



# GastroPlus<sup>®</sup>



Jim Mullin

Part I: Basic model considerations  
& food effect predictions



Ke Szeto

# 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)

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

Xinyuan Zhang,<sup>1</sup> Robert A. Lionberger,<sup>1,2</sup> Barbara M. Davit,<sup>1</sup> and Lawrence X. Yu<sup>1</sup>

### GastroPlus simulations of fed and fasted: PO 200, 300, & 400 mg Carbamazepine

Table III. Comparison of Predicted vs. Observed Mean Plasma PK Parameters

Parameters		Suspension		IR tablet		XR tablet		XR capsule	
		Obs.	Pred.	Obs.	Pred.	Obs.	Pred.	Obs.	Pred.
Dose (mg)		200		400		400		300	
$C_{max}$ (ng/mL)	Fasted	3066.7	2914.6	3610.1	3713.7	3005.2	3105.2	2066.1	2120.8
	Fed	2580.0	2506.9	5920.0	5501.2	3329.9	3786.6	2661.9	2798.5
$AUC_t$ ( $\mu\text{g}\times\text{h/mL}$ )	Fasted	166.6	163.3	279.8	301.6	270.4	263.7	194.3	190.2
	Fed	165.2	152.9	401.3	348.6	286.1	288.6	221.1	222.2
$AUC_{inf}$ ( $\mu\text{g}\times\text{h/mL}$ )	Fasted	179.8	177.3	298.6	330.0	285.8	272.2	226.4	203.8
	Fed	180.7	166.2	444.9	379.3	304.7	297.5	246.7	237.0
$T_{max}$ (hr)	Fasted	1	1.45	24	16	24	28.9	26	20.98
	Fed	4	3.66	12	4.8	24	16.4	15	17.8
$POT_{20}$ (hr)	Fasted	[0.6,8.5]	[0.8,10]	[3.7,41]	[2.9,40]	[10,42]	[13,44]	[8.2,54]	[8.9,48]
	Fed	[1.1,16]	[1.4,16]	[3.5,28]	[2.3,19]	[8.1,42]	[10,34]	[7.3,44]	[8.8, 39]
$F_a$ (%)	Fasted	N.A.	99.9	N.A.	93.0	N.A.	71.6	N.A.	76.6
	Fed	N.A.	99.9	N.A.	99.8	N.A.	78.2	N.A.	89.0
Correlation coefficient ( $R^2$ )	Fasted	0.956		0.975		0.974		0.977	
	Fed	0.940		0.876		0.954		0.991	

$POT_{20}$  peak occupancy time, time span over which the concentration is within 20% of  $C_{max}$ ;  $F_a$  fraction absorbed; N.A. not available

# Example: Class II Drug – impact of particle size changes under fasted vs. fed conditions

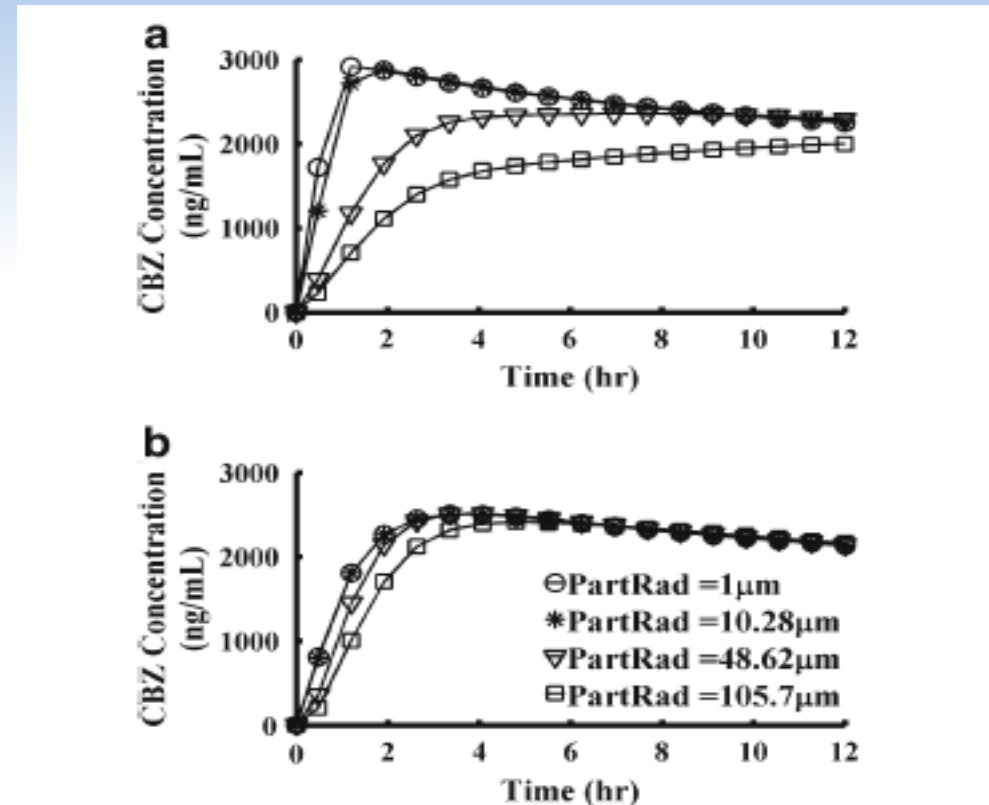
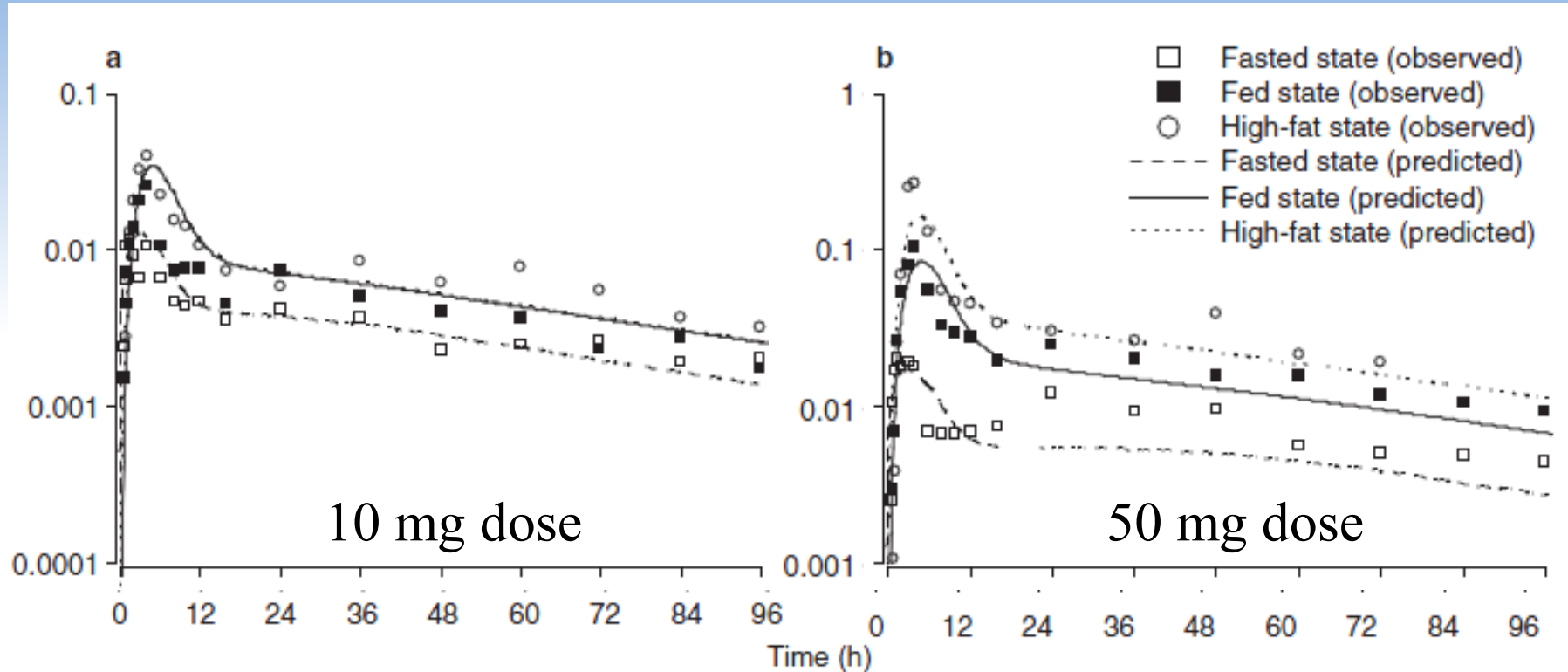


Fig. 7. PK profiles sensitivity to the mean particle radius is different under fasted and fed state. **a** IR suspension, fasted state; **b** IR suspension, fed state. Legend in (b) is also applied to (a)

# Cmd3

(basic pKa = 4.2, log P = 6.6, FaSSIF = 1.3 mg/mL)



1.5- to 2.4-fold increase in AUC under fed conditions due to increased solubility at higher bile salt concentrations

What changes are accounted for  
between fasted and fed state  
models in GastroPlus?

# Fed State – GI Physiology

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 <sup>-1</sup> )	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	175.2	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 0.12147 0.46632

Physiology: Human - Physiological - Fed

Qh (L/min): 1.4

Percent Fluid in SI: 40

Colon: 10

Main changes between Fasted and Fed state:

- Higher stomach volume
- Changes in pH (stomach and upper SI)
- Longer gastric emptying
- Higher bile salt concentrations
- Higher liver blood flow (needs to be manually changed with PBPK models)

# Solubility – Bile Salt

To account for physiological ‘distribution’ of bile salts, GastroPlus uses published equation based on concentration of bile salts in media and compound’s affinity to bile salt micelles

$$Sol_{bile,pH} = Sol_{aq,pH} \left( 1 + \frac{Mwt_{H_2O}}{\rho_{H_2O}} \times SR \times C_{bile} \right)$$

*in vivo* solubility in compartment with specific pH and bile salt concentration

Buffer solubility at given pH calc from reference solubility, pKa(s) and solubility factor(s)

Molecular weight and density of water

Bile salt solubilization ratio – represents drug’s affinity to bile salt micelles  
*This parameter is required to account for physiological effect of bile salts on solubility and dissolution rate*

*in vivo* concentration of bile salts in given compartment (physiological parameter)



# Built-in Fed Physiologies for Different Meal Types

GastroPlus(TM): GastDemo0.mdb (C:\Users\jmullin\Docum...\CodeR...\Gastr..)

