



Best Practices for Membrane & Biphasic *In Vitro* Dissolution with DDDPlus™ & GastroPlus®

Jim Mullin (jim@simulations-plus.com)

What is DDDPlus™?

Reimagining how companies design and analyze *in vitro* dissolution studies

- Now utilized by >50 companies globally plus the US and China FDAs

Provides models for most dosage forms and experimental conditions

- Immediate/delayed/controlled release oral products plus long-acting injectable formulations
- USP, ASD, biphasic, membrane dissolution apparatus

Significant momentum behind the DDDPlus/GastroPlus marriage to capture IVIVE of precipitation kinetics and establish product specs



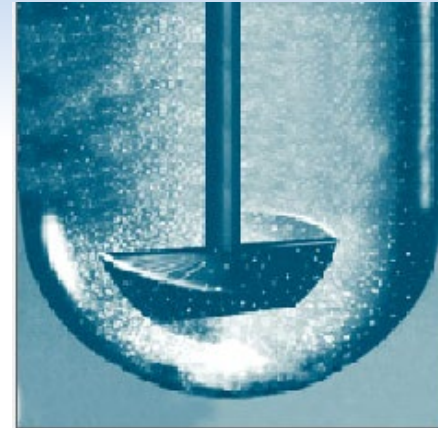
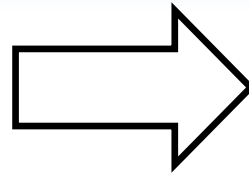
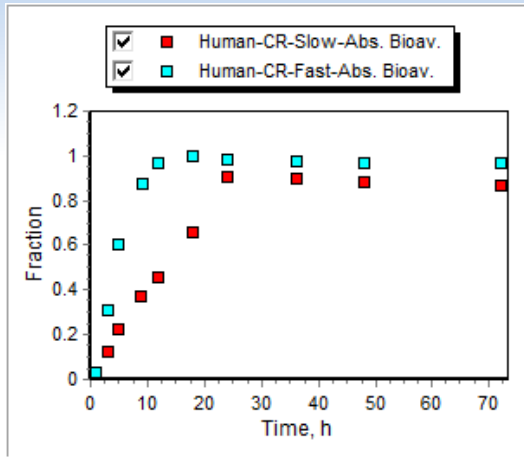
Introduction

- What is DDDPlus
 - How can it be utilized in drug development?
 - What can DDDPlus predict?
 - What inputs are necessary?
 - What results can be obtained?
- Case Study – synergy of DDDPlus and GastroPlus
 - Biphasic and membrane dissolution experiments for determination of precipitation kinetics.
 - *In Vitro to In Vivo* (IVIVE) extrapolation of precipitation

How Can Simulation Software be Used?

Dissolution Method Development

- How can I design my *in vitro* experiment to correlate with the *in vivo* release?



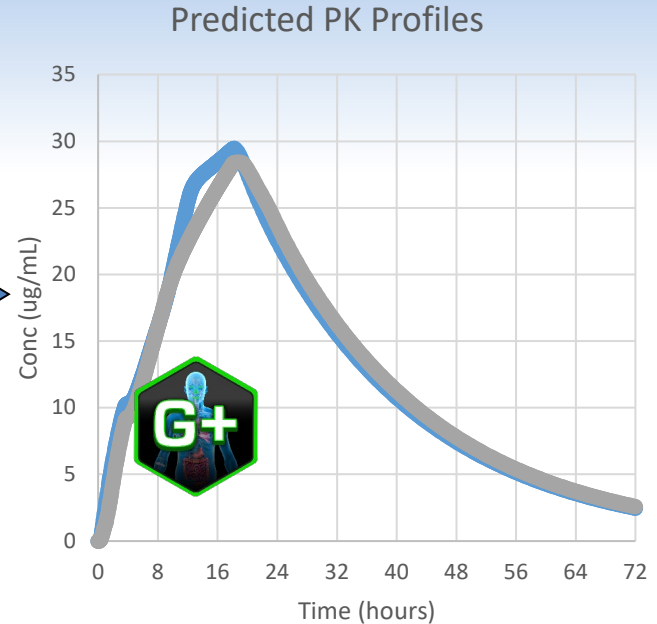
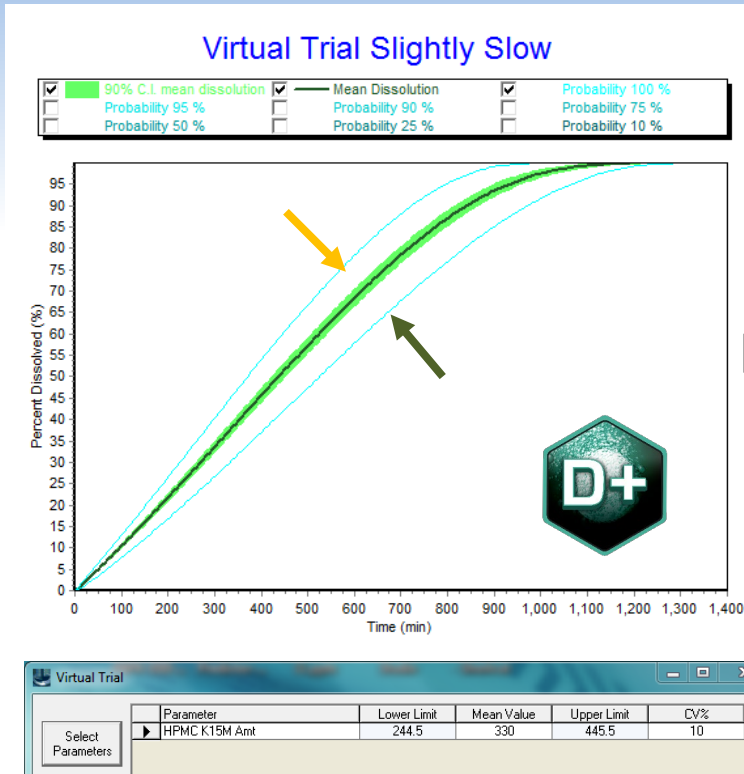
Formulation Design

- How do I modify my formulation to produce an *in vitro* dissolution rate that meets my target?

Establish design or safe spaces

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

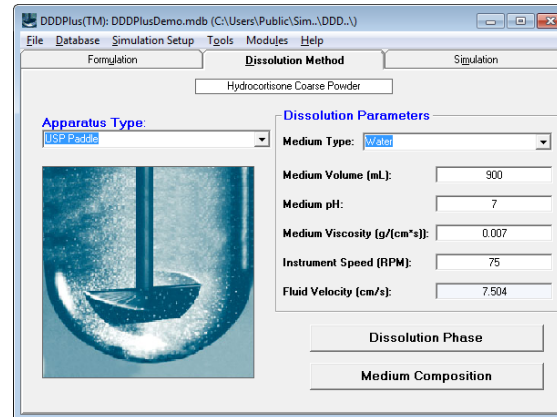
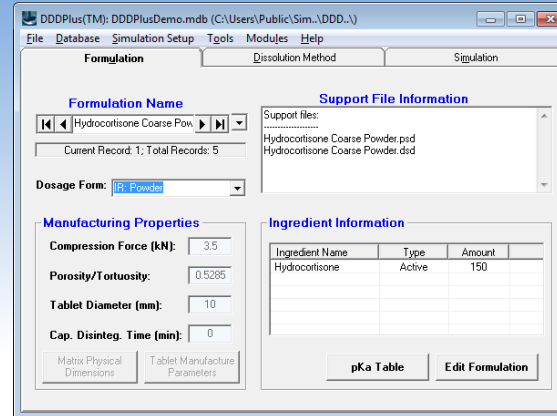
100th percentile ('extreme') dissolution profiles
loaded into GastroPlus to predict PK



What does DDDPlus consider?

DDDPlus is a state-of-the-art formulation simulation computer program that contains equations to account for the following:

- Dissolution rate for active pharmaceutical ingredient (API) and excipients
- Multiple particle size distribution for API and excipients
- Variety of dosage form models
- Solubility-dynamic microclimate pH calculation for API and excipients
- pH of buffers from composition of acids, bases, and salt equivalents.
- Selection of USP and biorelevant experimental apparatus and conditions



DDDPlus User Inputs

Formulation:

Compression force

Tablet diameter

Disintegration time

Porosity

Matrix dimensions (for controlled-release)

Ingredient(s):

Name

Salt type (if any)

Diffusion coefficient

Solubility factor

Particle size distribution

Type (API, wetting agent, disintegrant, other)

Mol. Weight

pKa(s)

Solubility @ **reference pH**

(type, mean, standard deviation, # bins)

Dissolution Method:

Vessel Type

Medium Volume

Buffer

Different experimental phases

Speed

Initial pH

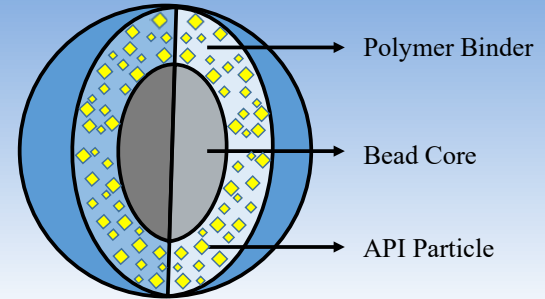
Surfactants

IMPORTANT: applying measured *in vitro* dissolution data to optimize model parameters is a key step in developing a validated model!

