	PE (%)	AUC (ng h/mL)	PE (%)
Phosphate buffer pH 4 (50 rpm)	10.28	-6.55	60.77	-5.08
Phosphate buffer pH 1.2 (50 rpm)	10.27	-6.45	60.77	-5.08
Phosphate buffer pH 6.8 (50 rpm)	9.94	-3.01	60.74	-5.03
Water	10.33	-7.07	60.77	-5.08
Phosphate buffer pH 6.8 (100 rpm)	9.51	1.41	60.70	-4.96

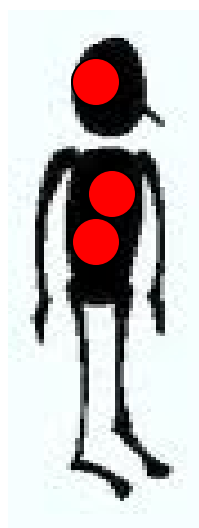
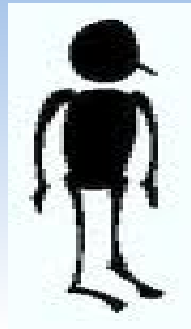
Table V. Percent Prediction Error (PE) for Cmax and AUC of Test Tablet

Observed values : Cmax=10.31 (ng/ml), AUC=62.80 (ng h/mL)

Dissolution Media	Cmax (ng/ml)	PE (%)	AUC (ng/mL)	PE (%)
Phosphate buffer pH 4 (50 rpm)	10.26	0.48	60.77	3.23
Phosphate buffer pH 1.2 (50 rpm)	10.19	1.16	60.77	3.23
Phosphate buffer pH 6.8 (50 rpm)	10.09	2.13	60.75	3.26
Water	10.35	-0.39	60.77	3.23
Phosphate buffer pH 6.8 (100 rpm)	9.88	4.15	60.73	3.29

Virtual Bioequivalence Trials

A population has variability as function of age, gender, weight, height (BMI), disease state.



You

Virtual Bioequivalence Trials

Bioequivalence trials are run to demonstrate bioequivalence between a test formulation and a reference formulation.

To demonstrate bioequivalence, the test product must duplicate the C_{max} and AUC of the reference product within 80-125% at 90% confidence intervals under both fasted and fed conditions.

The number of subjects in the trial can affect the outcome. If the number of subjects is too small, the trial might fail when the product is actually bioequivalent. If the number is too large, time and money are wasted.

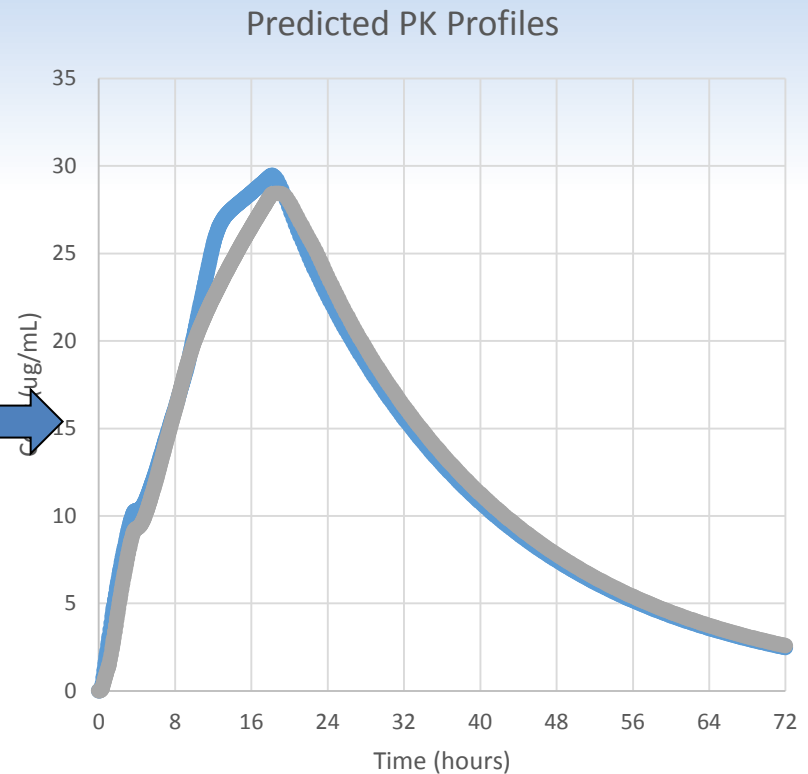
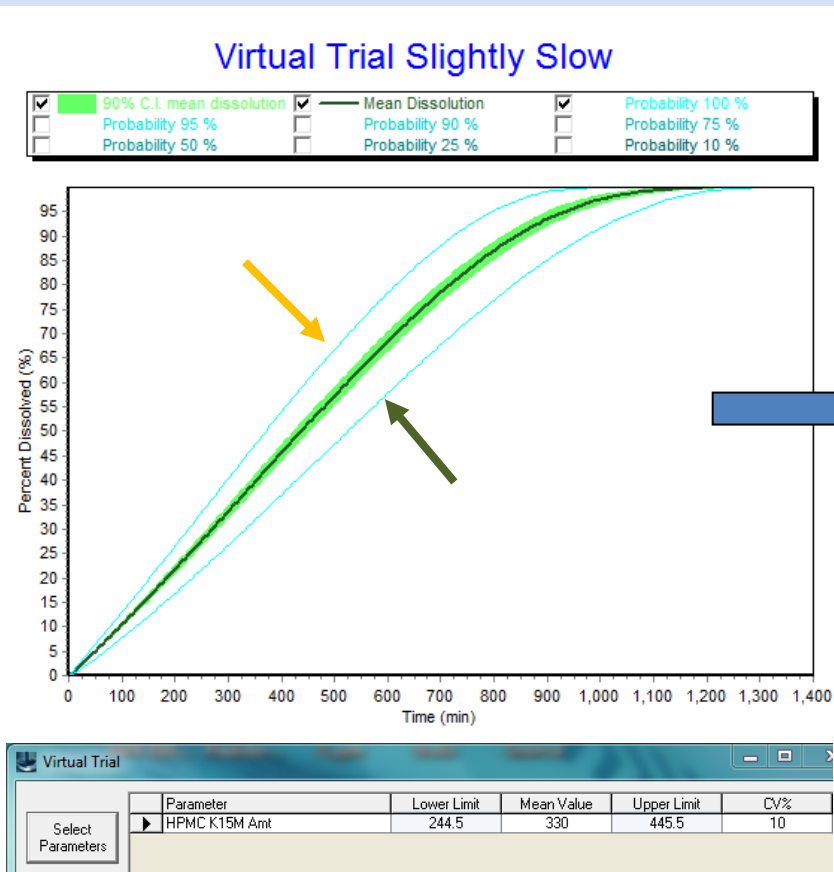
Failure of a bioequivalence trial is very expensive and time-consuming.

Virtual bioequivalence trials can help to predict whether a formulation is *likely* to pass or fail. They are not perfect, but they provide an important decision-making tool to use with all other information.