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

Compound: Propranolol HCl

Gut Physiology-Hum Pharmacokinetics Simulation Graph

Compartmental Parameters

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

Peff	ASF	pH	Transit Time (h)	Volume (mL)	Length (cm)	Radius (cm)	SEF	Bile Salt (mM)
0	0.0	4.90	2.45	978.5	29.19	9.87	1.000	0.0
0	2.721	5.40	0.28	44.57	14.56	1.56	4.235	22.28
0	2.668	5.40	0.94	166.6	60.26	1.48	3.949	18.09
0	2.665	6.00	0.74	131.0	60.26	1.32	3.488	14.99
0	2.640	6.60	0.58	102.0	60.26	1.16	3.029	10.14
0	2.621	6.90	0.42	75.35	60.26	1.00	2.569	7.093
0	2.589	7.40	0.29	53.57	60.26	0.84	2.109	1.049
0	0.352	6.40	4.36	50.49	13.50	3.45	1.790	0.0
0	0.823	6.80	13.07	53.55	28.35	2.45	2.480	0.0

Enzyme and Transporter Regional Distributions

C1-C4: 0.06944 0.43028 0.12147 0.4663

Physiology: Human - Physiological - Fed

ASF Model: Opt logD Model SA/V 6.1

Fed Meal Options

Percent Fluid in SI: 40 Colon: 10

Biorelevant solubilities from ADMET Predictor v6.1

pKa Table | logD: Struct-6.1 Diss Model: Johnson PartSize-Sol: ON BileSalt-Sol: ON | Diff: ON ConstRad: ON Precip: Time Ppara: OFF EHC: OFF ACAT: Co

- Link gastric emptying time to meal calories
- Account for effect of fat content on bile salt concentration

Fed State Model

Fed State Model: Default

Meal Calories: 233.68 % Fat in Meal: 30.00

Current gastric transit time of 1.00 hr.

Current duodenum bile salt concentration is 14.44 mM.

Cancel OK

Fed State Model

Fed State Model: Default

Meal Calories: 233.68

Current gastric transit time of 1.00 hr.

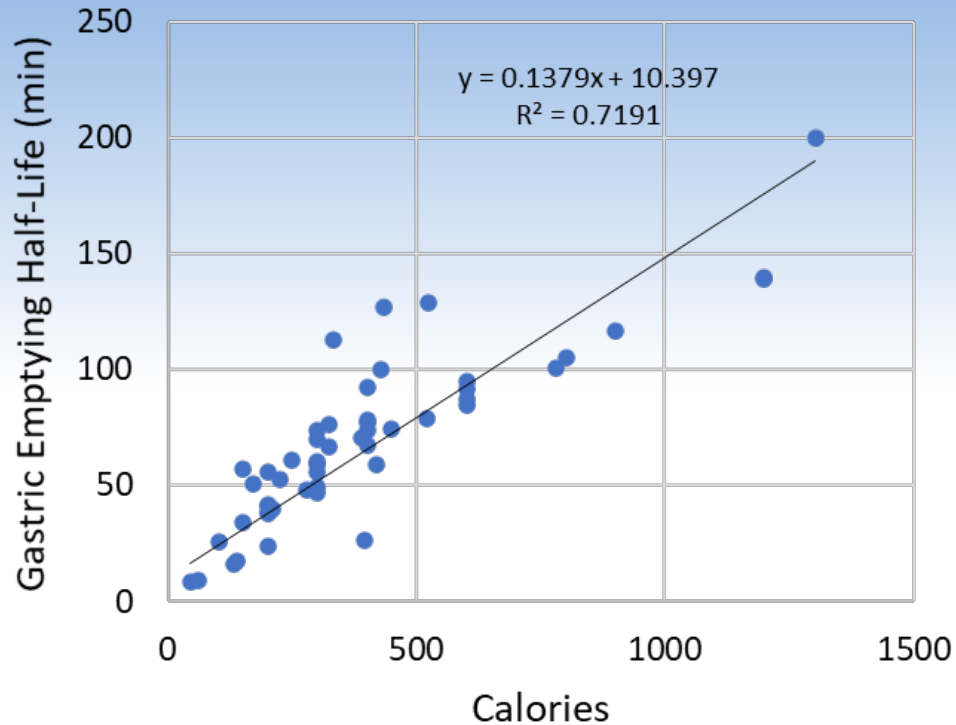
Current duodenum bile salt concentration is 14.44 mM.

Default  
User-Defined Fat and Calories  
FDA Breakfast Meal  
Low Fat - Low Calorie Meal  
Low Fat - Moderate Calorie Meal  
Low Fat - High Calorie Meal  
Moderate Fat - Low Calorie Meal  
Moderate Fat - Moderate Calorie Meal