Formulation Models

Immediate Release (*IR*)

- Powder
- Solution (for precipitation studies)
- Tablet
- Capsule
- Bead Coating

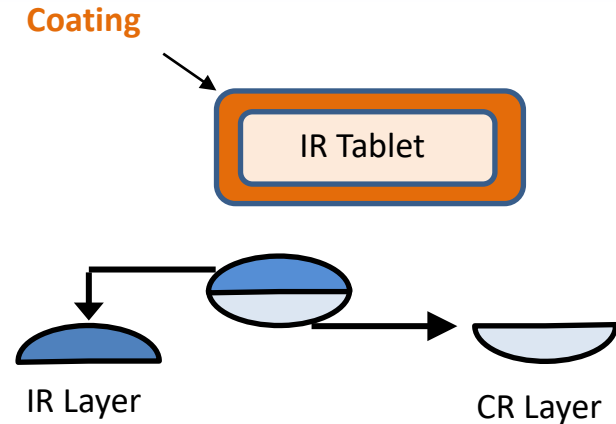


Delayed Release (*DR*)

- Coated Tablet

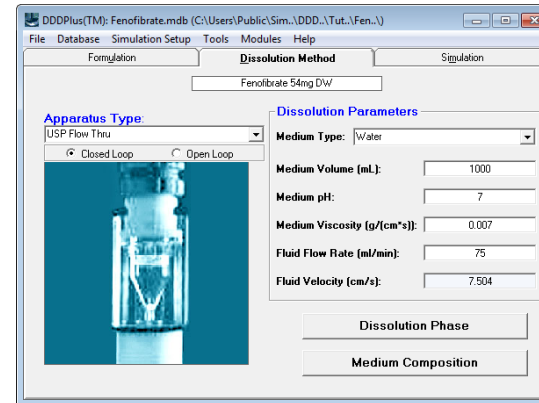
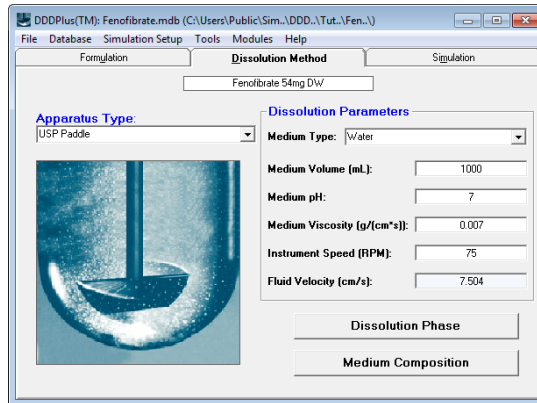
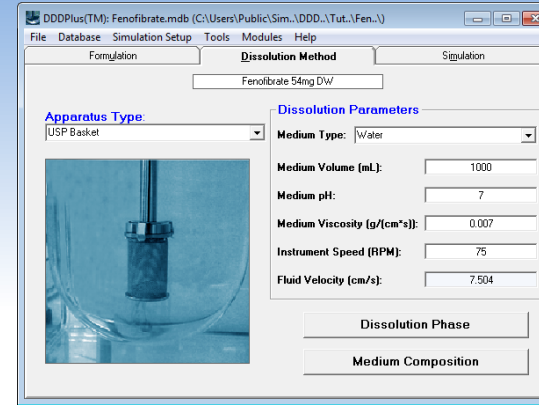
Controlled Release (*CR*)

- Polymer Matrix (Swellable)
- Polymer Matrix (Non-Swellable)
- Bilayer Tablet
- Bead Coating
- Long-Acting Injectable (LAI) Microsphere



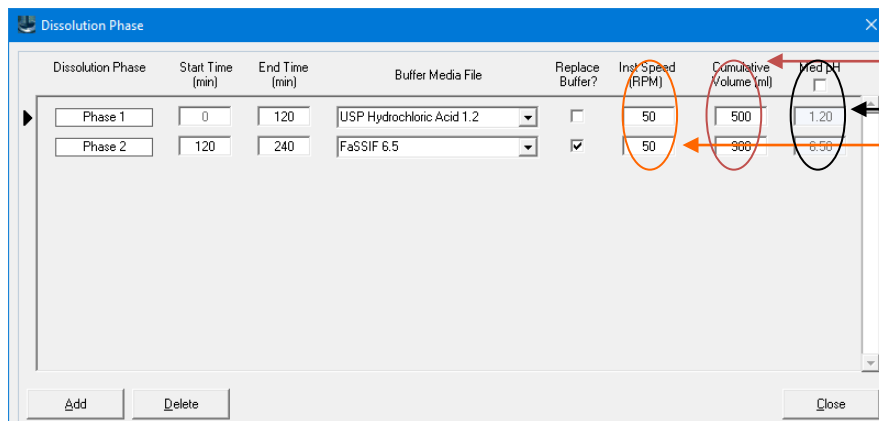
Experimental Apparatus Options

- USP 1 Basket
- USP 2 Paddle
- USP 4 Flow-through
 - ✓ Open or closed loop
- Pion μ Diss Profiler™
- Artificial Stomach Duodenum (ASD)
- Membrane Dissolution
- Biphasic Dissolution



Improved Multi-stage Dissolution Experiments

- Redesigning *in vitro* experiments can help when an *in vitro-in vivo* correlation cannot be achieved
 - *in vivo* environment sometimes cannot be accurately reproduced using standard *in vitro* conditions
- DDDPlus lets you optimize *in vitro* experimental conditions to produce a desired dissolution profile by incorporating multiple “phases”
 - Instrument speed, medium volume, and pH can vary for each “phase” you define, as well as the start and end times for each phase
 - **New!** Fully mechanistic pH buffer and microclimate pH calculations based on buffer compositions and if replacement or mixing of buffers is occurring



By adjusting:

Medium volume

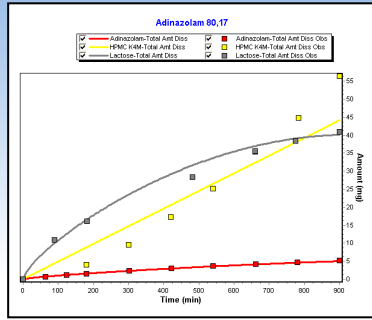
pH

Instrument Speed

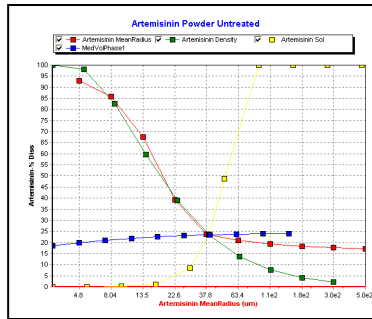
Full pH calculations of all the buffer mixtures are accounted for or you can replace the media with fresh buffer.

DDDPlus™ Outputs

Dissolution profiles for all formulation ingredients

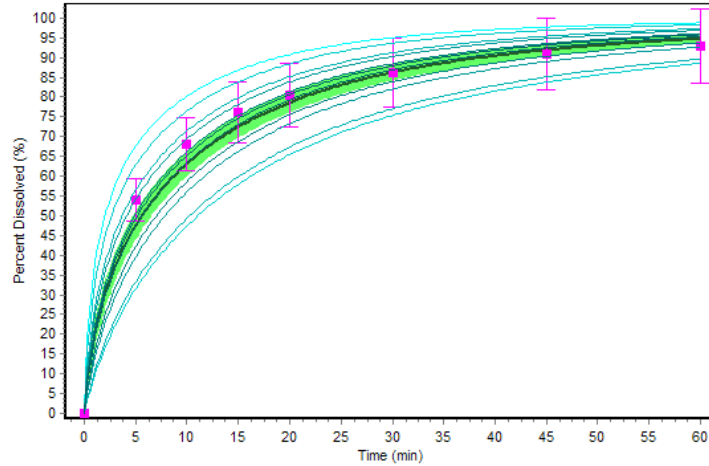
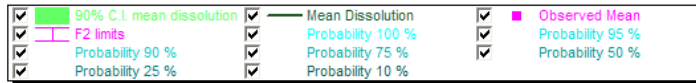


Parameter Sensitivity Analysis “Spider” Plots

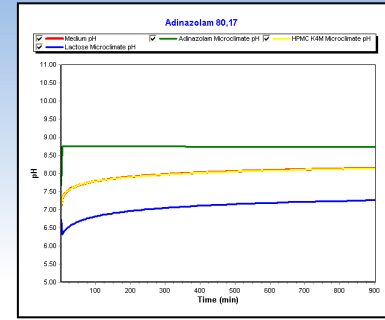


Virtual Trials

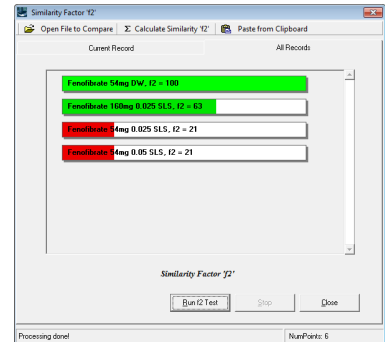
Virtual Trial Hydrocortisone Coarse Powder



Medium/microclimate pH changes



f1 (difference) & f2 (similarity) calculations



Case Study – Synergy of DDDPlus and GastroPlus

- Leverage DDDPlus to determine dissolution, precipitation kinetics, and formulation solubility for:
 - Compounds:
 - Dipyridamole
 - Ketoconazole
 - Itraconazole
 - Theory of membrane dissolution, biphasic dissolution, and precipitation models
 - DDDPlus precipitation model results for biphasic and membrane dissolution
- Utilize GastroPlus to predict *In Vivo* PK with DDDPlus precipitation kinetics from DDDPlus

Membrane Dissolution Experiment

- Membrane dissolution is a useful tool for:
 - Assessing formulation solubility for amorphous or enabled formulations vs. crystalline drug
 - Fitting precipitation kinetics for weakly basic compounds with simultaneous absorption.
- Input Parameters:
 - Donor and receiver diffusion layer thickness
 - Partition coefficient into membrane lipid phase
 - Solubility in receiver solution (assumes sink – proprietary buffer).
 - System parameters – membrane area, thickness, and porosity.
- Output Results:
 - Formulation solubility determined from drug absorption rate
 - Precipitation kinetics

DDDPlus Membrane Dissolution Options

Membrane Flux Parameters

Turn on Membrane Flux Dissolution

Receiver Volume (mL): 20.00

Receiver Viscosity (g/(cm*s)): 0.01000

Receiver Diffusion Layer Thickness (um): 10.00

Donor Diffusion Layer Thickness (um): 10.00

Membrane Thickness (um): 125.00

Membrane Area (cm²): 1.54

Membrane Porosity (fraction): 0.760

Membrane Partition Coefficient Conversion

LogP_{mem:w} = 0.660 × LogP_{o:w} + 1.170

Cancel Close

Formulation Composition

File Structure Import Tools

Selection and Preview Grid

Ingredient Name	Ingredient Type	Amount (mg)	MW/t (g/mol)	Diff. Coeff. (cm ² /s)	De
Hydrocortisone	Active	150	362.47	0.587	1.2

Physicochemical Information

Ingredient Name: Hydrocortisone | Ingredient Type: Active | Mol. Weight (g/mol): 362.47

Ref. Solubility (mg/ml): 0.361 | pH for Ref. Solubility: 7.2

Density (g/ml): 1.27 | Diff. Coeff. (cm²/s¹⁵): 0.587

logP: 1.58 | @ pH: 1 | Receiver Sol. (mg/ml): 100

Membrane Sol. (mg/ml): 100.97

Diff. Coeff. Mem. (cm²/s^{10^5}): 0.587 | Diff. Coeff. Rec. (cm²/s^{10^5}): 0.587

Biorelevant Solubility | Surfactant Solubility | Precipitation Model

Formulation Specific Information

Amount (mg): 150 | Salt Type: None | No. Moles: 1

Particle Size Distribution...