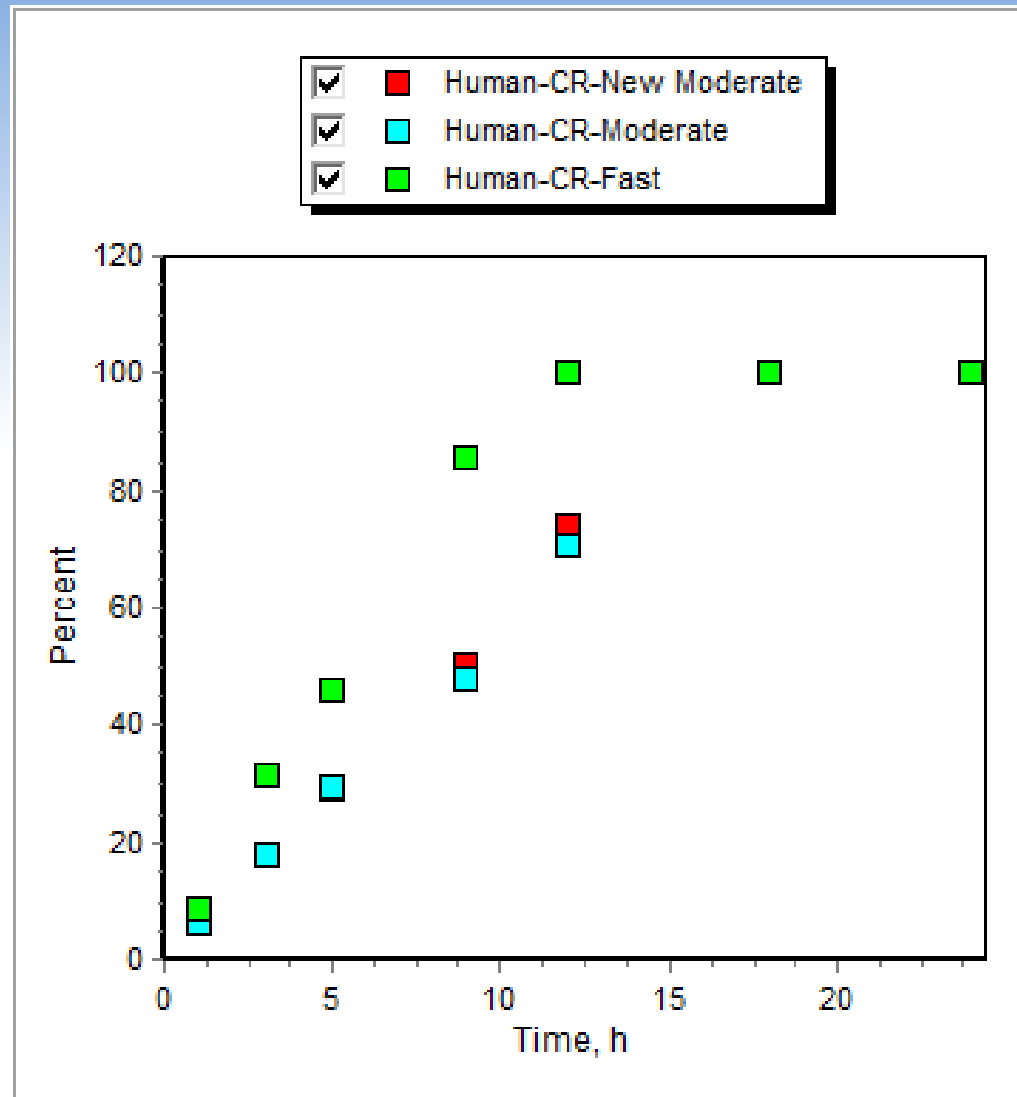
Establish dissolution specifications

10% variability around HPMC content
25 virtual lots simulated in DDDPlus

100th percentile ('extreme') dissolution profiles
loaded into GastroPlus to predict PK

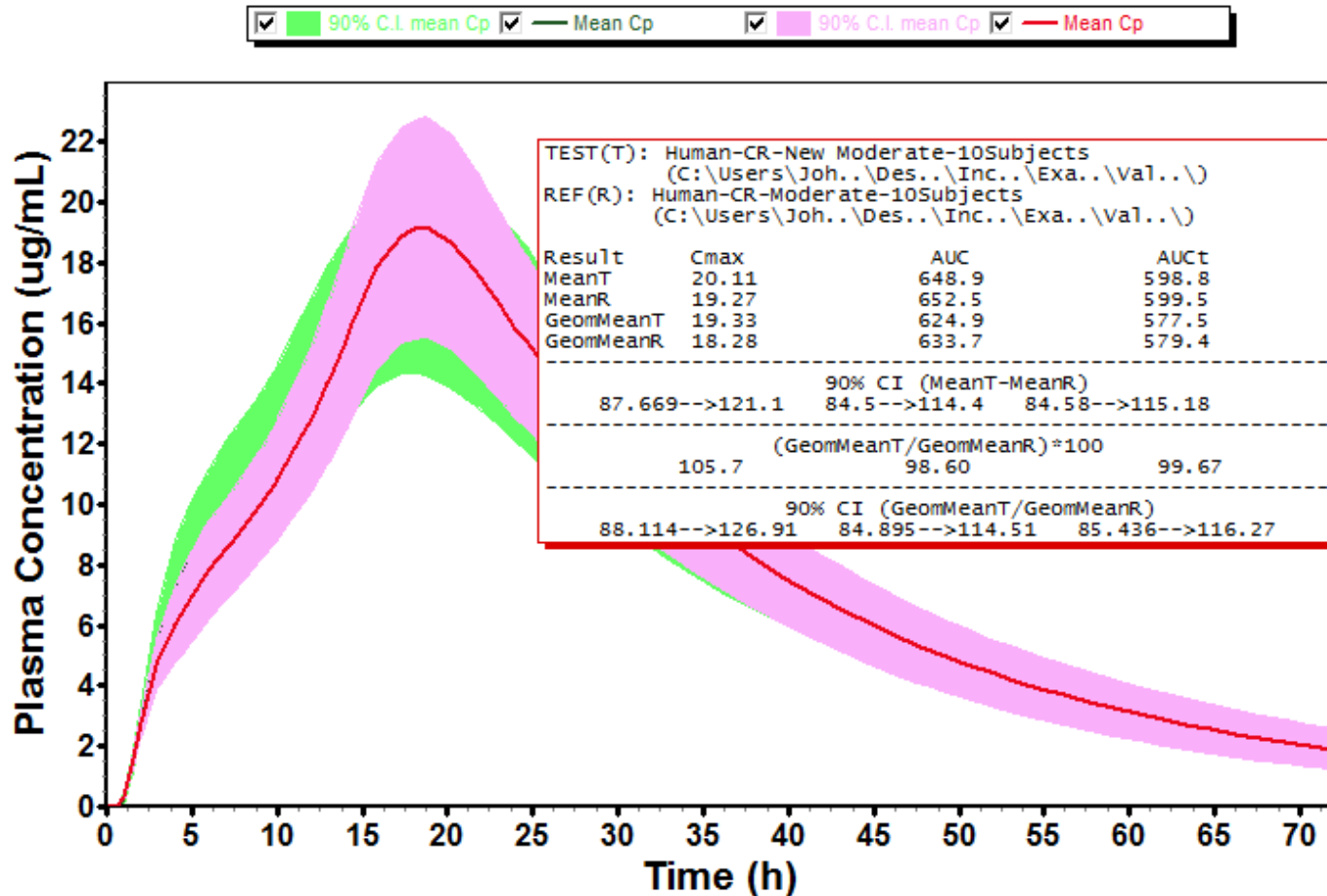


The Population Simulation



The Population Simulation

Population Simulation: Human-CR-New Moderate



Understanding food effects

BCS Predicts the likelihood and direction of a food effect 60 – 70% of the time.