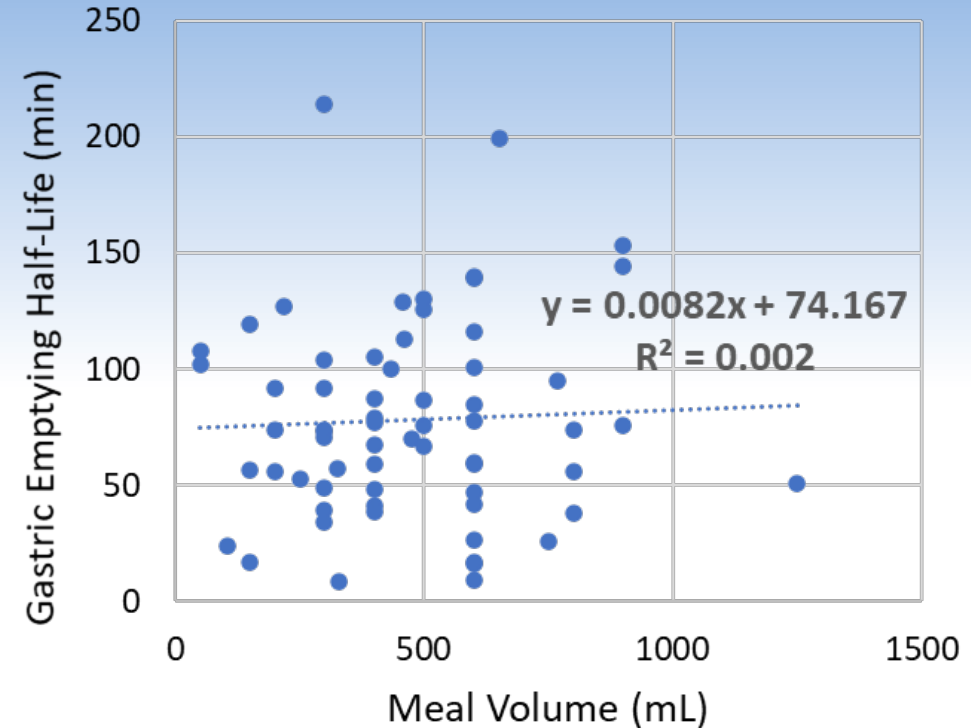
Cancel OK

# Adjusting Fed State Based on Calories and Meal Volume

## Gastric Emptying vs. Meal Calories

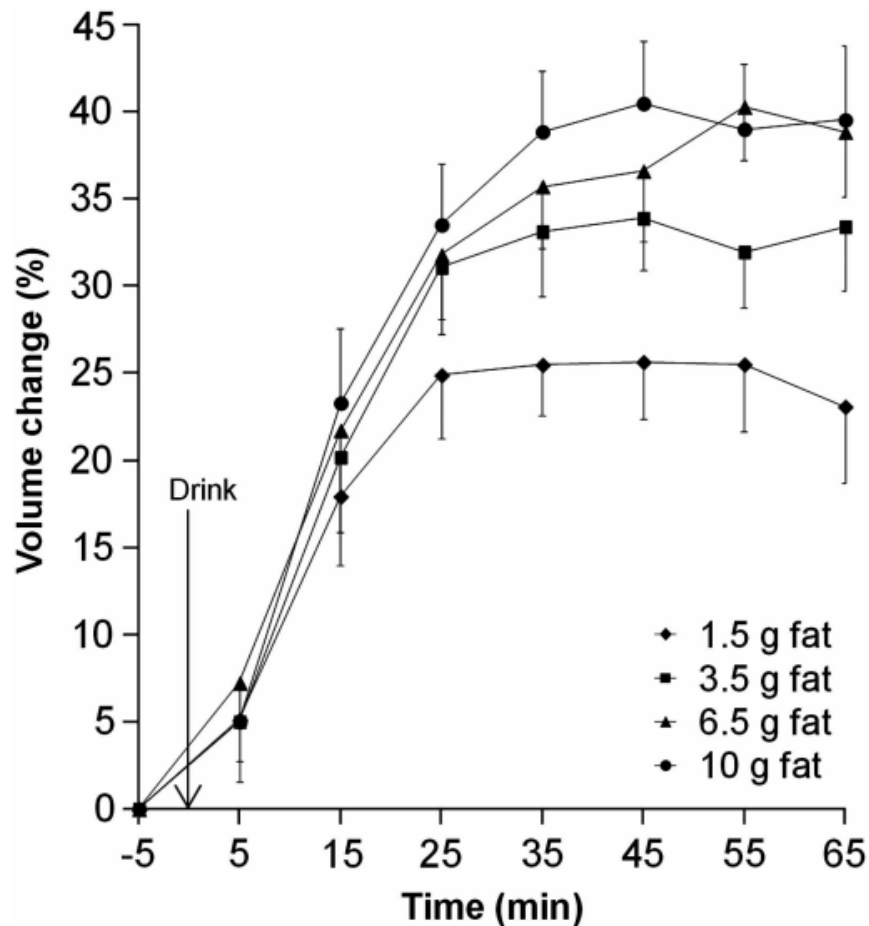


## Gastric Emptying vs. Meal Volume



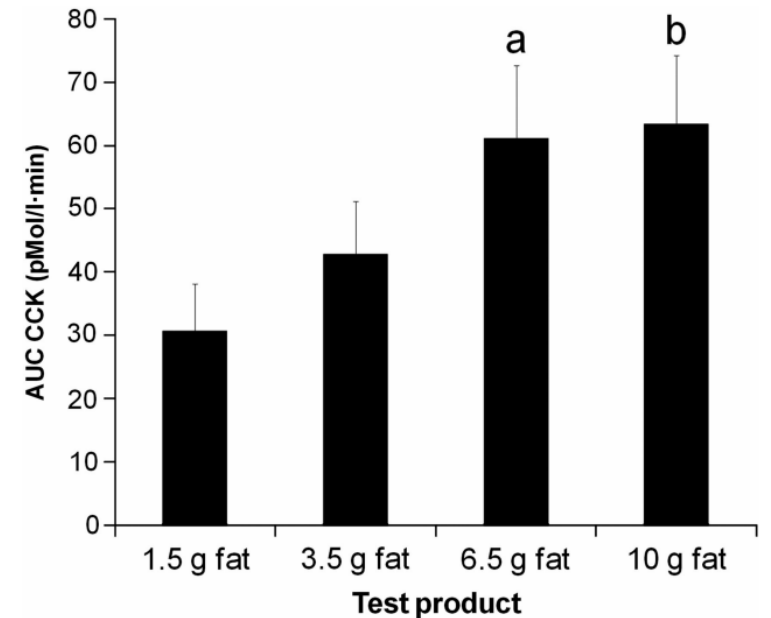
- Meal value is not well correlated with emptying time
- Gastric Emptying is linearly related to calories and can be slightly modified by meal type (fat, protein, carbohydrates)
  - Smaller impact and not currently considered due to lack of data

# Adjusting Fed State for Biliary concentration



**Figure 3.** Plot of the % gall bladder volume change with time for healthy volunteers after they consumed each of the four milk-based beverages of increasing fat content in Study 2. Values are mean  $\pm$  s.e.m.,  $n = 21$ .

- With increasing fat, gall bladder excretion volume increases
- Gall bladder secretion is mediated by hormone cholecystinin CCK



**Figure 4.** Area under the curve (AUC) for plasma cholecystinin (CCK) for healthy volunteers after they consumed each of the four milk-based beverages of increasing fat content in Study 2. Values are mean  $\pm$  s.e.m.,  $n = 17$ . <sup>a</sup>Significant difference versus 1.5g fat drink ( $P < 0.05$ ); <sup>b</sup>Significant difference versus 1.5g fat drink ( $P < 0.01$ ).

# Results of Dynamic Bile-Salt Model

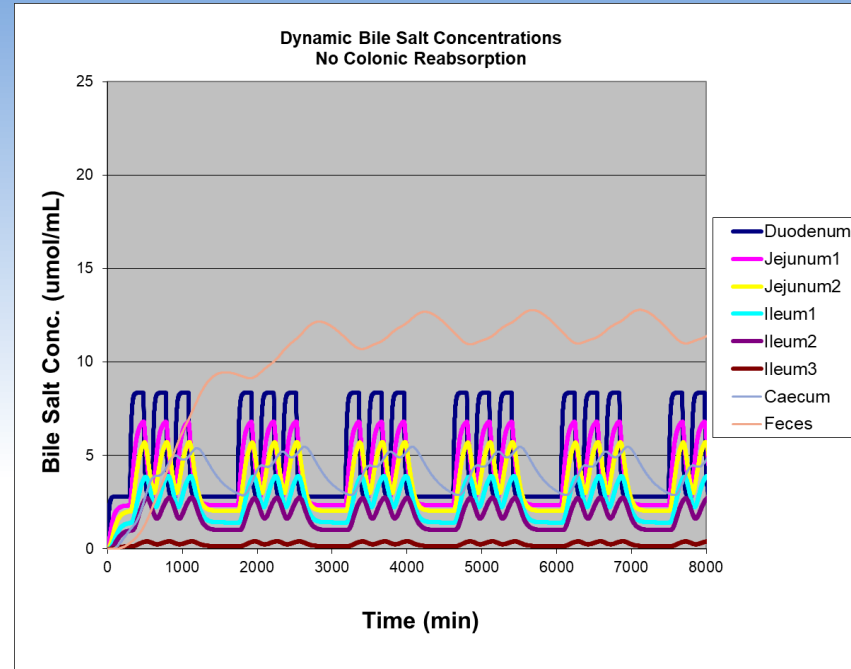
## Bile Salt Concentration Spreadsheet Model:

- A caecum and feces (large intestine) compartment were added using same volumes and transit times as G+
- No bile reabsorption assumed for caecum and colon

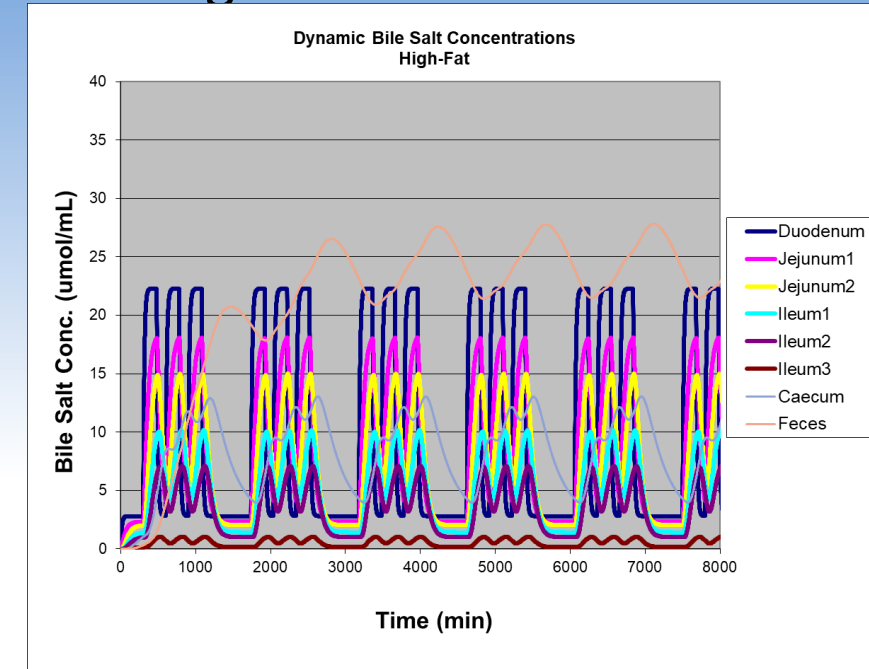
Feces Bile Concentration Reference:

Reddy, CANCER RESEARCH, 1981, 41, 3766-3768

## Low-Fat Bile Prediction



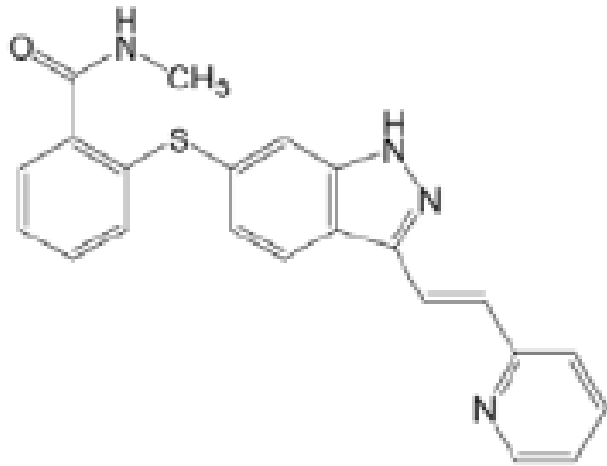
## High-Fat Bile Prediction



- Assumptions, 3 meals and fed-state occurs for 3 hours duration
- Then the bile production rate was fit to make fold change in bile in feces the same as literature data

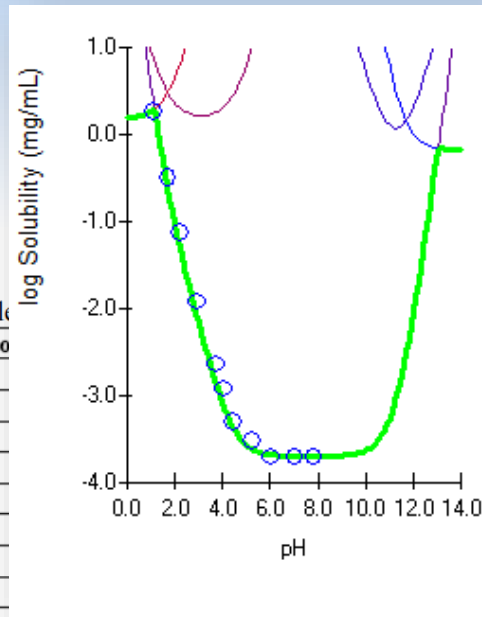
			Bile Concentration in Each Compartment (mM)					
	% Fat	Fold Change Bile in Feces	Duodenum	Jejunum1	Jejunum2	Ileum1	Ileum2	Ileum3
<b>Low Fat</b>	10	1.000	8.36	6.83	5.72	3.90	2.74	0.41
<b>Normal</b>	30	1.476	14.55	11.83	9.84	6.67	4.67	0.69
<b>Bile High Fat</b>	55	2.072	22.28	18.09	14.99	10.14	7.09	1.05