Mean Radius (um)	Stan. Dev.	Num. of Bins
17.94	2.4	16

Distribution Type: Log Normal (Geometric Inputs) | Rmin: 1.298 | Rmax: 248.003

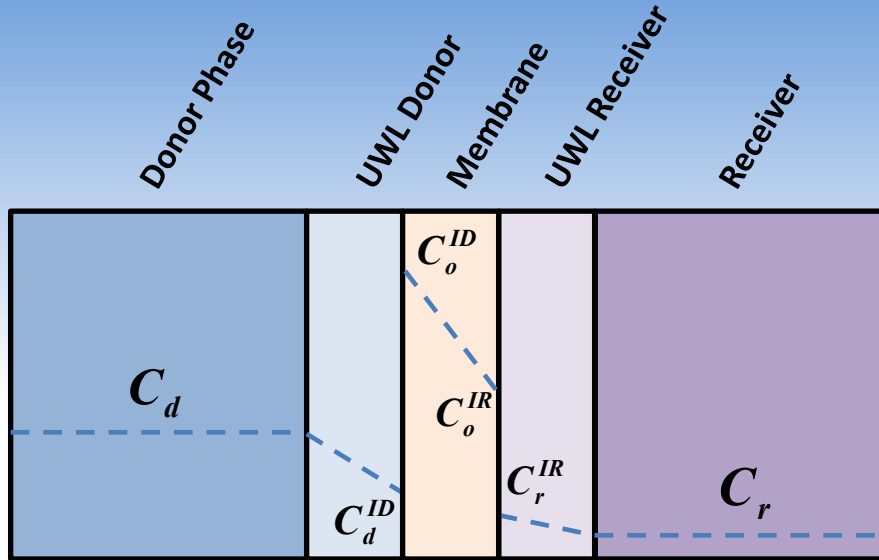
Constants

Calibration Constant: 1

Δ Add Ingredient | Delete Ingredient | Import Excipients | Close

Dissolution Model: Johnson-Spherical

Transmembrane Flux for Membrane Assays



$$j_{overall} = \frac{D_A D_o D_R K_{od}}{D_A D_R h_o + D_o (D_R h_d K_{od} + D_A h_r K_{or})} (C_d - K_r C_r)$$

Where: $K_r = \frac{K_{or}}{K_{od}}$

Steady-State Flux Equations:

$$j_{d,uwl} = \frac{D_A}{h_d} (C_d - C_d^{ID}) \quad j_o = \frac{D_o}{h_o} (C_o^{ID} - C_o^{IR})$$

$$j_{r,uwl} = \frac{D_R}{h_r} (C_r^{IR} - C_r) \quad j_{r,uwl} = j_o = j_{d,uwl}$$

Where: $C_d^{ID} K_{od} = C_o^{ID}$ and $C_r^{IR} K_{or} = C_o^{IR}$

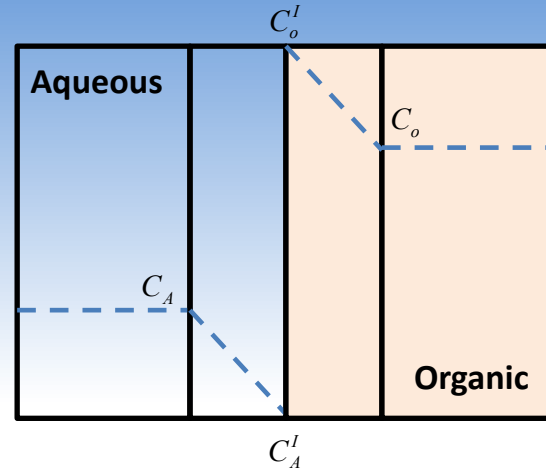
- C_d Concentration in donor compartment
- C_d^{ID} Concentration at donor/membrane interface
- C_o^{ID} Concentration in organic/donor interface
- C_o^{IR} Concentration in organic/receiver interface
- C_r^{IR} Concentration at receiver/membrane interface
- C_r Concentration in receiver compartment
- D_o, D_A Diffusion coefficient
- h_d, h_o, h_r Diffusion layer thickness in donor/organic/receiver
- $K_{od} = \frac{C_o^{eq}}{C_d^{eq}}$ Organic to donor partition coefficient
- $K_{or} = \frac{C_o^{eq}}{C_r^{eq}}$ Organic to receiver partition coefficient

Biphasic Dissolution Experiment

- Biphasic dissolution is a useful tool for:
 - Assessing formulation solubility for amorphous or enabled formulations vs. crystalline drug
 - Fitting precipitation kinetics for weakly basic compounds with simultaneous absorptive phase
- Input Parameters:
 - Aqueous and organic diffusion layer thickness
 - Partition coefficient into organic phase (LogP surrogate for decanol)
 - System parameters – aqueous to organic contact area
- Output Results:
 - Formulation solubility determined from drug absorption rate
 - Precipitation kinetics

Biphasic Dissolution Model

- Allows simultaneous prediction of precipitation in aqueous phase while drug can be extracted into organic
- This will hopefully lead to better *in vitro* to *in vivo* extrapolation for precipitation
- IR formulations only



$$\frac{dm_o}{dt} = \frac{k_A k_o K_{o:w}}{k_A + k_o K_{o:w}} A_I \left(C_A - \frac{C_o}{K_{o:w}} \right)$$

m_o Mass in organic

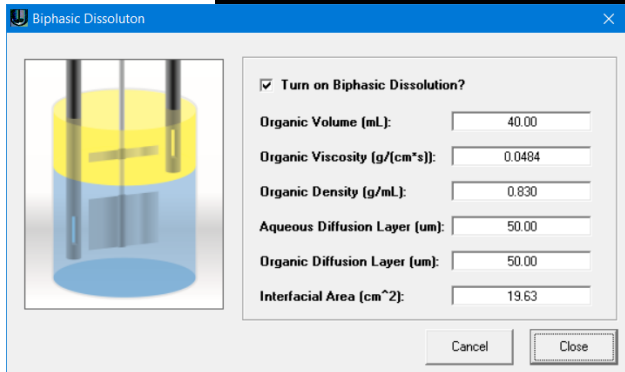
C_o Concentration in Organic

C_A Concentration in Aqueous

$k_A k_o$ Mass Transfer Coefficient in Aqueous and Organic

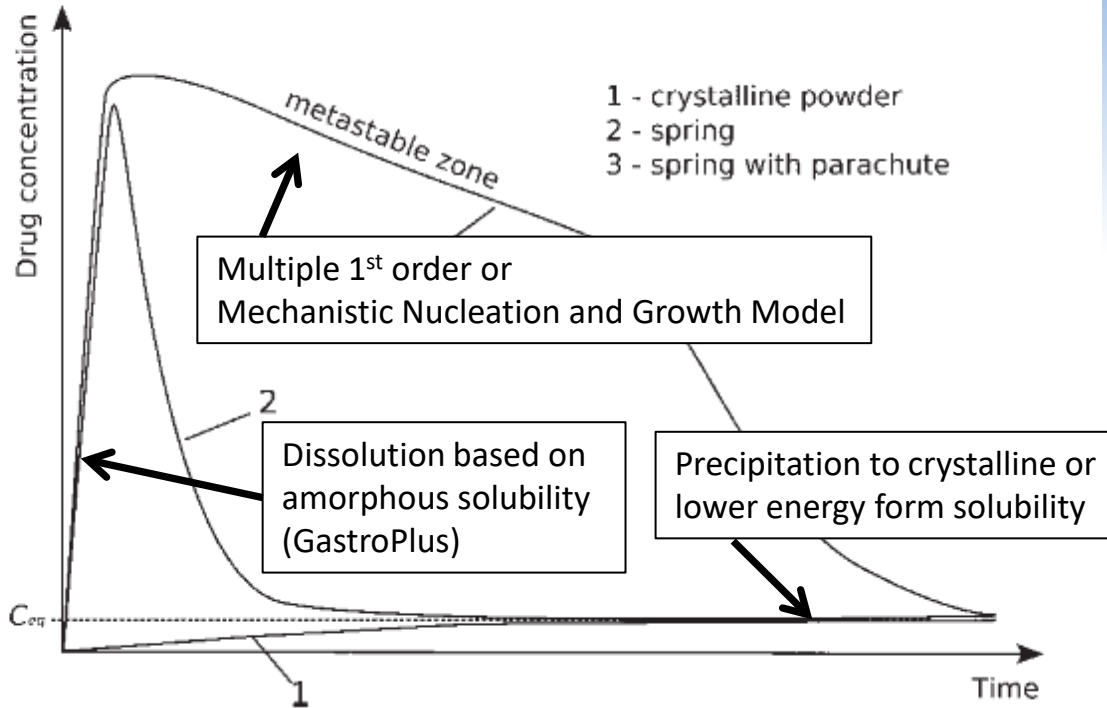
$K_{o:w}$ Organic/Water Partition Coefficient