Table I. Relationship Between Food Effect on the Extent of Absorption (AUC) and BCS Classification of Compounds

Food Effect/BCS	Class 1	Class 2	Class 3	Class 4	Total
Negative	9 (30%)	0 (0%)	14 (61%)	1 (9%)	24
No effect	20 (67%)	8 (29%)	7 (30%)	2 (18%)	37
Positive	1 (3%)	20 (71%)	2 (9%)	8 (73%)	31
Total	30	28	23	11	92

The number of compounds in each BCS class for a specific food effect category is listed and the percentage is provided in the parentheses.

- 67% of Class I drugs had **no** food effect.
- 71% of Class II drugs had a **positive** effect.
- 61% of Class III drugs had a **negative** effect.
- 73% of Class IV drugs had a **positive** effect.

Based on maximum absorbable dose (MAD), dose number, and log D.

Gu CH, Pharm. Res. 24 (6):1118 (2007)

Fed State – Light and High Fat Meal

GastroPlus(TM): AZD0865-VL.mdb (C:\Doc...\Viera1\Des...\GPv...\GP8.0\GP8.1\...)

File Edit Database Simulation Setup Controlled Release Tools Modules (Optional) Help

Compound Gut Physiology-Hum Pharmacokinetics Simulation Graph

Compartmental Parameters

Hum PO 1 mpk soln. Reset All Values

Excrete all un-absorbed drug at the end of gut transit time
 Zero-order gastric emptying

Compartment Data													Enzyme and Transporter Regional Distributions	
Compartment	Peff	ASF	pH	Transit Time (h)	Volume (mL)	Length (cm)	Radius (cm)	SEF	Bile Salt (mM)	Pore R (Å)	Poros/L (cm ⁻¹)	Comp. Type	3A4 Expr	3A4 Turn
Stomach	0	0.0	4.90	1.00	1000.0	30.00	10.00	1.000	0.0	2.200	2.580	Stomach	0.0	5.0E-4
Duodenum	0	2.630	5.40	0.26	40.23	15.00	1.60	4.235	14.44	10.41	48.64	Intestinal	2.09E-3	5.0E-4
Jejunum 1	0	2.616	5.40	0.95	175.3	62.00	1.50	3.949	12.02	9.640	38.90	Intestinal	3.26E-3	5.0E-4
Jejunum 2	0	2.615	6.00	0.76	139.9	62.00	1.34	3.489	10.46	8.400	26.09	Intestinal	3.26E-3	5.0E-4
Ileum 1	0	2.594	6.60	0.59	108.5	62.00	1.18	3.029	7.280	7.160	16.46	Intestinal	1.03E-3	5.0E-4
Ileum 2	0	2.574	6.90	0.43	79.48	62.00	1.01	2.569	5.990	5.920	9.540	Intestinal	1.03E-3	5.0E-4
Ileum 3	0	2.513	7.40	0.31	56.29	62.00	0.85	2.109	0.730	4.680	4.896	Intestinal	1.03E-3	5.0E-4
Caecum	0	1.416	6.40	4.50	52.92	13.75	3.50	1.790	0.0	3.920	2.915	Colon	3.1E-4	5.0E-4
Asc Colon	0	3.044	6.80	13.50	56.98	29.02	2.50	2.480	0.0	3.500	3.220	Colon	3.1E-4	5.0E-4

C1-C4: 0.06944 0.43028 0.12

Physiology: Human - Physiological - Fed

ASF Model: Opt logD Model SA/W 6.1

All properties are predictions from ADMET Predictor v6.0
 Changed pKa from AP value of 5.7 to 6.1 from Carlet-PharmRes-2010-27-21
 Changed log P from AP value of 2.44 to 4.2 from Carlet-PharmRes-2010-27-2
 Changed aqueous solubility from AP value of 19 ug/mL to 1.9 ug/mL at pH 8

pKa Table | logD: Struct-6.1 | Diss Model: Wang-Flan | PartSize-Sol: ON | BileSalt-Sol: ON | Diff: ON | ConstRad: OFF | Precip: Time | Ppara: Zhim | EHC: OFF

Main changes between Fasted and Fed state:

- Higher stomach volume
- Changes in pH (stomach and upper SI)
- Longer gastric emptying
- Higher bile salt concentrations

Fed State – Light and High Fat Meal

GastroPlus(TM): AZD0865-VL.mdb (C:\Doc..\Viera1\Des..\GPv..\GP8.0\GP8.1\...)

File Edit Database Simulation Setup Controlled Release Tools Modules (Optional) Help

Compound Gut Physiology-Hum Pharmacokinetics Simulation Graph

Compartmental Parameters

Hum PO 1 mpk soln. Reset All Values

Excrete all un-absorbed drug at the end of gut transit time
 Zero-order gastric emptying

Compartment Data													Enzyme and Transporter Regional Distributions	
Compartment	Peff	ASF	pH	Transit Time (h)	Volume (mL)	Length (cm)	Radius (cm)	SEF	Bile Salt (mM)	Pore R (Å)	Poros/L (cm ⁻¹)	Comp. Type	3A4 Expr	3A4 Turn
Stomach	0	0.0	4.90	1.00	1000.0	30.00	10.00	1.000	0.0	2.200	2.580	Stomach	0.0	5.0E-4
Duodenum	0	2.630	5.40	0.26	48.25	15.00	1.60	4.235	14.44	10.41	48.64	Intestinal	2.09E-3	5.0E-4
Jejunum 1	0	2.616	5.40	0.95	175.3	62.00	1.50	3.949	12.02	9.640	38.90	Intestinal	3.26E-3	5.0E-4
						2.00	1.34	3.489	10.46	8.400	26.09	Intestinal	3.26E-3	5.0E-4
						2.00	1.18	3.029	7.280	7.1				
						2.00	1.01	2.569	5.990	5.9				
						2.00	0.85	2.109	0.730	4.6				
						3.75	3.50	1.790	0.0	3.9				
						3.02	2.50	2.480	0.0	3.5				

Gastric emptying is expected to vary between high-fat and light meals

The fat in high-fat meal may aid in dissolution of highly lipophilic compounds.