# Case Study: Axitinib



**Table 21.** Solubility of Axitinib in Aqueous Media as a Function of pH at 20°C for at k

Aqueous Solution	Solubility (micro)
Water pH 7.5	0.2
0.1N HCl, pH 1.1	1841
0.06N HCl pH 1.7	320
0.02M Sodium Phosphate/0.02M Sodium Acetate pH 2.2	75
0.02M Sodium Phosphate/0.02M Sodium Acetate pH 2.9	12
0.02M Sodium Phosphate/0.02M Sodium Acetate pH 3.7	2.3
0.02M Sodium Phosphate/0.02M Sodium Acetate pH 4.0	1.2
0.02M Sodium Phosphate/0.02M Sodium Acetate pH 4.4	0.5
0.02M Sodium Phosphate/0.02M Sodium Acetate pH 5.2	0.3
0.02M Sodium Phosphate/0.02M Sodium Acetate pH 6.0	0.2
0.02M Sodium Phosphate/0.02M Sodium Acetate pH 7.0	0.2
0.02M Sodium Phosphate/0.02M Sodium Acetate pH 7.8	0.2



Property	Value	Ref
LogP	3.91	AP
pKa	1.71, 4.5 (Base) 11.68, 10.84 (Acid)	AP
Reference solubility	1.48E-3 @ pH 7.02	AP
Exp Sol.	1.841 mg/mL @ pH 1.1 0.0002 mg/mL @ pH 7.8	[1] [1]
Solubility Factor	3332.93 (AP) 7600(Exp)	AP, [1]
FaSSIF Sol.	0.001	[2]
FeSSIF Sol.	0.114	[2]
Human Peff (from Pampa)	2.89	[2]
Blood:plasma concentration ratio ( $R_{bp}$ )	0.8	[1]
Plasma protein binding ( $F_{up}$ )	1	[1]
Particle Size	4 $\mu$ m (SD = 3)	[3]
<b>Metabolism</b>		
CL (3A4)	21 L/hr	[1]
Vd	68 L	[1]

[1] FDA application 202324 Clin Pharm and Biopharm. Review

[2] Dodd, Journal of Pharmaceutical Sciences, 2019, 108, 87-101

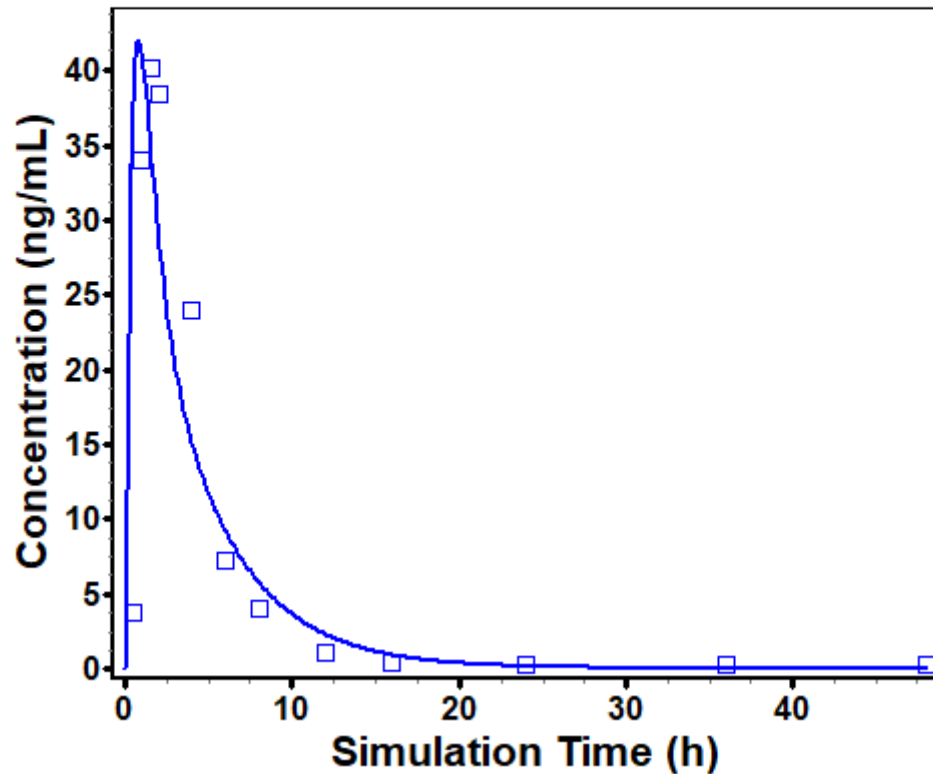
[3] Reynor, Drug Metabolism and Disposition (2013): dmd-113.

Axitinib is a case study for gastric emptying because of high permeability and high solubility in gastric. Precipitation doesn't occur even though Axitinib is a low solubility weak base.

# Case Study: Axitinib Fasted State and Baseline Fed Model

axitinib 5mg Fasted Tab

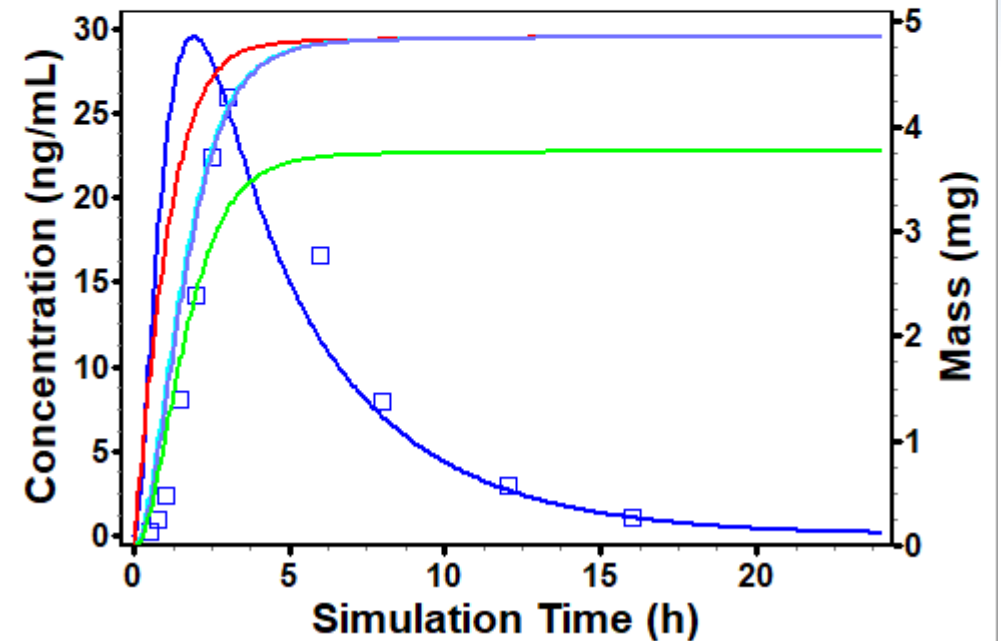
- Cp-Venous Return-axitinib 5mg Fasted Tab
- Cp-Venous Return-axitinib 5mg Fasted Tab Obs
- ▭ Cp-Venous Return-axitinib 5mg Fasted Tab Err



Pithavala, Cancer Chemother Pharmacol (2012) 70:103–112

axitinib 5mg Fed Tab High Fat

- Cp-Venous Return-axitinib 5mg Fed Tab High Fat
- Cp-Venous Return-axitinib 5mg Fed Tab High Fat Obs
- ▭ Cp-Venous Return-axitinib 5mg Fed Tab High Fat Err
- Amount Dissolved-axitinib 5mg Fed Tab High Fat
- Amount Absorbed-axitinib 5mg Fed Tab High Fat
- Amount Portal Vein-axitinib 5mg Fed Tab High Fat
- Tot entered SC-axitinib 5mg Fed Tab High Fat