A_I Interfacial Area



Precipitation and Supersaturating Drug Delivery Systems

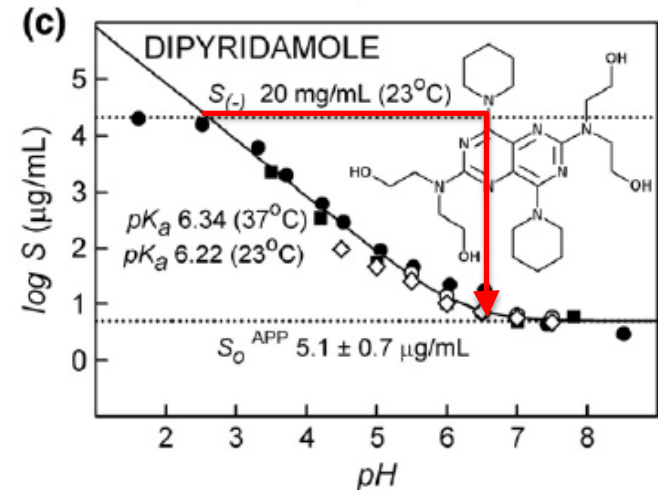
Schematic diagram of "Spring and Parachute"



Key Variables:

- C/S – Supersaturation Ratio

Basic compounds with high Salt Forms of Acidic Drugs
gastric vs. intestinal solubility



Brouwers J, J. Pharm. Sci. 98(8):2549 (2009)

Precipitation Model Options in DDDPlus/GastroPlus

Precipitation Model: First Order

Mechanistic Model Options:

Nucleation Type: Homogeneous Model Version: Lindfors

General Nucleation Inputs:

Nucleation Model: Diffusional Interf Iension (J/m²): 0.0189

Surface Integration Factor (Lambda, um): 0.06 Exp Correction Factor: 0.06

Precipitation Time Model Options:

Mean Precipitation Time (sec): 396

Precipitate Will:

form new particles with radius [um]: 4.27

"grow" particles in first bin only

"grow" all particles

OK Cancel

First order precipitation:

- Precipitate will form new particles with user-defined radius.
- Precipitate will “grow” in smallest particle size distribution bin only.
- Precipitate will “grow” all particles.
- Default precipitation time (900 s) determined from exponential fit to transfer assay data for dipyridamole from Kostewicz, 2004.
- Precipitation time as function of luminal pH.

Mechanistic Nucleation and Growth (MNG) precipitation model

Dipyridamole Transfer Assay First Order Precipitation

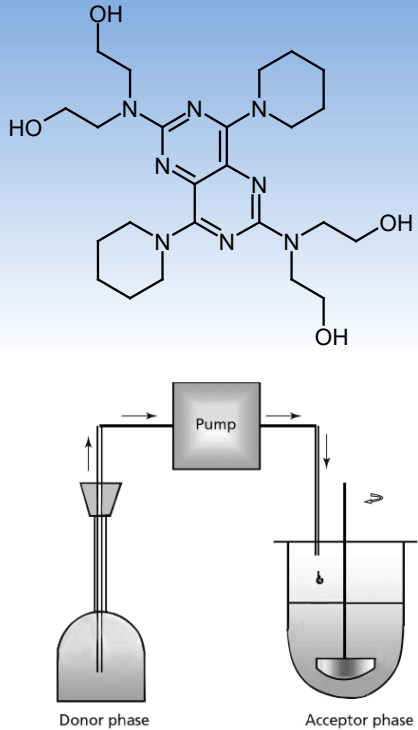
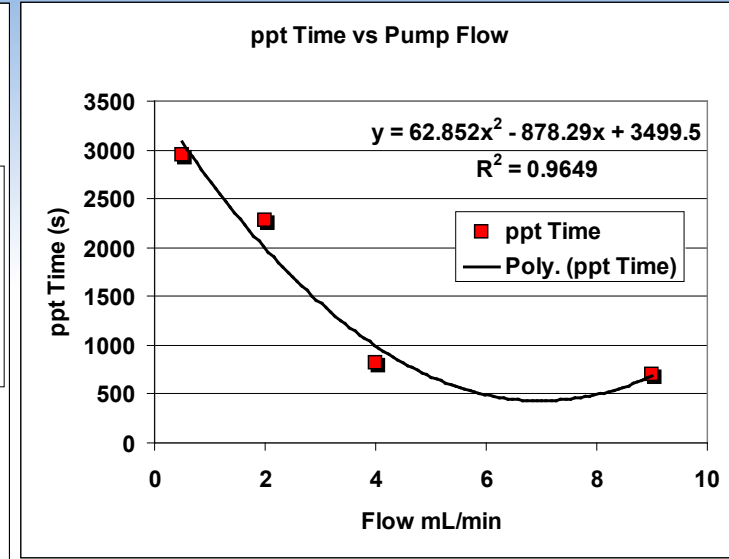
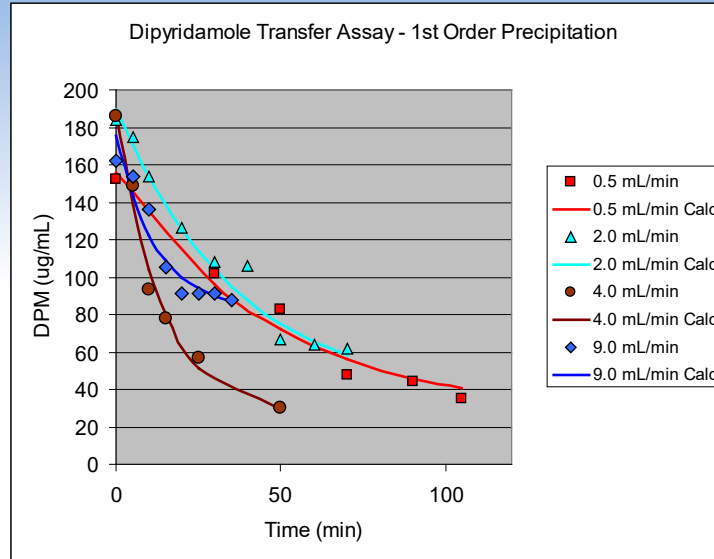


Figure 2 Experimental set-up to examine precipitation.



@ 4.25 mL/min the ppt. Time = 900 s

GastroPlus First Order Precipitation:

$$\frac{dM_i}{dt} = \left(\frac{V_i}{T_{precip}} \right) (C_i - S_i)$$

M_i – Mass in compartment i

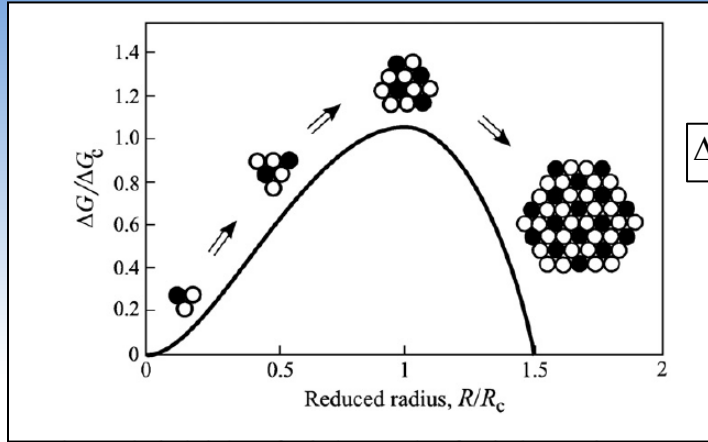
V_i – Volume of compartment i

T_p – Precipitation Time

C_i – Concentration in compartment i

S_i – Solubility in compartment i

Classical Nucleation and Growth



$$\Delta G_{system} = \Delta G_{cluster} + \Delta G_{bulk}$$

Rodriguez-Hornedo N, J. Pharm. Sci. 88(7):651 (1999)
 De Yoreo JJ, Rev. Mineralogy Geochem. 54:57 (2003)
Lindfors L, J. Colloid Interfac. Sci. 325 (2):404 (2008)
 Sugano K., Int. J. Pharmaceut. 378:142 (2009))

The nucleation rate J (the net production of critical clusters per unit time and unit bulk volume) is a function of the steady state concentration of the critical clusters and the transport of monomers to the critical clusters.

$$J = \text{Pre-exponential Term} * e^{\text{Exponential Term}}$$

Pre-exponential Term:

$$D_{mono} N_{Av} c^2 \left(\frac{k_B T}{\gamma} \right)^{1/2} \ln \left(\frac{c}{S} \right) \frac{R^*}{R^* + \lambda}$$

Exponential Term:

$$- \text{ExpCorr} \frac{16\pi}{3} \left(\frac{\gamma}{k_B T} \right)^3 \frac{v_m^2}{\ln \left(\frac{c}{S} \right)^2}$$

- D_{mono} = diffusion coefficient of the monomer (3.42E-4 cm²/min)
- N_A = Avogadro's number (6.02E+23 moles/mole)
- c = Conc. of free monomer (moles/cm³)
- S = Solubility at the current pH
- k_b = Boltzman's constant (1.38E-21 cJoules/Deg. K)
(Note: Joule = Newton-meter)
- $T = 310^\circ \text{K}$
- γ = Interfacial tension (Newtons/cm)
- v_m = Molecular volume = ($V_m/N_A = \text{XX cm}^3/\text{molec} / 6.02\text{E}+23 \text{ molec/mole}$)
- R^* = Critical radius (cm)
- λ = Effective radius from Lindfors (cm)
- ExpCorr = exponential correction factor

Case Study - Data




pharmaceutics

O'Dwyer, *Pharmaceutics* 12.3 (2020): 272.