Qh (L/min): 1.4
 Percent Fluid in SI: 40
 Colon: 10

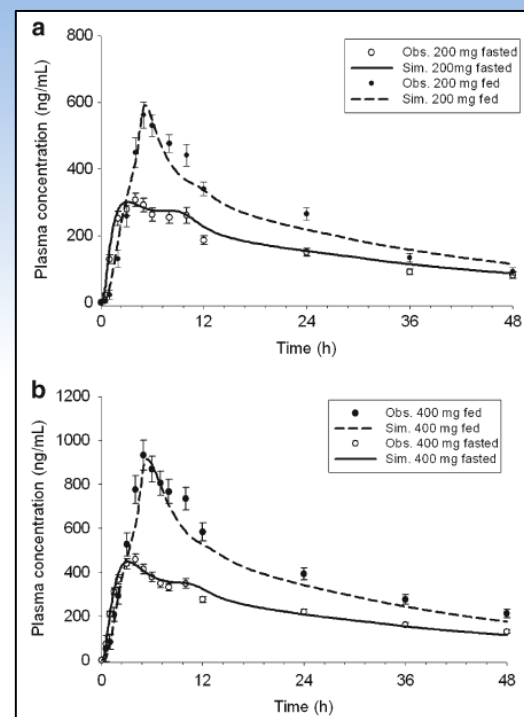
Physiology: Human - Physiological - Fed
 ASF Model: Opt logD Model SA/W 6.1

All properties are predictions from ADMET Predictor v6.0
 Changed pKa from AP value of 5.7 to 6.1 from Carlet-PharmRes-2010-27-2119-Predicting Intestinal Precipitation
 Changed log P from AP value of 2.44 to 4.2 from Carlet-PharmRes-2010-27-2119-Predicting Intestinal Precipitation
 Changed aqueous solubility from AP value of 19 ug/mL to 1.9 ug/mL at pH 8. from from Carlet-PharmRes-2010-27-2119-Predicting Intestinal Precipitation.

pKa Table | logD: Struct-6.1 | Diss Model: Wang-Flan | PartSize-Sol: ON | BileSalt-Sol: ON | Diff: ON | ConstRad: OFF | Precip: Time | Ppara: Zhim | EHC: OFF

Analyzing multiple dimensions: Design of Experiments (DoE) Approach

Parameters	Value(s)
Compound parameters	
M_w ; g/mol	>475
cLogP:	>4
pK_a (base):	3.2, 6.2
Dosage:	IR capsule
Solubility (mg/mL):	1.8 (pH 1), 0.3 (pH 2), 0.001 (pH 6.8)
Biorelevant solubility (mg/mL):	0.023 (fasted); 0.190 (fed)
Mean precipitation time (s):	450 s (fasted); 2,000 s (fed)
Effective permeability (cm/s):	1.48×10^{-4}
Particle radius of API (μm):	19
Physiological parameters	
Stomach pH	1.2 (Fasted); 1.2–4.9 (Fed)
Duodenum/jejunum pH	6.0–6.4 (Fasted); 5.4–6.0 (Fed)
Ileum pH	6.6–7.4 (Fasted); 6.6–7.4 (Fed)
Cecum–colon pH	6.4–6.8
Stomach transit time (h)	2.0 (Fasted); 5.4 (Fed)
Small intestine transit time (h)	3.3
Cecum transit time (h)	4.2
Ascending colon transit time (h)	12.6
Pharmacokinetics	
First pass extraction (%):	9.0
Blood/plasma ratio:	0.68
Plasma unbound (%):	1.6
Clearance (L/h/kg)	0.070
V_c (L/kg)	0.4
k_{12} (1/h)	0.64
k_{21} (1/h)	0.17
V_t (L/kg)	1.5



- Baseline models in GastroPlus were developed to predict the food effect for a weak base compound across different doses
- Is there an optimal combination of formulation parameters that allow us to reach our target endpoint (e.g., $F_a\%$, C_{max} , AUC)?

Analyzing multiple dimensions: Design of Experiments (DoE) Approach

- Is there an optimal combination of formulation parameters that allow us to reach our target endpoint (e.g., Fa%, Cmax, AUC)?
- Can we “design out” the food effect?

Compound: Hum 200 mg IR Cap - Fasted

Pharmacokinetics

ACAT	ACAT-Compound	Compound	Formulation
Dosing Hum 200 mg l ~ap - Fasted	Manufacture Hum 200 mg l ~ap - Fasted		

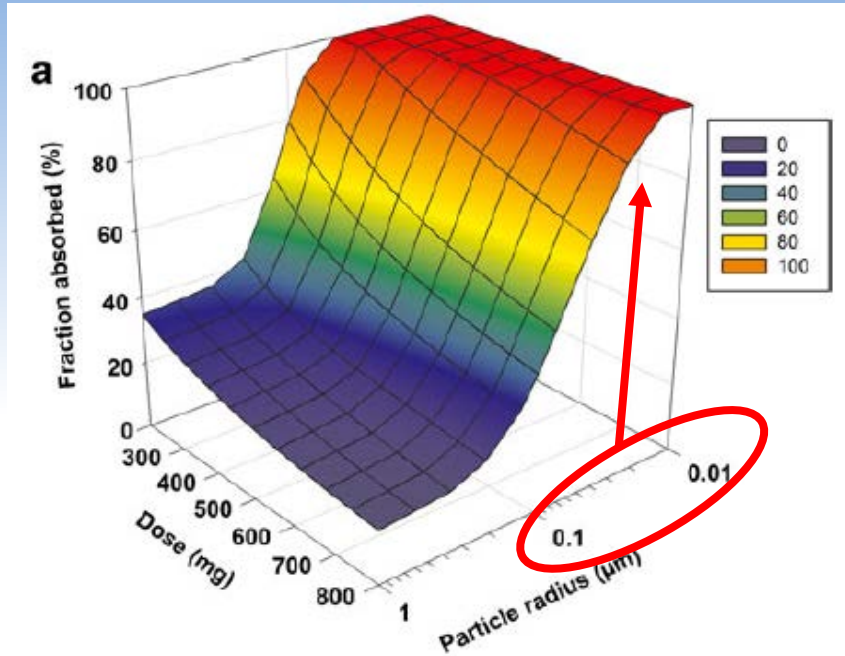
Initial Dose
 Dose Volume
 Infusion Rate

Particle Shape
 Part Radius SD
 Particle Radius
 Precip Radius
 Particle Density
 Oral ResidenceT
 Oral Lag Time
 Gastric Ret