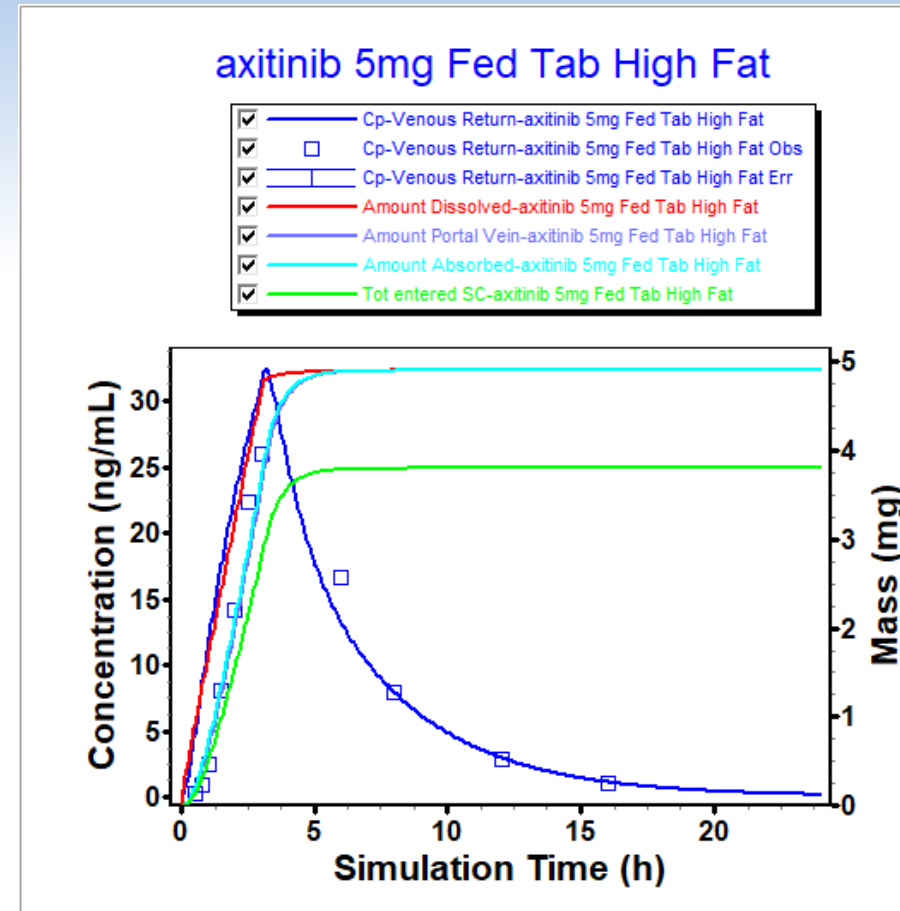
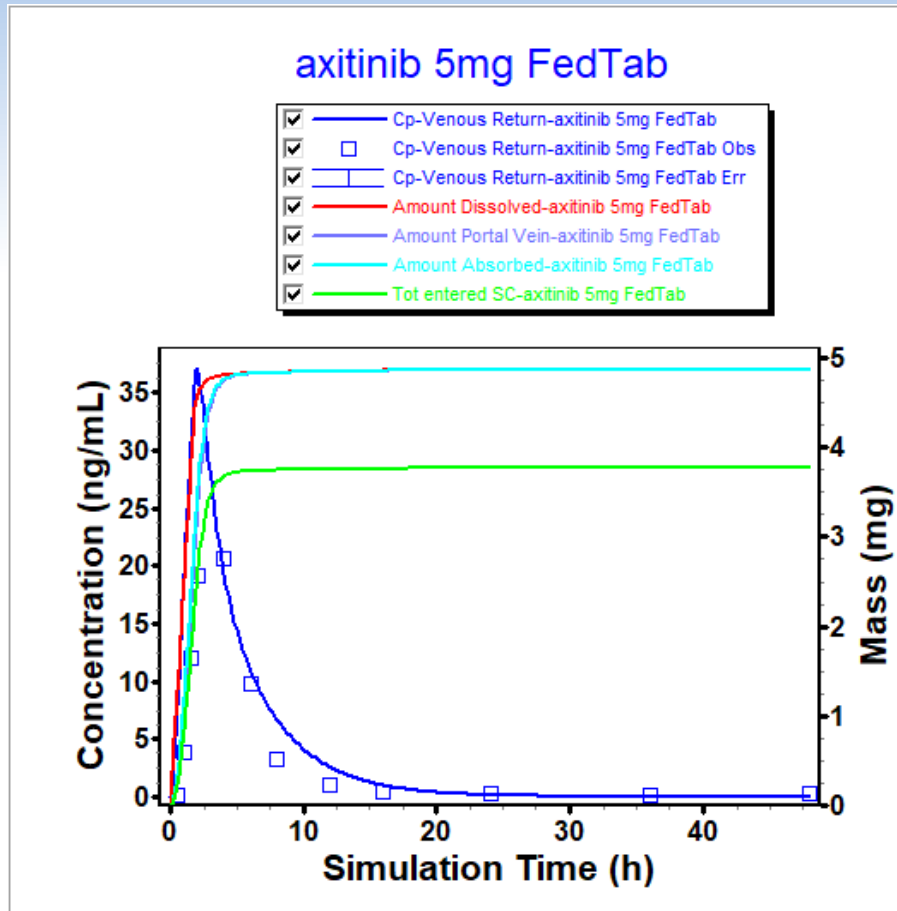
Gastric emptying is too fast using default values

# Case Study: Axitinib Fed State

## Zero Order Emptying

Normal Meal (500 cal)  
Gastric Emptying = 1.84 hr

Fed High-Fat (900 cal) Bile Salt Increased  
Gastric Emptying = 3.11 hr

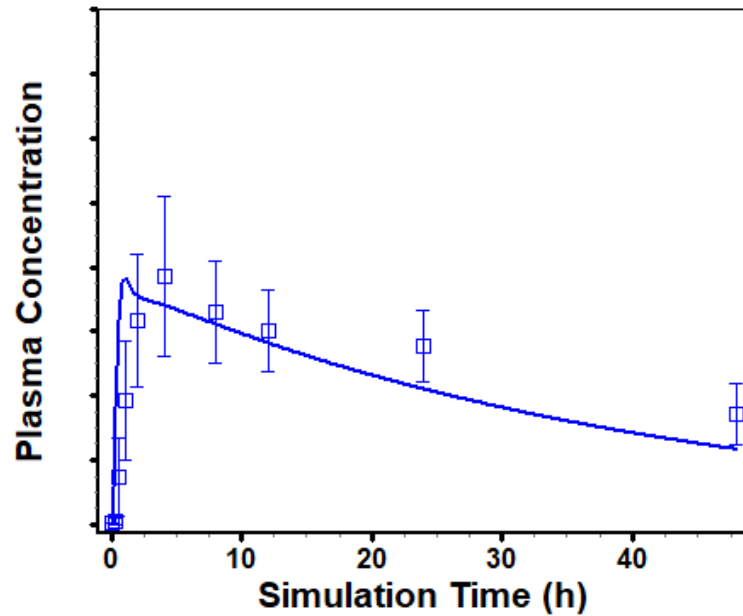


# Case Study: High Fat Meal Prediction

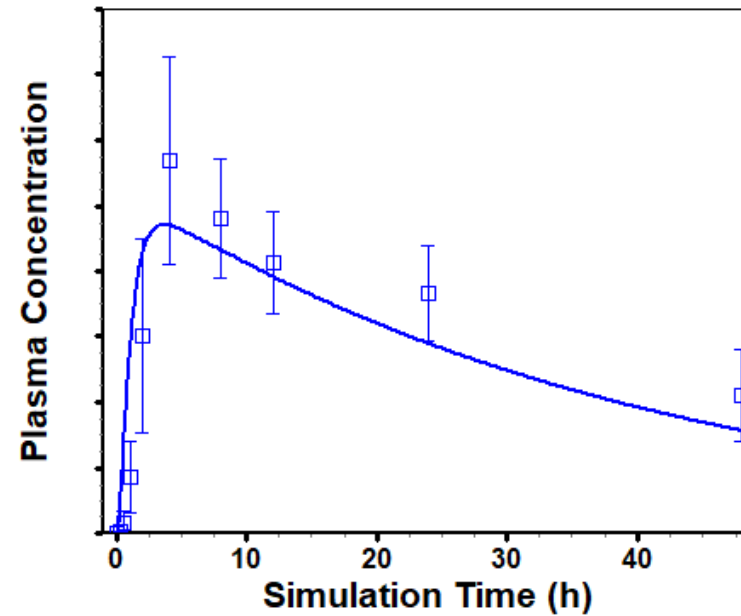
Default Fed in GP9.6

Default for FDA standard breakfast

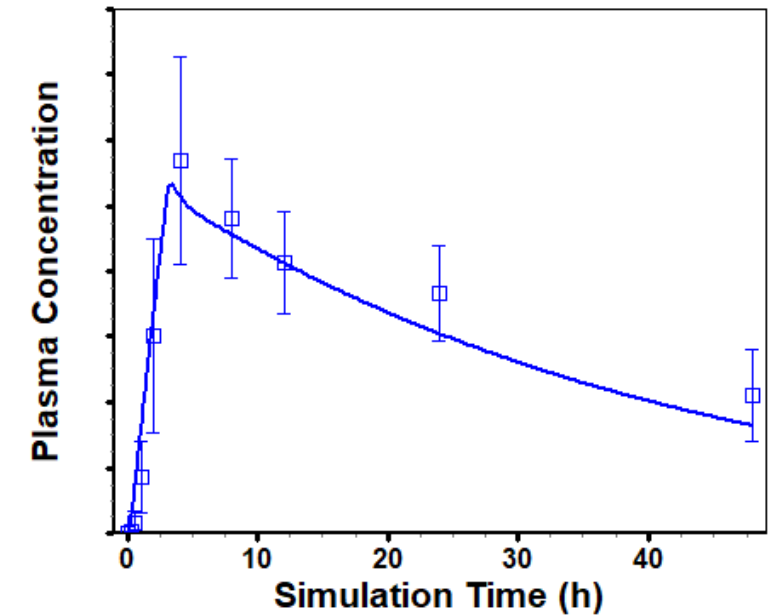
Fasted



Fed



Fed





# New Fed State Meal Option Validation Summary

	Meal Type	Best Fed Model Setting
Dolutegravir	High Fat / High Calorie	Zero Order GP 9.7
	Mod. Fat / Mod. Calorie	Zero Order GP 9.7
	Low Fat / Low Calorie	Zero Order GP 9.7
Lapatinib	Low Fat / Low Calorie	Default Gastroplus 9.6
	High Fat / High Calorie	Default Gastroplus 9.6
Ixazomib	High Fat / High Calorie	Exponential GP 9.7
Axitinib	Normal Meal	Exponential GP 9.7
	High Fat / High Calorie	Zero Order GP 9.7
Internal Study 1	Low Fat / Low Calorie	Zero Order GP 9.7
	High Fat / High Calorie	Zero Order GP 9.7
Internal Study 2	FDA High Fat Breakfast	Zero Order GP 9.7
Internal Study 3	High Fat / FDA Breakfast	Zero Order GP 9.7
Internal Study 4	High Fat / FDA Breakfast	Zero Order GP 9.7

- Validation studies indicate zero-order emptying is most predictive of plasma concentration in fed-state

# Q&A

Questions & Answers



**GastroPlus<sup>®</sup>**



# GastroPlus<sup>®</sup> Demo – Compound X Positive Food Effect

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*Research Article*

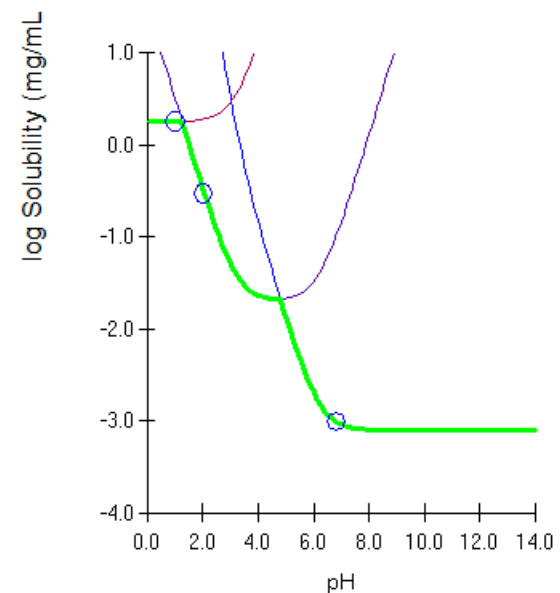
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## **Application of Physiologically Based Absorption Modeling to Formulation Development of a Low Solubility, Low Permeability Weak Base: Mechanistic Investigation of Food Effect**