Article

On the Usefulness of Two Small-Scale In Vitro Setups in the Evaluation of Luminal Precipitation of Lipophilic Weak Bases in Early Formulation Development

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- Comprehensive dataset for biphasic and membrane dissolution with corresponding *iv vivo* predictions
- Precipitation parameters were estimated based on aqueous concentration or organic concentrations for itraconazole. A full mechanistic interpretation was not performed.

Dose and Formulation Models

Dosing Biphasic Experiment:

- Dipyridamole
 - 10 mg crystalline suspension
 - 0.25 mg/mL dose (40 mL media)
- Ketoconazole
 - 20 mg crystalline suspension
 - 0.5 mg/mL dose (40 mL media)
- Itraconazole
 - 5 mg dose sporanox solution
 - 5 mg dose sporanox capsule
 - 0.125 mg/mL dose (40 mL media)

Dosing Membrane Experiment:

- Dipyridamole
 - 5 mg crystalline suspension
 - 0.25 mg/mL dose (20 mL media)
- Ketoconazole
 - 10 mg crystalline suspension
 - 0.5 mg/mL dose (20 mL media)
- Itraconazole
 - 2.5 mg dose sporanox solution
 - 2.5 mg dose sporanox capsule
 - 0.125 mg/mL dose (20 mL media)

Biphasic Dissolution Model Settings

Medium Composition - Acetate_Phosphate_PH2 - Phase 1 - Medium Composition pH: 2

Ingredient	Concentration (M)	Surfactant
Phosphoric Acid H3PO4	0.005	<input type="checkbox"/>
Sodium Acetate CH3COONa	0.005	<input type="checkbox"/>
▶ Hydrochloric Acid HCl	0.013	<input type="checkbox"/>
*		<input type="checkbox"/>

Medium Composition - FassiV2_10X - Phase 2 - Medium Composition pH: 13.21

Ingredient	Concentration (M)	Surfactant
Maleic acid	0.1912	<input type="checkbox"/>
Sodium Hydroxide NaOH	0.547	<input type="checkbox"/>
Sodium Chloride	0.6862	<input type="checkbox"/>
Sodium Taurocholate	0.03	<input checked="" type="checkbox"/>
▶ Phosphatidylcholine	0.002	<input checked="" type="checkbox"/>
*		<input type="checkbox"/>

Membrane Dissolution Model Settings

Left Window: 0.01 M Hydrochloric Acid

Medium: 0.01 M Hydrochloric Acid
Phase: Phase 1

Ingredient	Concentration (M)	Surfactant
▶ Hydrochloric Acid HCl	0.01	<input type="checkbox"/>
*		<input type="checkbox"/>

Medium Composition pH: 2

Buttons: Surfactant, Calculate pH, Edit, Close

Right Window: FaSSiF_V2_4X

Medium: FaSSiF_V2_4X
Phase: Phase 2

Ingredient	Concentration (M)	Surfactant
Maleic acid	0.07652	<input type="checkbox"/>
Sodium Hydroxide NaOH	0.17575	<input type="checkbox"/>
Sodium Chloride	0.27448	<input type="checkbox"/>
Sodium Taurocholate	0.012	<input checked="" type="checkbox"/>
▶ Phosphatidylcholine	0.0008	<input checked="" type="checkbox"/>
*		<input type="checkbox"/>

Medium Composition pH: 12.34

Buttons: Surfactant, Calculate pH, Edit, Close

Annotations: Fiber C (left), 15 mL (left), < Buffer (right), Red box around 'Surfactant' button (right)

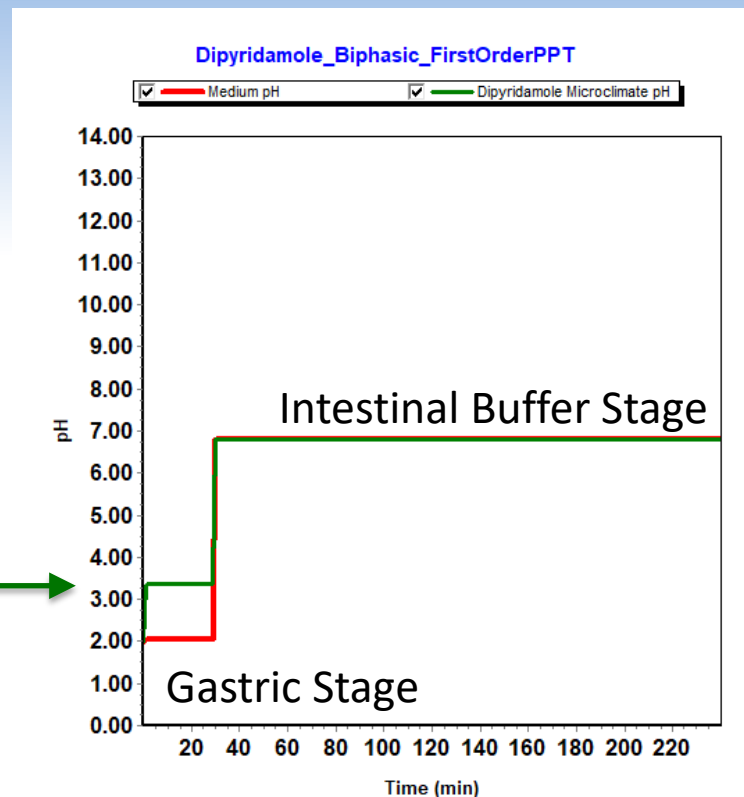
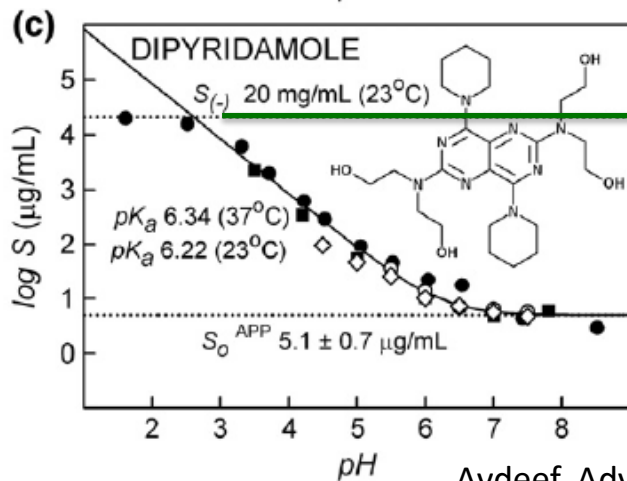
Dipyridamole pH Prediction

Multi-stage Biphasic Dissolution

Free base dissolving will change the local microclimate and solubility

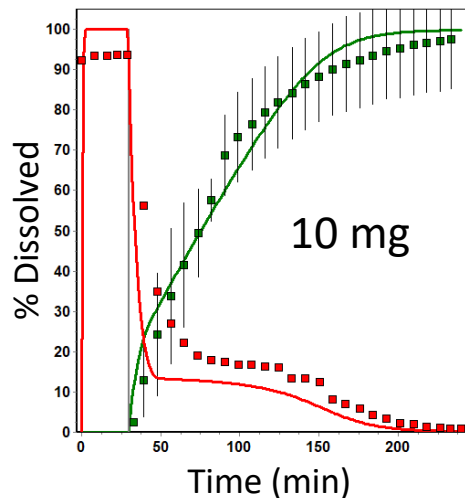
— Microclimate pH

— Media pH

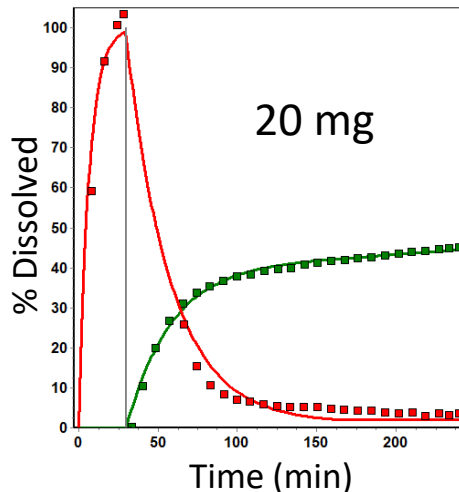


Biphasic First Order Precipitation Results

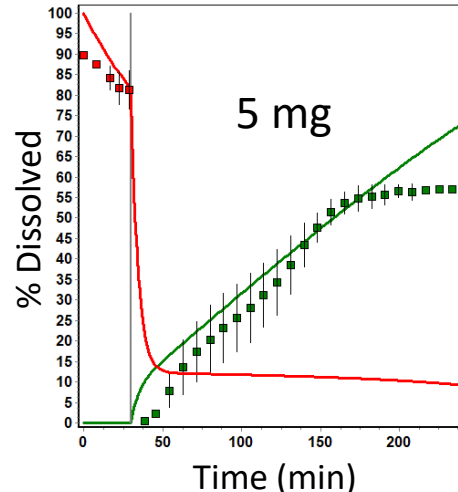
Dipyridamole



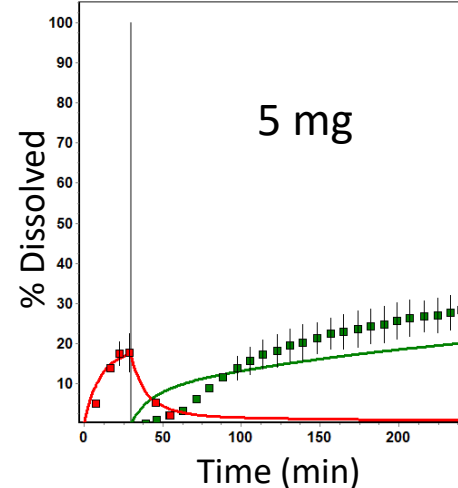
Ketoconazole



Itraconazole Solution



Itraconazole Capsule



PPT Time = 396 sec
PPT Size = 4.27 μm

Particle Size = 10 μm
PPT Time = 2857 sec
PPT Size = 1 μm

Ingredient	
Itraconazole	
Time, min	Prec. Time (s)
0	3800
30	350

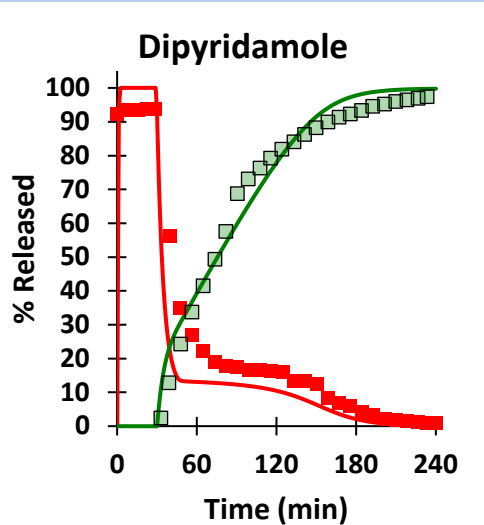
Particle Size = 13 μm
PPT Time = 900 sec
PPT Size = 1 μm