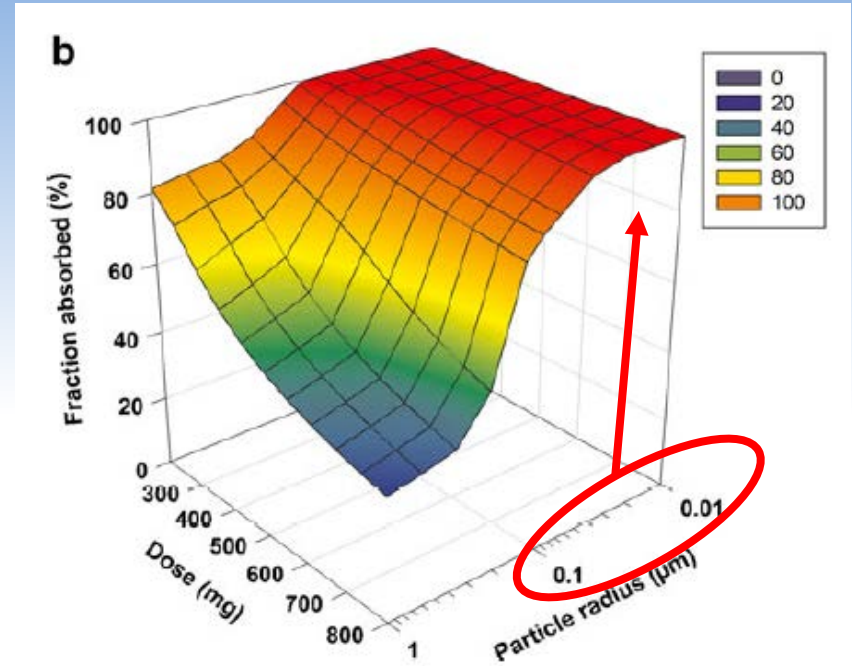
Parameter Sensitivity Analysis Setup

Select Parameters	Parameter	Lower Bound	Baseline Value	Upper Bound	Number of Test	Spacing of Param Values
	Dose of Hum 200 mg IR Cap - Fast	50	200	1000	5	Logarithmic
<input checked="" type="checkbox"/> Run 3D PSA	▶ Mean Drug Particle Radius of Hum	0.5	19	50	5	Logarithmic

3D Parameter Sensitivity Analysis



Fasted



Fed

- Parameter sensitivity analysis was run across dose and particle size together
- API particle size reduction may be useful to mitigate the food effect

Pharmaceutical Development

Research Article

Incorporation of Physiologically Based Pharmacokinetic Modeling in the Evaluation of Solubility Requirements for the Salt Selection Process: A Case Study Using Phenytoin

Po-Chang Chiang^{1,3} and Harvey Wong^{2,3}

Received 23 May 2013; accepted 26 July 2013

Abstract. In the pharmaceutical industry, salt is commonly used to improve the oral bioavailability of drugs. Currently, there is a limited understanding on the solubility requirement for improvement in oral exposure. Despite the obvious need, there is very little research mainly due to the complexity of such a system. To our knowledge, no model to guide this important process and salt solubility requirement still remains. Physiologically based pharmacokinetic (PBPK) modeling offers a means to dynamically model the processes determining oral absorption. A sensitivity analysis was performed to determine a solubility requirement for phenytoin to achieve optimal oral bioavailability for a given dose. Based on the analysis, it is predicted that a salt with solubility greater than 0.3 mg/mL would show no further increases in oral bioavailability. A screen was performed using a variety of phenytoin salts. The piperazine and hydrochloride salts showed the highest aqueous solubility and were tested *in vivo*. Consistent with the model, no significant differences in oral bioavailability for these two salts despite an increase in solubility. Our study illustrates that higher solubility salts sometimes provide no additional improvements in oral bioavailability and PBPK modeling can be utilized as an important tool to provide guidance to the salt selection and define a salt solubility requirement.

KEY WORDS: bioavailability; oral absorption; pharmacokinetic; physiological model; solubility.

Incorporating PBPK to assist with salt selection (Chiang et al., 2013)

AAAPS PharmSciTech (© 2013)
DOI: 10.1208/s12249-013-0018-2

Research Article

Theme: Leveraging BCS Classification and *in-silico* Modeling for Product Development
Guest Editors: Divyakant Desai, John Crison, and Peter Timmins

Utility of Physiologically Based Modeling and Preclinical *In Vitro/In Vivo* Data to Mitigate Positive Food Effect in a BCS Class 2 Compound

Binfeng Xia,¹ Tycho Heimbach,^{1,4} Tsu-han Lin,¹ Shoufeng Li,² Hefei Zhang,³ Jennifer Sheng,³ and Handan He¹

Received 25 March 2013; accepted 31 July 2013

Abstract. Physiologically based pharmacokinetic (PBPK) modeling has become a useful tool to estimate the performance of orally administered formulations. However, the development of a PBPK model to support formulation development with a pH-dependent and limited solubility compound, such as dry filled capsules displayed a developed and assessed in *in vitro* and *in vivo* data, the PBPK model predicted values were within $\pm 30\%$ of the observed values, illustrating that enhanced solubility formulation was found to be not statistically significant. GastroPlus population simulations also suggested that the amorphous formulation is promising in mitigating a clinically significant food effect. Overall, these efforts supported the rationale of clinical investigation of the new formulation, and more importantly, highlighted a practical application of PBPK modeling solving issues of undesirable food effects in weakly basic compounds based on preclinical *in vitro/in vivo* data.

KEY WORDS: amorphous formulation; food effects; physiologically based pharmacokinetic (PBPK) modeling; population simulation; precipitation.

Predicting food effect for BCS Class II compounds (Xia et al., 2013)

The AAAPS Journal (© 2012)
DOI: 10.1208/s12248-012-9372-3

Mini-Review

Theme: Facilitating Oral Product Development and Reducing Regulatory Burden through Novel Approaches to Assess Bioavailability/Bioequivalence
Guest Editors: James Polli, Jack Cook, Barbara Davis, and Paul Dickinson

The Use of Modeling Tools to Drive Efficient Oral Product Design

Neil R. Mathias^{1,2} and John Crison¹

Received 24 February 2012; accepted 10 May 2012

Abstract. Modeling and simulation of drug dissolution and oral absorption has become a valuable tool over the last decade to understand drug behavior *in vivo* based on the physicochemical properties of Active Pharmaceutical Ingredients (API) and dosage forms. As *in silico* and *in vitro* data become more sophisticated and our knowledge of physiological processes has grown, modeling has become a valuable confluence, tying-in *in vitro* data with *in vivo* data while offering mechanistic insights into drug performance. To a formulation scientist, this unveils not just the parameters that significantly impact dissolution/absorption, but helps probe explanations around why a formulation performs. In formulation development, modeling can be effectively used to guide: API selection (form and properties), influence clinical study design, assess dosage form performance, optimize formulation, form design, and breakdown clinically relevant conditions on dosage form performance (pH effect for patients on pH-elevating treatments, and food effect). This minireview describes examples of these applications in guiding product development including those with strategies to mitigate observed clinical exposure liability or mechanistically probe product *in vivo* performance attributes.

KEY WORDS: dissolution and absorption; drug formulation; drug development; GastroPlus; modeling and simulation.