Hefei Zhang,<sup>1,2</sup> Binfeng Xia,<sup>1</sup> Jennifer Sheng,<sup>1</sup> Tycho Heimbach,<sup>1</sup> Tsu-Han Lin,<sup>1</sup> Handan He,<sup>1</sup> Yanfeng Wang,<sup>1</sup> Steven Novick,<sup>1</sup> and Ann Comfort<sup>1</sup>

*Received 2 October 2013; accepted 23 December 2013*

- **Compound X (BCS Class II/IV)**
- Lipophilic ( $\log P > 4$ ) and moderate base (pKa 3.2 and 6.2)
- Low (0.001 mg/mL), pH dependent aqueous solubility
- Moderate intestinal permeability ( $1.48 \times 10^{-4}$  cm/s)
- Class II (or IV)
- 70% dose recovered in feces as parent
- Estimated bioavailability of compound is ~30%

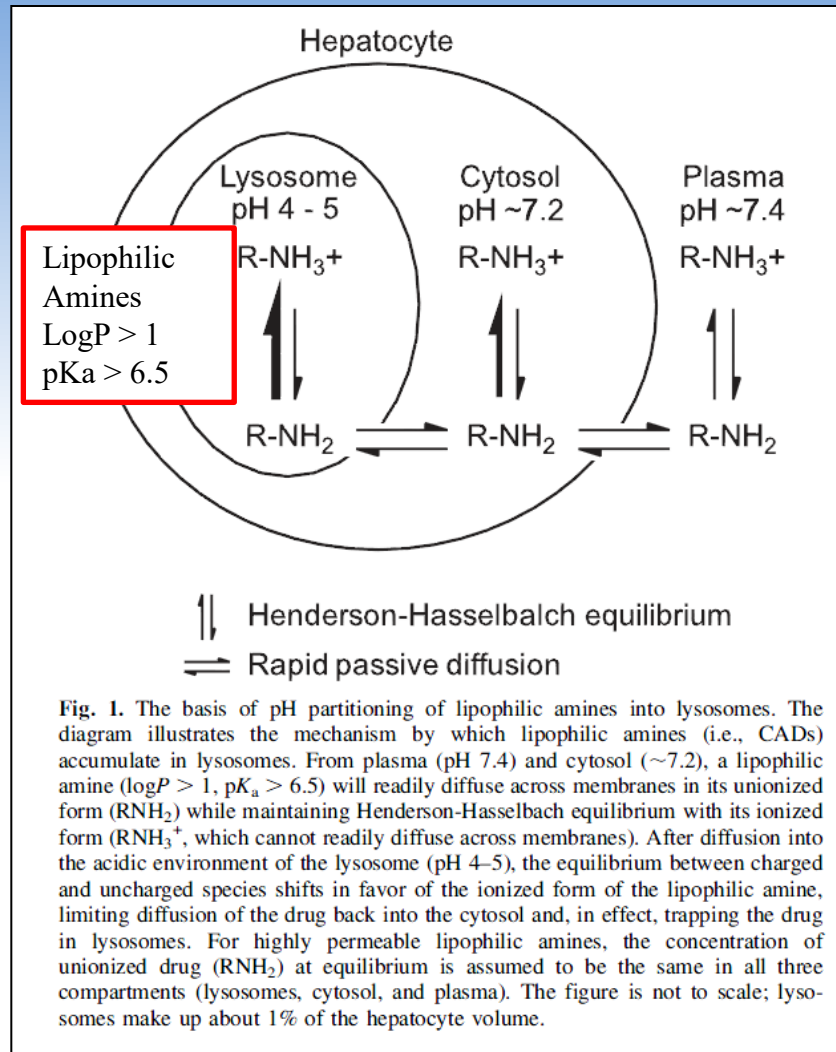


# Physicochemical and pharmacokinetic parameters for Compound X

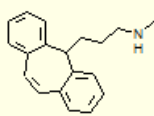
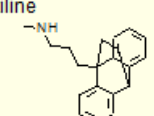
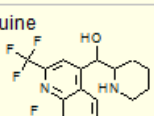
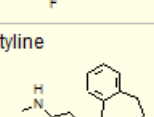
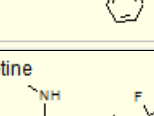
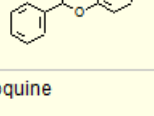
**Table II.** Physicochemical Parameters, Default Physiological Values, and Pharmacokinetic Parameter Used in the Simulation at Various Doses

Parameters	Value(s)
<b>Compound parameters</b>	
$M_w$ : 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):	$1.48 \times 10^{-4}$
Particle radius of API ( $\mu\text{m}$ ):	19
<b>Physiological parameters</b>	
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
<b>Pharmacokinetics</b>	
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

# Lysosomal Trapping of Lipophilic Cations



Ref: Kazmi F., Drug Metab. Disp. 41(3):897 (2013)

Drug	Log P	Basic pKa	T <sub>max</sub> (h)
Protriptyline 	4.69	10.0	27
Maprotiline 	4.7	10.1	16
Mefloquine 	3.81	8.52	15
Nortriptyline 	4.46	9.65	7.8
Fluoxetine 	4.39	9.82	7
Chloroquine 	5.11	9.86	6

# Activity: Compound X

1. **Open** the “**Compound X**” database in your **Examples\Compound X** folder
2. Navigate to the **Hum 200 mg IR Cap – Fasted** record and review the model parameters. *What is unique about this compound?*
3. Let’s run a PSA around the fraction unbound in enterocytes. **Click** on the “**PSA**” button on the **simulation tab** and select “**FuEnt**” on the **pharmacokinetics tab**
4. Need to change **Fu,ent** to ~3% by going to the **Database**→**4 View/Edit Tables**→**1 View/Edit Drugs** Table menu
5. Let’s modify the **mean precipitation time** in the model to match better with the observed PK data
6. Navigate to the **Hum 200 mg IR Cap – Fed** record and enter this new **Mean Precipitation Time**

# 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

ACAI      ΔCAT-Compound      Compound      Formulation

Dosing Hum: 200 mg I ~ap - Fasted      Manufacture: Hum 200 mg I ~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 RetT

Parameter Sensitivity Analysis Setup

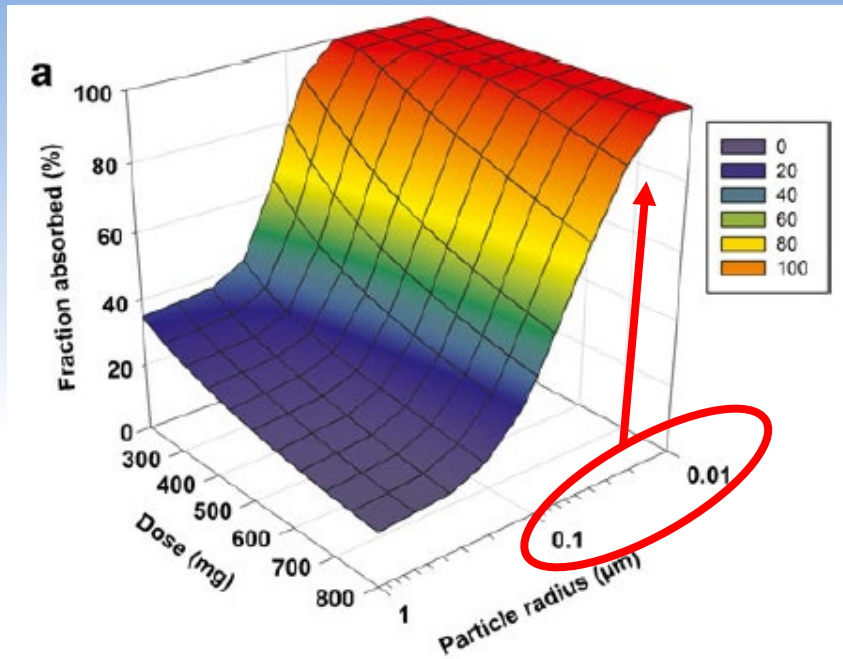
Select Parameters	Parameter	Lower Bound	Baseline Value	Upper Bound	Number of Test	Spacing of Param Values
	Dose of Hum 200 mg IR Cap - Fast	50	200	1000	5	Logarithmic
	Mean Drug Particle Radius of Hum	0.5	19	50	5	Logarithmic

Run 3D PSA

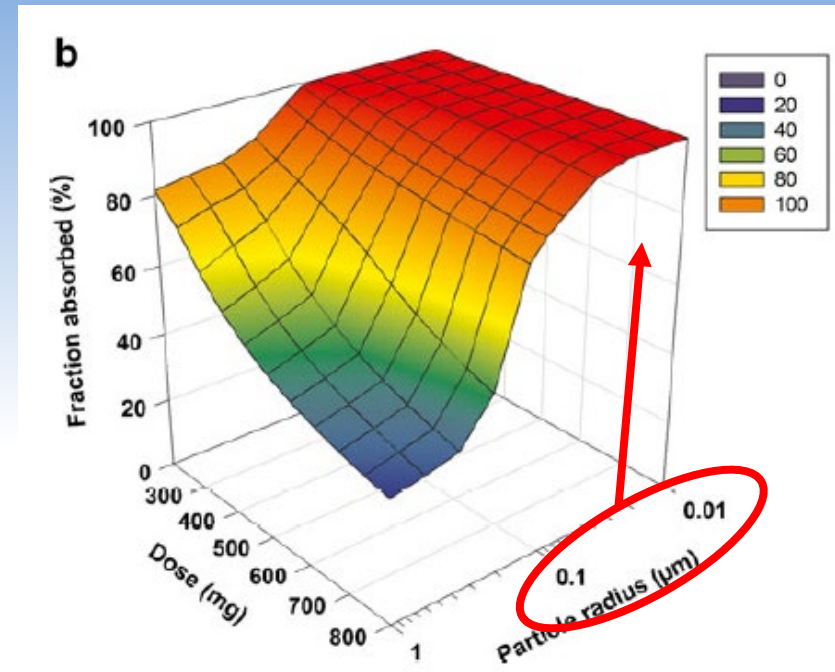
Zhang et al. AAPS PharmSciTech 2014 January 17