PPT Size = 1 μm

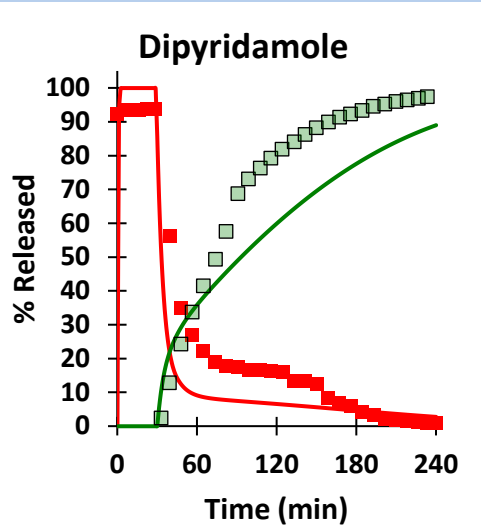
—■— Aqueous
—■— Organic

Parameter Sensitivity Analysis - Precipitate Size

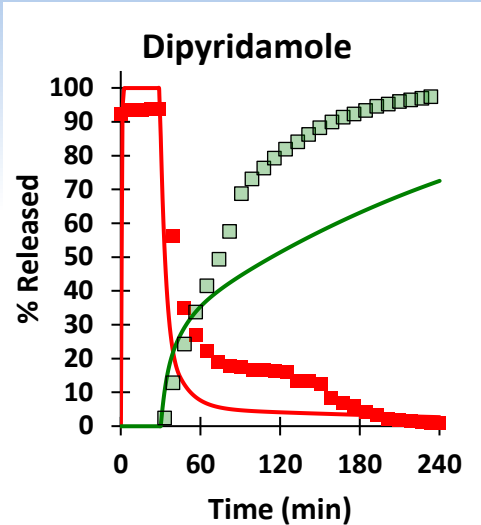
PPT Size = 4.27 μm



PPT Size = 15 μm



PPT Size = 30 μm

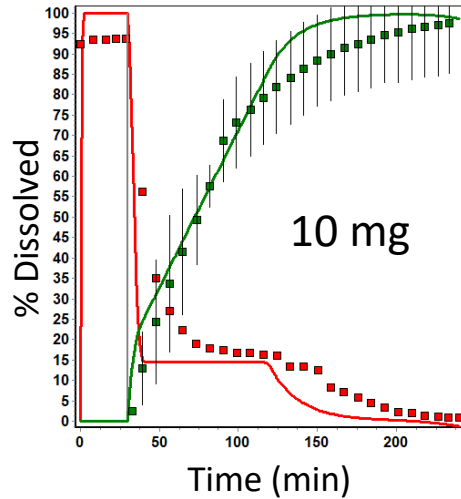


Redissolution of precipitate can be an important aspect that allows for appearance of drug in organic to be predicted correctly.

- Aqueous
- Organic

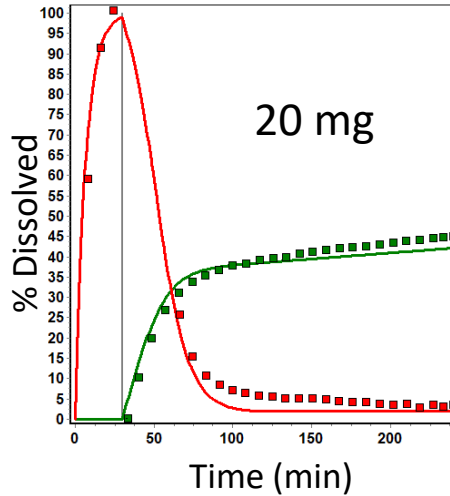
Biphasic Mechanistic Precipitation Results

Dipyridamole



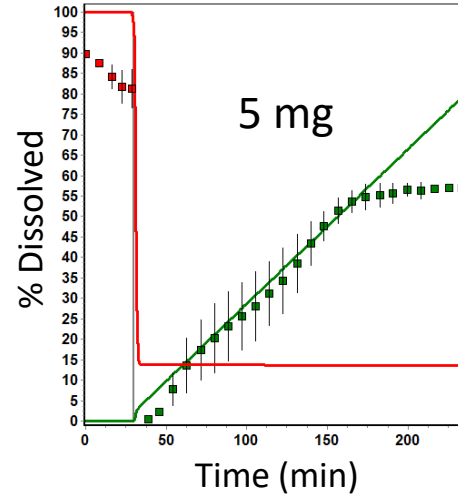
Surface Integration = 0.0371 μm
Exponential Correction = 0.0417

Ketoconazole



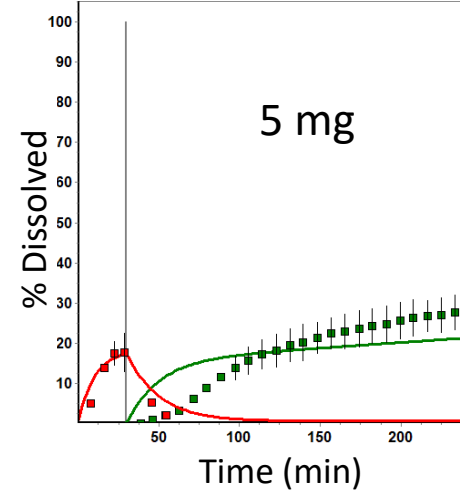
Surface Integration = 0.268 μm
Exponential Correction = 0.3

Itraconazole Solution



Surface Integration = 0.003 μm
Exponential Correction = 0.003

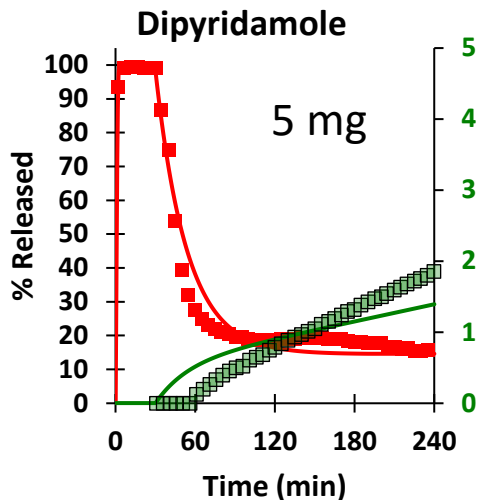
Itraconazole Capsule



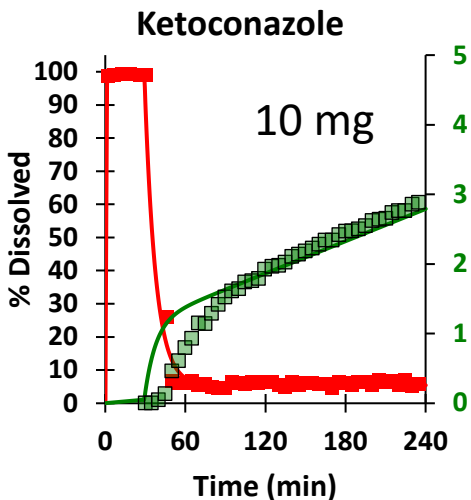
Surface Integration = 0.05 μm
Exponential Correction = 0.05

—■— Aqueous
—■— Organic

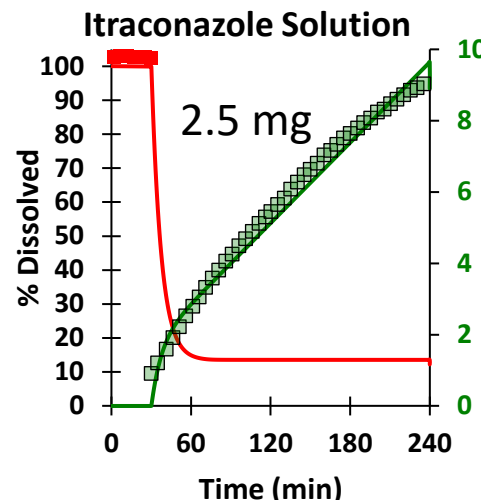
Membrane First Order Precipitation Results