Incorporating modeling & simulation to assist with oral product development (Mathias et al., 2012)

AAAPS PharmSciTech (© 2014)
DOI: 10.1208/s12249-014-0194-8

Research Article

Theme: Leveraging BCS Classification and *in-silico* Modeling for Product Development
Guest Editors: Divyakant Desai, John Crison, and Peter Timmins

Application of Absorption Modeling to Predict Bioequivalence Outcome of Two Batches of Etoricoxib Tablets

Amitava Mitra,^{1,3} Filippos Kesisoglou,¹ and Peter Dogterom²

Received 24 January 2014; accepted 7 August 2014

Abstract. As part of the overall product development and manufacturing strategy, pharmaceutical companies routinely change formulation and manufacturing site. Depending on the type and level of change in data and/or bioequivalence (BE) may be needed to support the change. In this report, we demonstrate that for certain weakly acidic drugs, absorption modeling could be used to predict the failure to show dissolution similarity under some conditions. Etoricoxib is described here, which was then used to *a priori* predict the BE of etoricoxib tablets manufactured at two different sites. Dissolution studies in 0.01 N HCl and pH 6.8 media failed to show comparability of the tablets and virtual trials conducted using the 0.01 N HCl C_{max} for all tablet strengths for batches manufactured at the two sites. The results were verified in a definitive bioequivalence study, which confirmed the model. Since the development of traditional *in vitro-in vivo* correlations (TIVVC) for immediate release (IR) products is challenging, in cases such as etoricoxib, absorption modeling could be used as an alternative to support waiver of a BE study.

KEY WORDS: bioequivalence; dissolution; modeling; pharmacokinetics; SUPAC.

Virtual bioequivalence trials to predict BE of different product batches (Mitra et al., 2014)

Pharmaceutical Development

RESEARCH ARTICLE – Pharmacokinetics, Pharmacodynamics and Drug Transport and Metabolism

Application of Physiologically Based Absorption Modeling for Amphetamine Salts Drug Products in Generic Drug Evaluation

ANDREW H. BABISKIN, XINYUAN ZHANG

Division of Quantitative Methods and Modeling, Office of Research and Standards, Office of Generic Drugs, Center for Drug Evaluation and Research, US Food and Drug Administration, Silver Spring, Maryland 20993

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Journal of Pharmaceutical Sciences (2015) 104, 2442–2450. © 2015 Wiley Periodicals, Inc. DOI 10.1002/jps.24474

Using M&S to predict virtual BE and assess dissolution specifications (Babiskin et al., 2015)

Extended-release (ER) drug products are widely used for the treatment of attention deficit disorder (ADHD). Physiologically based absorption modeling (PBAM) models for mixed AMP salts ER capsules and dextroamphetamine ER tablets were developed to predict virtual BE and assess BE guidance sets. Virtual BE simulations were conducted to assess BE in various populations generally conducted for approval. The models were also used to predict pharmacokinetic profiles falling within specification after the development of *in vitro-in vivo* models to test sensitivity of PK metrics to the changes in formulation variables. PBAM models are in the public domain in the USA J Pharm Sci.

Keywords: bioavailability; clinical trial simulation; modified release

RESEARCH PAPER

Use of *In Vitro-In Vivo* Correlation to Predict the Pharmacokinetics of Several Products Containing a BCS Class I Drug in Extended Release Matrices

Tahseen Mirza • Srikanth A. Bykadi • Christopher D. Ellison • Yongsheng Yang • Barbara M. Davitt • Mansoor A. Khan

Received: 28 February 2012 / Accepted: 9 August 2012
© Springer Science+Business Media, LLC 2012

ABSTRACT

Purpose To determine if an IVIVC model can predict PK profiles of varying formulations of a BCS Class I drug that is a salt of a weak base.

Method An IVIVC model (Level A) was created by combining deconvoluted *in vivo* absorption data obtained from oral administration of 50 mg, 100 mg, and 200 mg fast and slow extended release formulations with *in vitro* percent dissolved using residual regression analysis. The model was then used to predict the *in vivo* profiles of the test products that varied in formulation characteristics.

Results The model passed internal validation for predicted C_{max} and AUC. For external validation, *in vitro* data of five different test formulations was utilized. The model passed external validation for two test formulations that were different but belonging to the same release mechanism as that of the reference formulation. Three formulations failed external validation because they belonged to either a mixed or different release mechanism. The model and results were further confirmed

ABBREVIATIONS

AUC	area under the curve
BCS	biopharmaceutics classification system
C_{max}	maximum drug concentration observed in the blood plasma profile
FRA	fraction of drug absorbed into the body
FRD	fraction of drug dissolved during <i>in vitro</i> experimentation
IVIVC	<i>in vitro-in vivo</i> correlation
k_e	constant of elimination
MAPE	mean absolute percentage error
rpm	revolutions per minute
SURFC-MR	scale up post approval changes modified release
V_d	volume of distribution
%PE _{AUC}	percent error of AUC prediction
%PE _{C_{max}}	percent error of C_{max} prediction

INTRODUCTION

In vitro-in vivo correlation (IVIVC) has been defined by the United States Pharmacopoeia (USP) Subcommittee on Biopharmaceutics as “the establishment of a rational relationship between a biological property, or parameter derived from a biological property produced by a dosage form, and a physicochemical property or characteristic of the same dosage form” (1). The Food and Drug Administration defines IVIVC as “A predictive mathematical model describing the relationship between an *in vitro* property of an extended release dosage form (usually the rate or extent of drug dissolution or release) and a relevant *in vivo* response, e.g., plasma drug concentration or amount of drug absorbed” (2). In most cases, the *in vitro* property is the rate or extent of drug dissolution or release while the *in vivo* response is the plasma drug concentration

Generating mechanistic IVIVCs to predict test formulations (Mirza et al., 2012)

IVIVC model formulations

Khan

(OR)

Food and Drug Administration
Division of Biopharmaceutics II (CDER/DFSG/CD/DB)
7515 Standish Drive
Rockville, Maryland 20855, USA

Published online: 22 August 2012



Research Article

Utility of Physiologically Based Absorption Modeling in Implementing Quality by Design in Drug Development

Xinyuan Zhang,¹ Robert A. Lionberger,^{1,2} Barbara M. Davitt,¹ and Lawrence X. Yu¹

Received 16 September 2010; accepted 14 December 2010; published online 5 January 2011

Abstract. To implement Quality by Design (QbD) in drug development, scientists need tools that link drug product properties to *in vivo* performance. Physiologically based absorption models are potentially useful tools; yet, their utility of QbD implementation has not been discussed or explored much in the literature. We simulated pharmacokinetics (PK) of carbamazepine (CBZ) after administration of four oral formulations, immediate-release (IR) suspension, IR tablet, extended-release (XR) tablet and capsule, under fasted and fed conditions and presented a general diagram of a modeling and simulation strategy integrated with pharmaceutical development. We obtained PK parameters and absorption scale factors (ASFs) by deconvolution for other PK parameters. We explored three key areas used to help identify optimal critical formulations varied for the IR tablet that show decreased. Finally, virtual *in vivo* bioequivalence study may be a more sensitive predictive model is a potential

KEY WORDS: advanced compartmental absorption and transit (ACAT) model; gastroplus™; modified release (MR); quality by design (QbD).