# 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

Zhang et al. AAPS PharmSciTech 2014 January 17

# Negative food effects

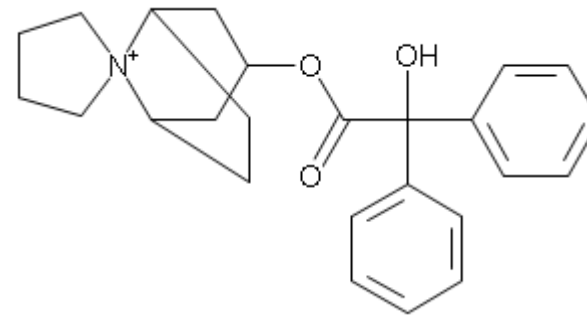
## Mechanistic investigation of food effect on disintegration and dissolution of BCS class III compound solid formulations: the importance of viscosity

Asma Radwan<sup>a</sup>, Gordon L. Amidon<sup>b</sup>, and Peter Langguth<sup>a,\*</sup>

<sup>a</sup>*Institute of Pharmacy and Biochemistry, Johannes Gutenberg University, Mainz, Germany*

<sup>b</sup>*College of Pharmacy, The University of Michigan, Ann Arbor, MI 48109-1065, USA*

- **Trospium HCl (BCS Class III)**
- Hydrophilic (log P = -1.22)
- High (~700 mg/mL)
- Low intestinal permeability ( $0.07 \times 10^{-4}$  cm/s)
- Class III
- Not metabolized
- Estimated bioavailability of compound is ~10%



# Trospium Solution Viscosity

Table 2. Physicochemical characteristics of trospium chloride solutions in different media. Except for solubility, determinations were performed once. For solubility, the means of  $n = 2-3$  and the individual values or the standard deviations are reported

Media	Viscosity $\eta$ (cP) at 1.29 s <sup>-1</sup>	Density (g/ml)	Solubility (g/ml)	Osmotic pressure (mmol/kg)
SIF	1	0.997	0.786 ± 0.021	95
0.1 N HCl	1	0.991	0.696 (0.703, 0.689)	188
0.5% HPMC (pH = 6.8)	20	0.993	0.667 ± 0.029	115
1% HPMC (pH = 6.8)	100	0.998	0.537 (0.540, 0.533)	125
2% HPMC (pH = 6.8)	1300	1.004	0.415 (0.420, 0.410)	200
Acetate buffer	1	0.990	0.765 (0.759, 0.770)	74
0.5% HPMC (pH = 4.6)	ND	0.991	0.658 (0.624, 0.693)	82
1% HPMC (pH = 4.6)	ND	0.994	0.559 (0.568, 0.550)	84
2% HPMC (pH = 4.6)	900	0.997	ND	86
0.25% guar (pH = 6.8)	15	0.998	0.787 (0.770, 0.803)	105
0.5% guar (pH = 6.8)	140	0.998	0.636 (0.643, 0.628)	106
0.75% guar (pH = 6.8)	1350	0.998	0.515 (0.520, 0.510)	109

ND, not determined.

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*Biopharm. Drug Dispos.* 33: 403–416 (2012)

DOI: 10.1002/bdd

# Trospium Solution Viscosity

Table 2. Physicochemical parameters were performed once.

Media

- SIF
- 0.1 N HCl
- 0.5% HPMC (pH = 6.8)
- 1% HPMC (pH = 6.8)
- 2% HPMC (pH = 6.8)
- Acetate buffer
- 0.5% HPMC (pH = 4.6)
- 1% HPMC (pH = 4.6)
- 2% HPMC (pH = 4.6)
- 0.25% guar (pH = 6.8)
- 0.5% guar (pH = 6.8)
- 0.75% guar (pH = 6.8)

ND, not determined.

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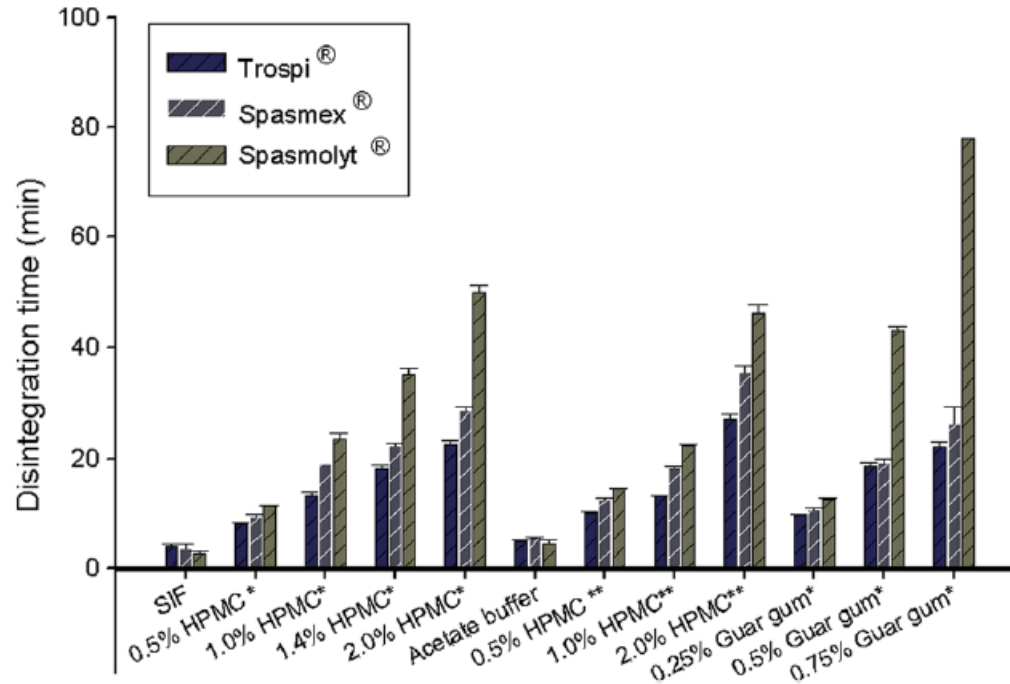


Figure 3. Disintegration times of various trospium chloride products in different disintegration media. The effects of increasing media viscosity on disintegration times were in all cases significant ( $p < 0.05$ ), whereas the effect of change of pH for HPMC solutions at the same concentrations of VEA was insignificant ( $p > 0.05$ ). \*pH 6.8; \*\*pH 4.6

lity, determinations  
ons are reported

: pressure (mmol/kg)

- 95
- 188
- 115
- 125
- 200
- 74
- 82
- 84
- 86
- 105
- 106
- 109

os. 33: 403–416 (2012)  
DOI: 10.1002/bdd

# Trospium *in vitro* dissolution

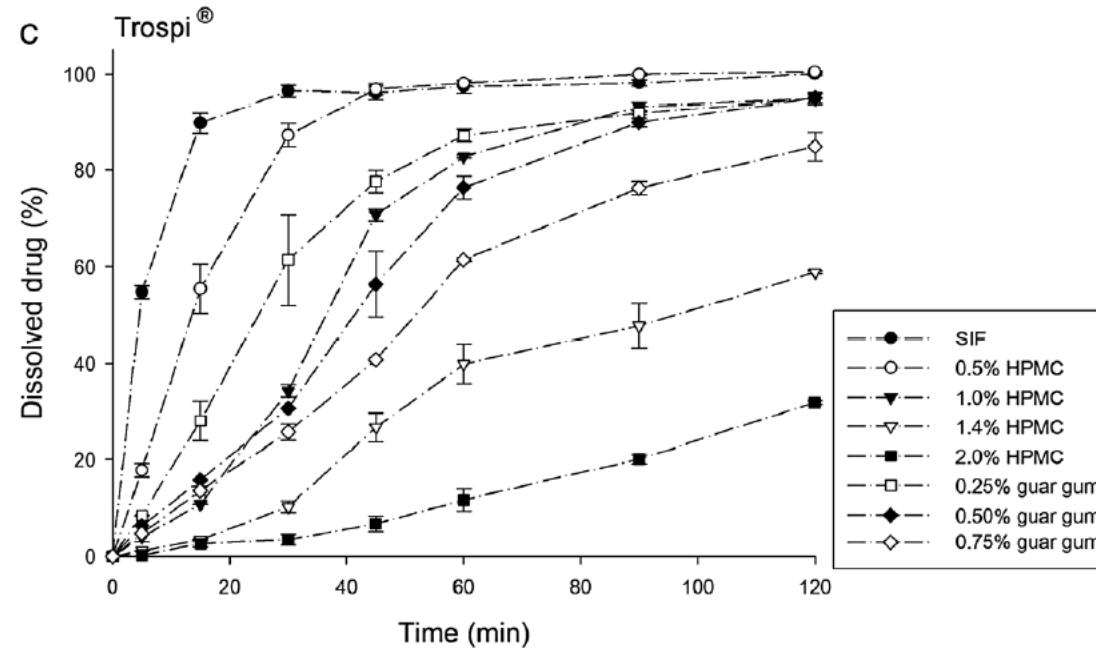


Figure 5. (a) Dissolution profiles for Spasmolyt<sup>®</sup> in viscous HPMC and guar solutions at pH 6.8, 50 rpm in USP-2 apparatus. In all cases values for  $f_2$  were  $< 50$ . Mean  $\pm$  SD,  $n = 3$ . (b) Dissolution profiles for Spasmex<sup>®</sup> in viscous HPMC and guar solutions at pH 6.8, 50 rpm in USP-2 apparatus. In all cases values for  $f_2$  were  $< 50$ . Mean  $\pm$  SD,  $n = 3$ . (c) Dissolution profiles for Trospi<sup>®</sup> in viscous HPMC and guar solution at pH 6.8, 50 rpm in USP-2 apparatus. In all cases values for  $f_2$  were  $< 50$ . Mean  $\pm$  SD,  $n = 3$