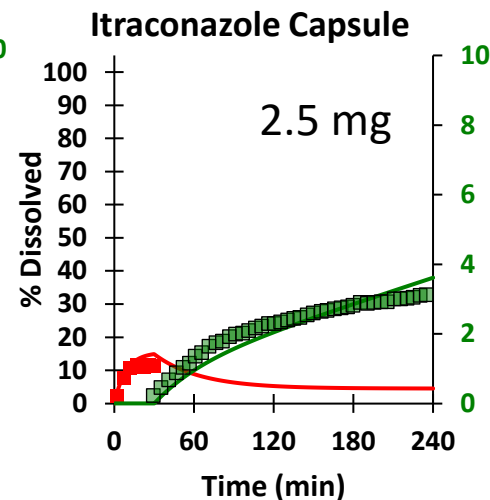
PPT Time = 1429 sec
PPT Size = 0.32 μm



PPT Time = 500 sec
PPT Size = 1 μm



PPT Time = 450 sec
PPT Size = 1 μm

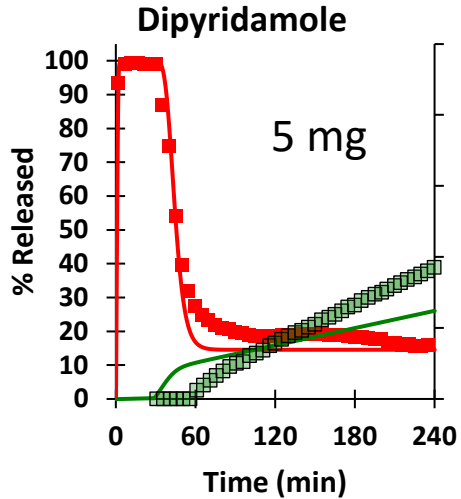


Particle Size = 13 μm
PPT Time = 2300 sec
PPT Size = 1 μm

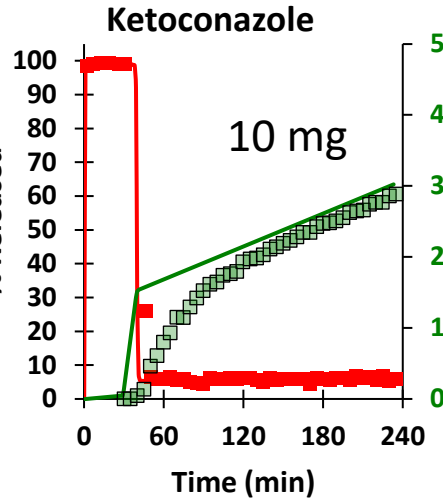
—■— Aqueous
—■— Receiver

Amount of drug appearing in receiver is based on the solubility difference between amorphous drug and cyclodextrin solution.

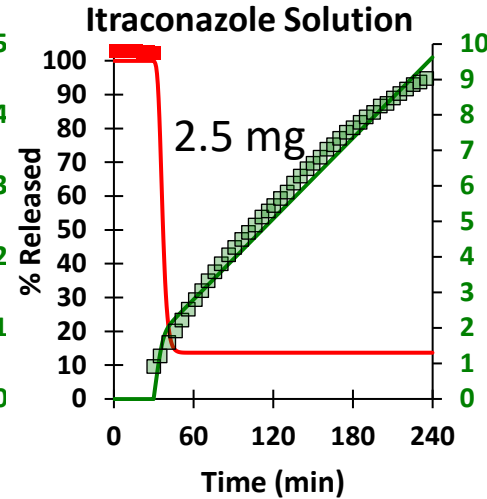
Membrane Mechanistic Precipitation Results



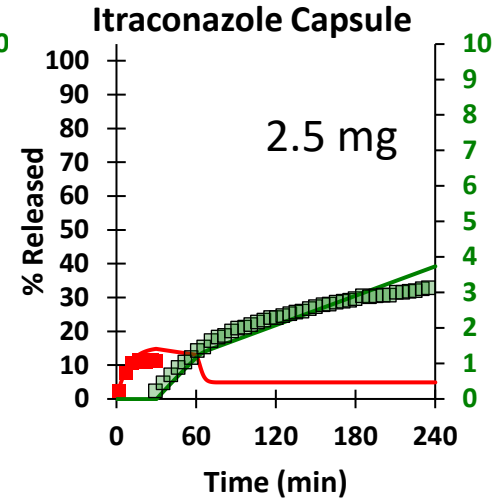
Surface Integration = 0.0639 μm
Exponential Correction = 0.0639



Surface Integration = 0.095 μm
Exponential Correction = 0.095



Surface Integration = 0.011 μm
Exponential Correction = 0.011

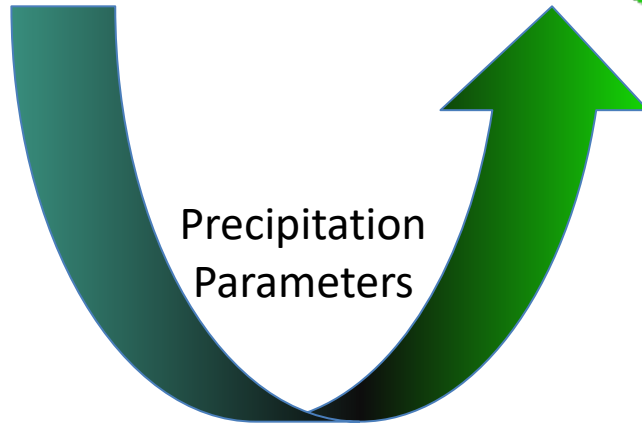


Surface Integration = 5.60e-4 μm
Exponential Correction = 5.60e-4

■ Aqueous
■ Receiver

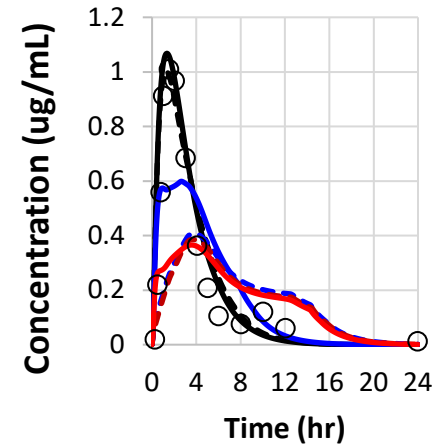
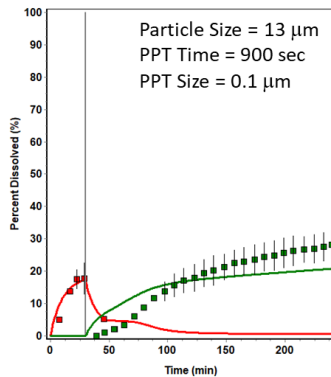
Slope of drug appearing in receiver is based on the solubility difference between amorphous drug and cyclodextrin solution.

In Vitro Dissolution in GastroPlus Models

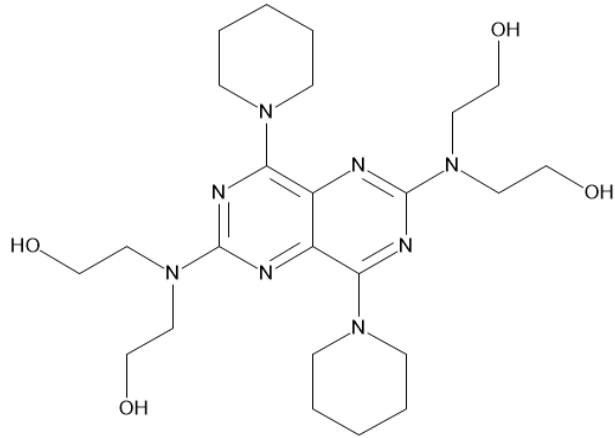


Precipitation
Parameters

First Order Precipitation



Dipyridamole GastroPlus Model



Molecular Weight: 504.64 g/mol

AP 10 = ADMET Predictor v.10

S+ stands for properties predicted with Simulations Plus models

S+Sw = solubility in pure water

S+SF = solubility factor

S+Peff = human jejunal permeability

S+PrUnbnd = percent unbound in human plasma

S+Rbp = human blood/plasma concentration ratio

S+logP = 3.04 (AP 10.0)

Exp LogP = 3.71 @ pH 7 (FDA Label)

S+pKa = 5.83, 3.56, 1.83, 1.63, 1.43, 1.13, 0.71, 0.29 (AP 10.0)

Exp pKa = 6.307 (Fit to multiple experimental sources)

S+Sw = 2.34 mg/mL @ pH = 8.75 (AP 10.0)

Exp Sw = 0.005 mg/mL @ pH = 11 (Fit to *in vitro* data)

S+Peff = Human: 0.45×10^{-4} cm/s (AP 10.0)

EXP Caco = 1.124×10^5 cm/s

Human Peff = 1.81×10^{-4} cm/s (Converted from Exp)

S+PrUnbnd (human) = 8.99 % (AP 10.0)

Exp PrUnbnd (%) = 0.20% (FDA Label)

S+Rbp (human) = 0.67 (AP 10.0)

Exp Rbp = 0.647 (Serebrauny 2009, calc)

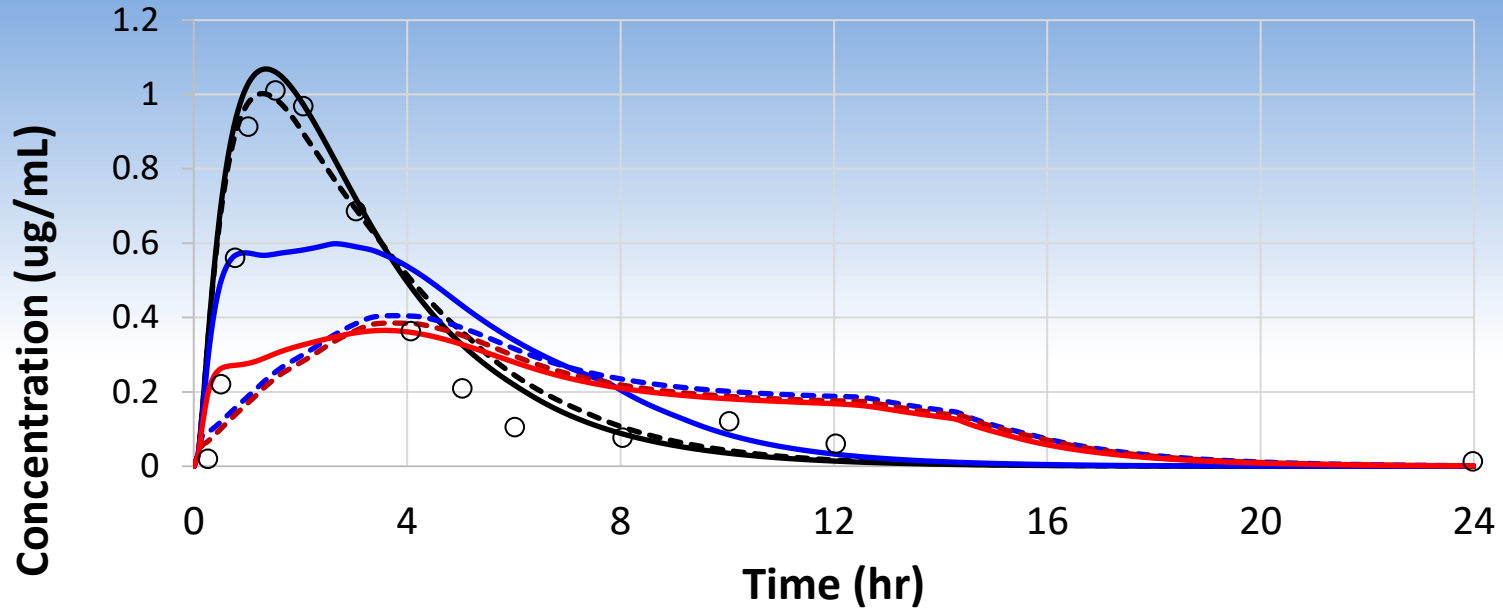
Clearance (L/hr/kg) = 0.195 (Nielson-Kudsk 1979)