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The role of predictive biopharmaceutical modeling and simulation in drug development and regulatory evaluation*

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Office of Generic Drugs, Food and Drug Administration, United States

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Drug development and review

ABSTRACT

Advances in predicting *in vivo* performance of drug products are developed and reviewed. Modeling and simulation in drug product development and regulatory drug review the development of biorelevant specifications, the development of products with rapid therapeutic onset, the development of products with improved safety and efficacy, better application of biopharmaceutical modeling in drug product development, regulatory challenges in bioequivalence demonstration of complex drug products also present exciting opportunities for creative modeling and simulation approaches. A collaborative effort among academia, government and industry in modeling and simulation will result in improved safe and effective new/generic drugs to the American public.

Published by Elsevier B.V.

Role of M&S in drug development and regulatory evaluation (Jiang et al., 2011)

*Re-engineered formulations and
“virtual” bioequivalence:
A successful biowaiver case study*

Objectives

- Post approval, sponsor's manufacturing process change resulted in different particle size distributions for new lots
- With GastroPlus, could they apply for a biowaiver by:
 - assessing the effects of changes in particle size distribution of the active pharmaceutical ingredient (API) on its oral bioavailability?
 - predicting the virtual bioequivalence between the “new” and “old” API lots?

Tasks

- Determine the most appropriate absorption/PBPK model for the API across several doses for the non-engineered lots
- Assess the effect of particle size on API exposure for the immediate release formulation
- Evaluate predicted bioequivalence of the tablets manufactured with particle-engineered (PE) API (narrower particle size distribution) versus the tablets manufactured with non particle-engineered (NPE) API (broader particle size distribution)

Formulation Specifications

Various Particle Size Used in Clinical Studies

NPE API Lot Number	d10 (µm)	d50 (µm)	d90 (µm)
NPE Lot 1	20	63	173
NPE Lot 2	8	179	512
NPE Lot 3	15	49	142
NPE Lot 4	31	86	348
NPE Lot 5	26	78	276
NPE Lot 6	9	29	101
NPE Lot 7	11	35	114
NPE Lot 8	12	37	124
NPE Lot 9	10	36	119
NPE Lot 10	13	45	138
NPE Lot 11	11	35	99

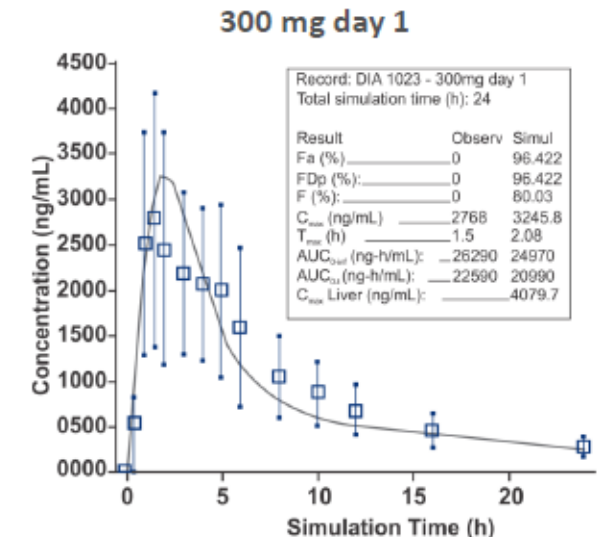
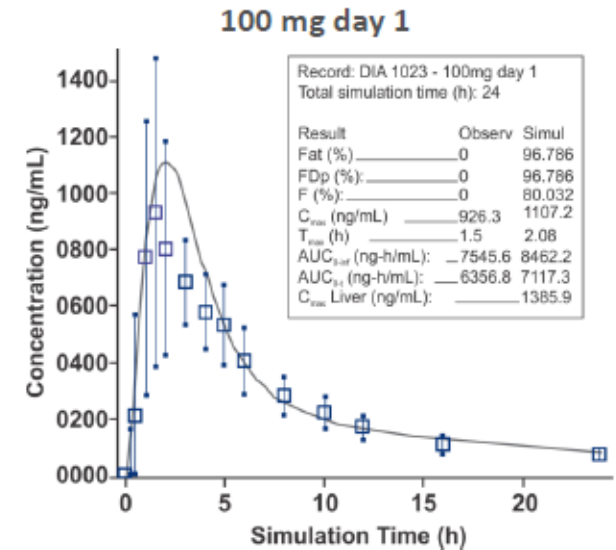
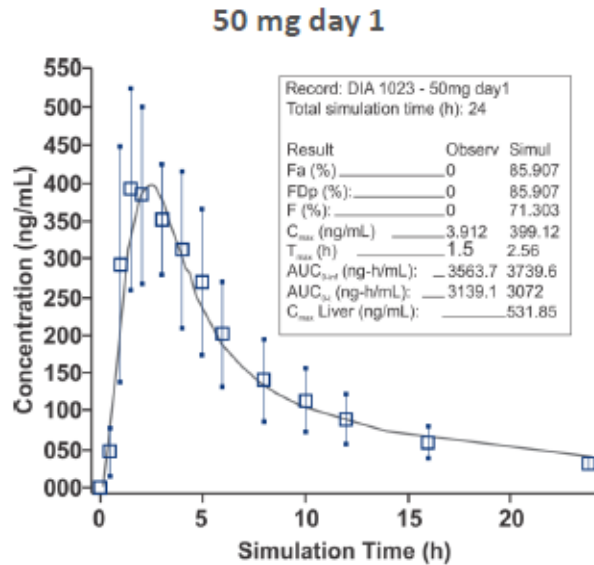
PE API Lot Number	d10 (µm)	d50 (µm)	d90 (µm)
PE Lot 1	16	40	88
PE Lot 2	20	49	102
PE Lot 3	22	53	108
PE Lot 4	19	39	71
PE Lot 5	17	35	67
PE Lot 6	23	48	93
PE Lot 7	21	44	87
PE Lot 8	21	45	90
PE Lot 9	24	50	94
PE Lot 10	21	45	89
PE Lot 11	19	42	88
PE Lot 12	22	47	95

API: active pharmaceutical ingredient; d10, d50, and d90: diameter for which 10%, 50%, and 90% (respectively) by volume of the particles are less than this value;
 NPE: non-particle-engineered; PE: particle-engineered

Part I: Model Validation

Model Validation

Simulated (lines) and experimental (points)