# Activity: Trospium

1. **Open** the “Trospium” database in your **Examples\Trospium** folder
2. Navigate to the **Human 60 mg Tablet Fasted** record and review the model parameters. **Run PSA** around select parameters to identify the key properties impacting absorption
3. Transfer the adjusted Peff to the **Human 60 mg Tab 0.5 pct HPMC – Fasted** record. With the *in vitro* dissolution data, use the **Z-Factor** model to **simulate** the PK
4. Transfer the adjusted Peff to the **Human 60 mg Tab 2 pct HPMC – Fed** record and **switch** the gut physiology to the **Fed model**. With the *in vitro* dissolution data, use the **Z-Factor** model to **simulate PK**
5. Investigate possible reasons for the slight overprediction of the PK under fed conditions

# Negative Food Effect

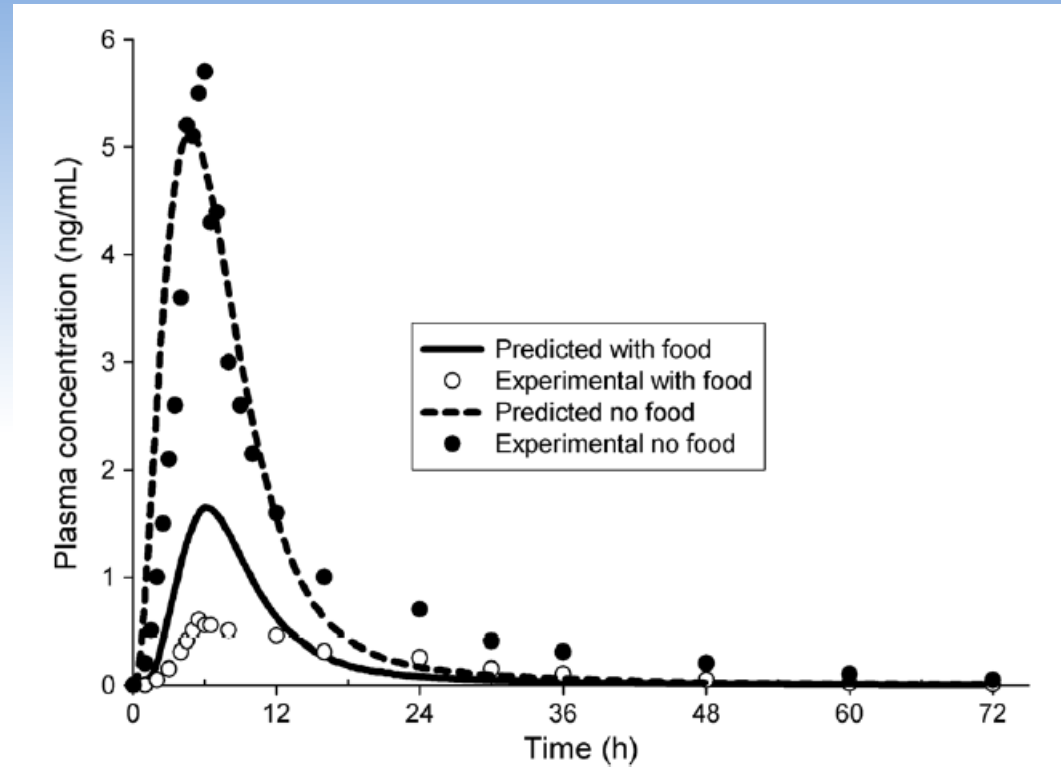
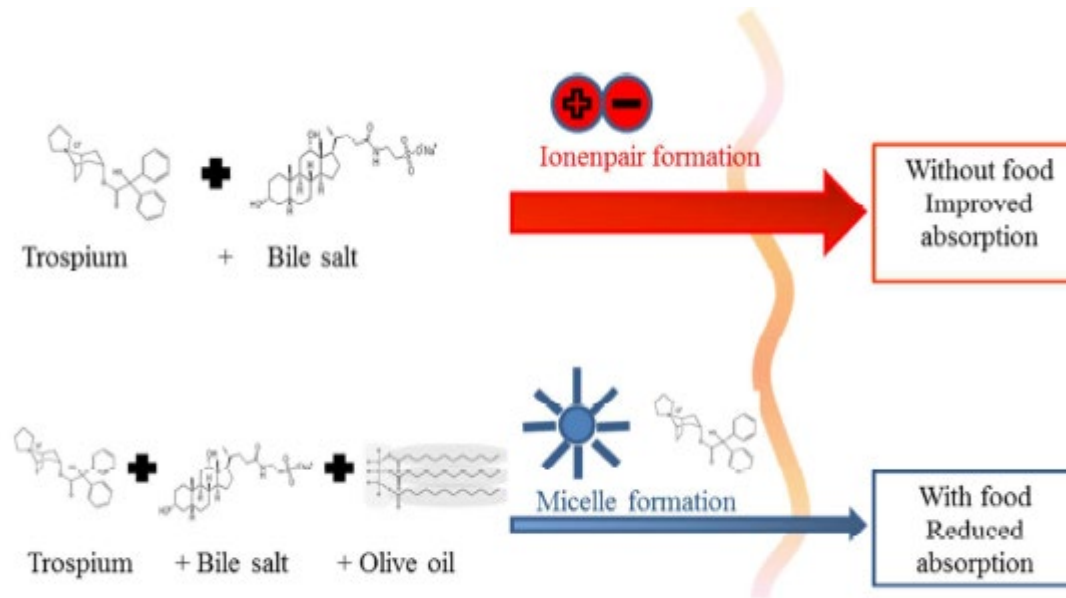


Figure 8. Simulated and predicted plasma concentration–time profiles for trospium in fasted and fed states in humans



Ion pairing with bile salts modulates intestinal permeability and contributes to food-drug interaction of BCS class III compound trospium chloride



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# Novartis Negative Food Effect: Caco-2 Experiment in FeSSIF Buffer

Contents lists available at ScienceDirect  
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Journal homepage: www.elsevier.com/locate/ejpb

Research paper  
Biorelevant media for transport experiments in the Caco-2 model to evaluate drug absorption in the fasted and the fed state and their usefulness  
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ABSTRACT  
In this work we developed and characterized transport media that simulate the composition of intestinal phase of stomach fluid in the fasted and fed state and are appropriate for evaluating intestinal drug permeability characteristics using the Caco-2 model (FeSSIF-10 $\mu$ M and FaSSIF-10 $\mu$ M, respectively). Media composition was based on factors (pH and bicarbonate) and actively reported data on total lipid concentrations in the mucosal phase of contents of the upper small intestine in the fasted and the fed state and was adjusted for oil culture compatibility. Permeation data were evaluated by compartmental kinetic modeling. Permeability coefficients of 12 hydrophilic drugs were not affected by media composition. In contrast, P values of a series of lipophilic compounds measured with FeSSIF-10 $\mu$ M and FaSSIF-10 $\mu$ M and relating to transport by diffusion were smaller than those obtained with a purely aqueous reference transport medium, as 10 $\mu$ M, following the rank order  $P_{FaSSIF-10\mu M} < P_{FeSSIF-10\mu M} < P_{10\mu M}$ . The degree of permeability values was stronger as lipophilicity of the compounds increased. Compared with values obtained using as 10 $\mu$ M, permeability were reduced, depending on the compound, by more than 20- to 100 fold when measured with FeSSIF-10 $\mu$ M, whereas compound ranking in regard to the permeability characteristics was also affected. The impact of advanced Factor on Bio through the in vitro, hence on drug absorption, is corroborated with the drug amount found in cellular particles needs to be taken into consideration in PBPK modeling especially when the food effect is evaluated.

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35 | Heimbach, Lin, G+ UaeGroup, 2015

### Compound U: ACAT model did not predict the negative food effect using the regular Permeability

Caco-2 Permeability	Low to moderate
Papp A-B and B-A, cmx10 <sup>-6</sup> /min	2.5, 12.8
Peff, cmx10 <sup>-4</sup> /s	1.445

Compound P displayed negative food effect in humans  
ACAT model could not predict the negative food effect using the original Papp data

37 | Heimbach, Lin, G+ UaeGroup, 2015

### Effect of FeSSIF components on Drug U permeability

Drug U: PAB (cmx10<sup>-6</sup>/min) is ~5-8X higher in FeSSIF/FaSSIF (10  $\mu$ M) compared to FaSSIF and FeSSIF.

Propranolol: PAB (cmx10<sup>-6</sup>/min) is ~2X higher in FaSSIF and FeSSIF compared to FeSSIF/FaSSIF (10  $\mu$ M).

- Absorptive permeability increased with increasing Drug U concentration possibly due P-gp saturation or excessive non-specific binding
- FaSSIF/FeSSIF absorptive permeability ratio (10  $\mu$ M) was ~5-8 for Drug U and approximately 2.0 for propranolol

38 | Heimbach, Lin, G+ UaeGroup, 2015

### GastroPlus PBPK Modeling with 8 fold lower Peff

	Fasted		Fed		Fed/Fasted	
	Sim	Obs	Sim	Obs	Sim	Obs
Cmax (ng/mL)	359	343	227	160	0.63	0.47
AUC (ng/mL.h)	9788	9737	7944	4406	0.46	0.45

Model	P <sub>app</sub> × 10 <sup>4</sup> cm/min ± SD		P <sub>app, FaSSIF/FaSSIF</sub> fold difference
	FeSSIF <sup>a</sup>	FeSSIF <sup>b</sup>	
HB&S	2.91 ± 0.06	1.10 ± 0.12	0.132 ± 0.007

- Early ACAT model could not predict the negative food effect using the original Papp/Peff data
- Ratio of biorelevant Papp in the FeSSIF/FaSSIF was applied to simulate the PK under fed condition
- Update ACAT model can 1) well simulate the PK profile under fasted condition; 2) correctly predict the negative food effect of Drug U in humans

39 | Heimbach, Lin, G+ UaeGroup, 2015

# Q&A

Questions & Answers



**GastroPlus<sup>®</sup>**