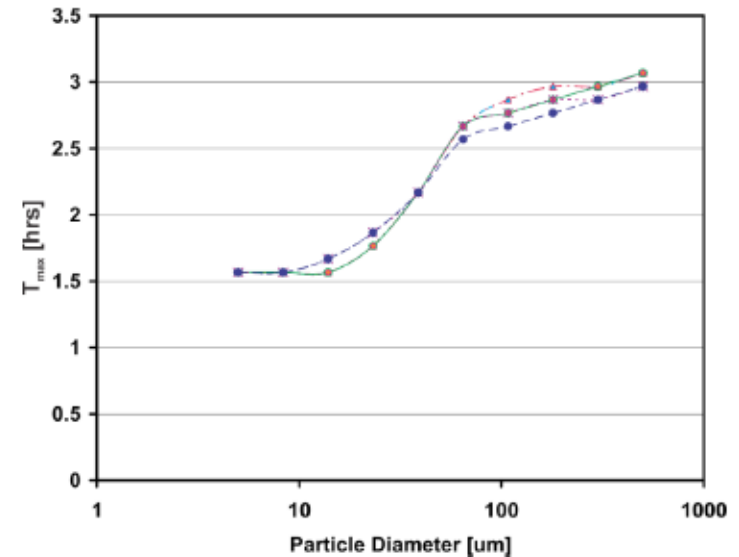
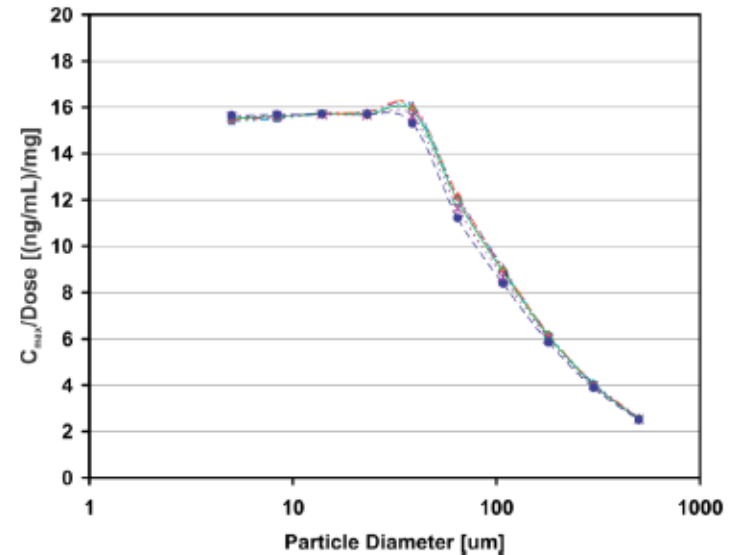
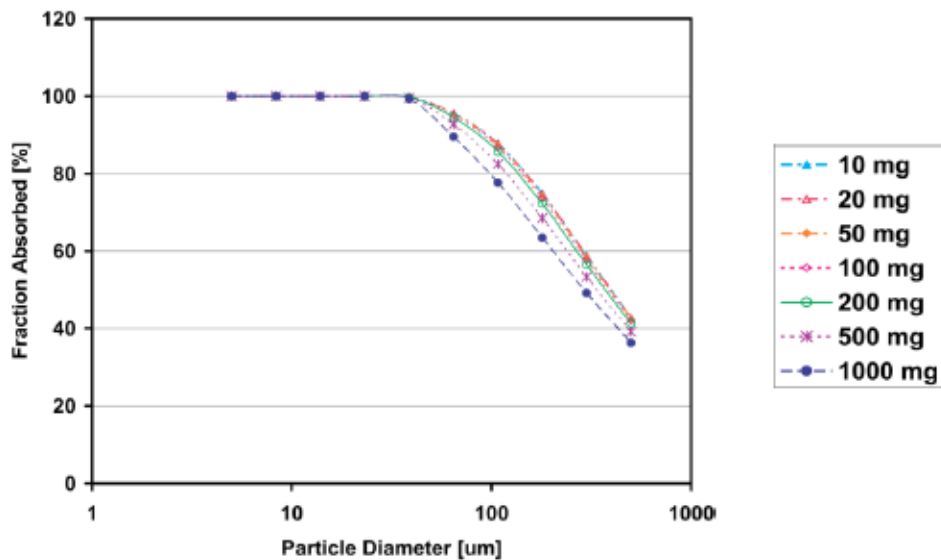
Excellent match between the measured and predicted Cp time profiles for 50, 100, and 300 mg doses

Cp-time: plasma concentration time

Part II: Parameter Sensitivity Analysis

Parameter Sensitivity Analysis

Change in $F_a\%$, C_{max} , T_{max} as a function of mean particle diameter



C_{max} : maximum observed plasma concentration; $F_a\%$: fraction absorbed;
 T_{max} : time to reach C_{max}

Part III: Virtual BE Simulations

Virtual Bioequivalence Study Simulations

- Using crossover virtual trial simulation comparing different formulations (PK parameters: C_{\max} and AUC)

API Lot	NPE or PE	d10 (μm)	d50 (μm)	d90 (μm)
Lot 1	NPE	26	78	276
Lot 2	NPE	11	35	99
Lot 3	NPE	14	43	116
Lot 4	NPE	11	32	91
Lot 5	PE	17	41	88

API: active pharmaceutical ingredient; AUC_{∞} : area under the plasma concentration-time curve; C_{\max} : maximum observed plasma concentration; d10, d50, and d90: diameter for which 10%, 50%, and 90% (respectively) by volume of the particles are less than this value; NPE: non-particle-engineered; PE: particle-engineered; PK: pharmacokinetics

Part III: Virtual BE Simulations

Virtual Bioequivalence Study Simulations

API Lot	PE/NPE	Dose (mg)	AUC _∞ (ng.h/mL) (N=250)		C _{max} (ng/mL) (N=250)	
			GM	GMR (90% CI)	GM	GMR (90% CI)
Lot 5	PE	50	4180	113.3 (110.7, 116.1)	551	139.3 (136.0, 142.7)
Lot 1	NPE	50	3688		395	
Lot 5	PE	100	8242	103.0 (100.9, 105.1)	551	106.4 (104.3, 108.6)
Lot 3	NPE	100	8001		395	
Lot 5	PE	300	24998	102.2 (99.8, 104.6)	3118	100.0 (97.7, 102.4)
Lot 2	NPE	300	24460		3117	
Lot 5	PE	100	8242	98.2 (96.2, 100.2)	1068	95.1 (93.2, 97.0)
Lot 4	NPE	100	8395		1123	
Lot 5	PE	300	24998	101.9 (99.8, 104.1)	3118	98.3 (96.3, 100.4)
Lot 4	NPE	300	24525		3171	

API: active pharmaceutical ingredient; AUC_∞: area under the plasma concentration-time curve from time 0 to infinite time; CI: confidence interval; C_{max}: maximum observed plasma concentration; GM: geometric mean; GMR: geometric mean ration; NPE: non-particle-engineered; PE: particle-engineered

Summary

- A mechanistic, physiologically-based absorption/PK model was constructed in GastroPlus and validated across three dose levels (50, 100, and 300 mg) using *in vivo* data collected from tablets manufactured with non-particle-engineered API.
- Parameter sensitivity analysis showed that mean particle size would be the main property that determines whether formulations are likely to be bioequivalent, regardless of dose.
- Virtual bioequivalence trial simulations showed that, for a sufficiently powered study, the population-derived C_{\max} and AUC values would be bioequivalent between the tablets manufactured with non-particle-engineered (NPE) vs. new-particle-engineered (PE) API, regardless of the dose.
- **Regulatory agencies approved the sponsor's biowaiver application**

How Modeling & Simulation Can Save Resources in Generic R&D

- Understand the mechanisms that affect the absorption/PK of reference products earlier
 - Gain unique insight into the release kinetics & establish better targets
- Guide formulation & dissolution method design
 - Improve chances for success in follow-up pilot studies
- Estimate population behaviors before running clinical trials (*virtual bioequivalence trials*)
 - Separate formulation & physiological effects
- Ultimate goal:

Reduce “trial and error”