Clearance (L/hr) = 12 (Australian Label)

Clearance average (L/hr) = 13.59

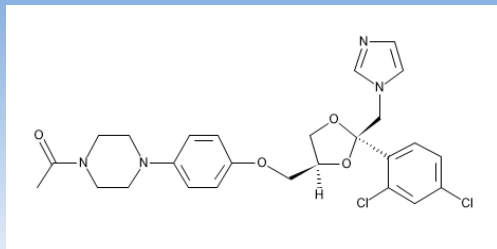
PBPK model adjusted to experimental Vdss
by adjusting LogP to 3.184 to calculate Kp

Dipyridamole 75 mg Tablet Precipitation *IVIVE*

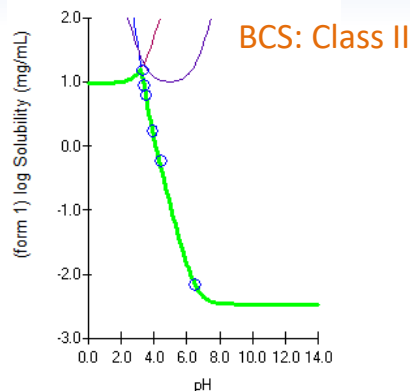


- No Precipitation
- - - Observed PPT Time = 22000 sec
- - - Membrane Disso Mech PPT
- Biphasic PPT Time = 396 sec
- Exp.
- Membrane Disso PPT Time = 1429 sec
- - - Biphasic Mech PPT

Ketoconazole GastroPlus Model



Molecular Weight: 531.44 g/mol



AP9.5= ADMET Predictor v. 9.5
S+ stands for properties predicted with Simulations Plus models
S+Sw = solubility in pure water
S+Peff = human jejunal permeability
S+PrUnbnd = percent unbound in human plasma
S+Rbp = human blood/plasma concentration ratio

S+logP = 3.74 (AP 9.5)

Exp LogP = 3.73 (Walter-JPharmPharmacol-40-689-1988)

S+pKa = 6.15 (base 1), 4.22 (base 2) (AP 9.5)

Exp pKa = 6.51 (base 1), 3.4 (base 2) (Walter-JPharmPharmacol-40-689-1988)

S+Sw = 0.11 mg/mL @ pH = 8.23 (AP 9.5)

Exp Sw = 0.0069 @ pH = 6.5 (Poelma-JPharmPharmacol-43-5-317)

S+Peff = Human: 3.38×10^{-4} cm/s (AP 9.5)

Exp Caco = 1.35×10^5 cm/s (Ingels-International Journal of Pharmaceutics-274-221-2004)

Human Peff = 3.22×10^{-4} cm/s (Converted from caco2 permeability value from Ingels data (Ingels-International Journal of Pharmaceutics-274-221-2004)

S+PrUnbnd (human) = 4.38 % (AP 9.5)

Exp PrUnbnd (%) = 1 % (Daneshmend-ClinPharmacokinet-14-13-1988)

S+Rbp (human) = 0.69 (AP 9.5)

Exp Rbp = 0.6 (Matthew-PharmRes-10-3-418-1993)

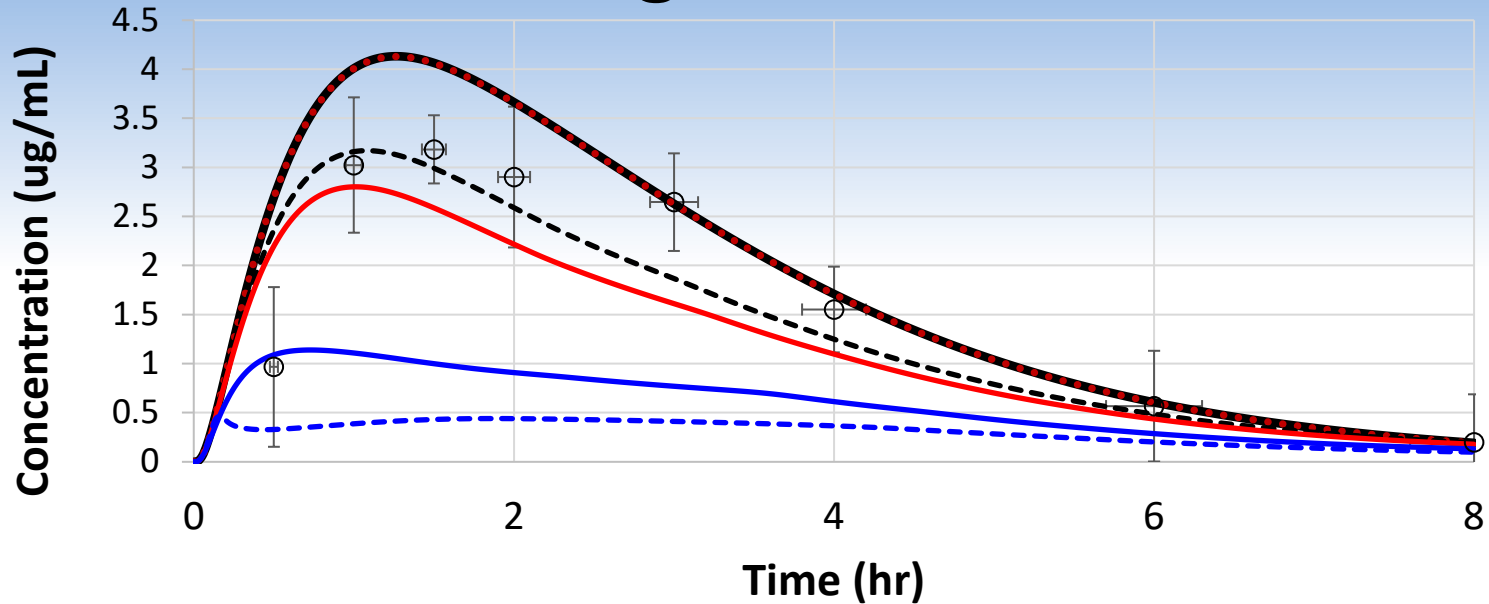
Changed log D (7.0) = 2.6 to calculate Kps and then changed back to 3.73 to run simulation

Metabolism CYP3A4: Km = 15 nM with Vmax = 0.048 nmol/min/mg Prot.

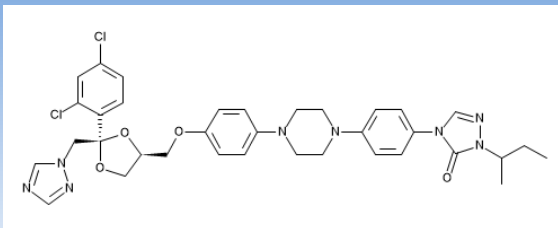
Transporters Pgp: Gut Km = 4.57 mg/L, Vmax = 0.006 mg/s

PBPK Km = 4.57 mg/L, Vmax = 0.001 m/s/mg trans

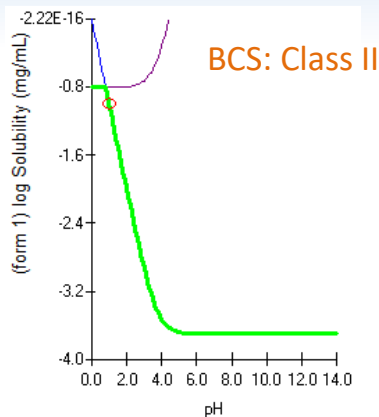
Ketoconazole IVIVE Precipitation 200 mg IR Tablet



Itraconazole GastroPlus Model



Molecular Weight: 705.65 g/mol



AP9.5= ADMET Predictor v. 9.5
S+ stands for properties predicted with Simulations Plus models
S+Sw = solubility in pure water
S+Peff = human jejunal permeability
S+PrUnbnd = percent unbound in human plasma
S+Rbp = human blood/plasma concentration ratio

S+logP = 4.89 (AP 9.5)

Exp LogP = 5.66 (FDA SPORANOX® (itraconazole) Capsule Label Information)

S+pKa = 4.57, 3.69, 2.93, 1.28 (AP 9.5)

Exp pKa = 3.7 (FDA SPORANOX® (itraconazole) Oral Solution Information" 2002)

S+Sw = 0.00733 mg/mL @ pH = 7.07 (AP 9.5)

Exp Sw = 0.1 mg/mL @ pH = 1.2 @ 60 min (Yin-DrugDesDevelTher-9-2801-2015)

GSE Sw = 0.000189 mg/mL (Yalkowski Equation)

S+Peff = Human: 1.85 x10⁻⁴ cm/s (AP 9.5)

Mem+ Caco = 2.66 x10⁵ cm/s

Human Peff = 3.85 x 10⁻⁴ cm/s (Converted from MembranePlus Peff)

S+PrUnbnd (human) = 2.75 % (AP 9.5)

Exp PrUnbnd (%) = 0.20% (FDA Label)

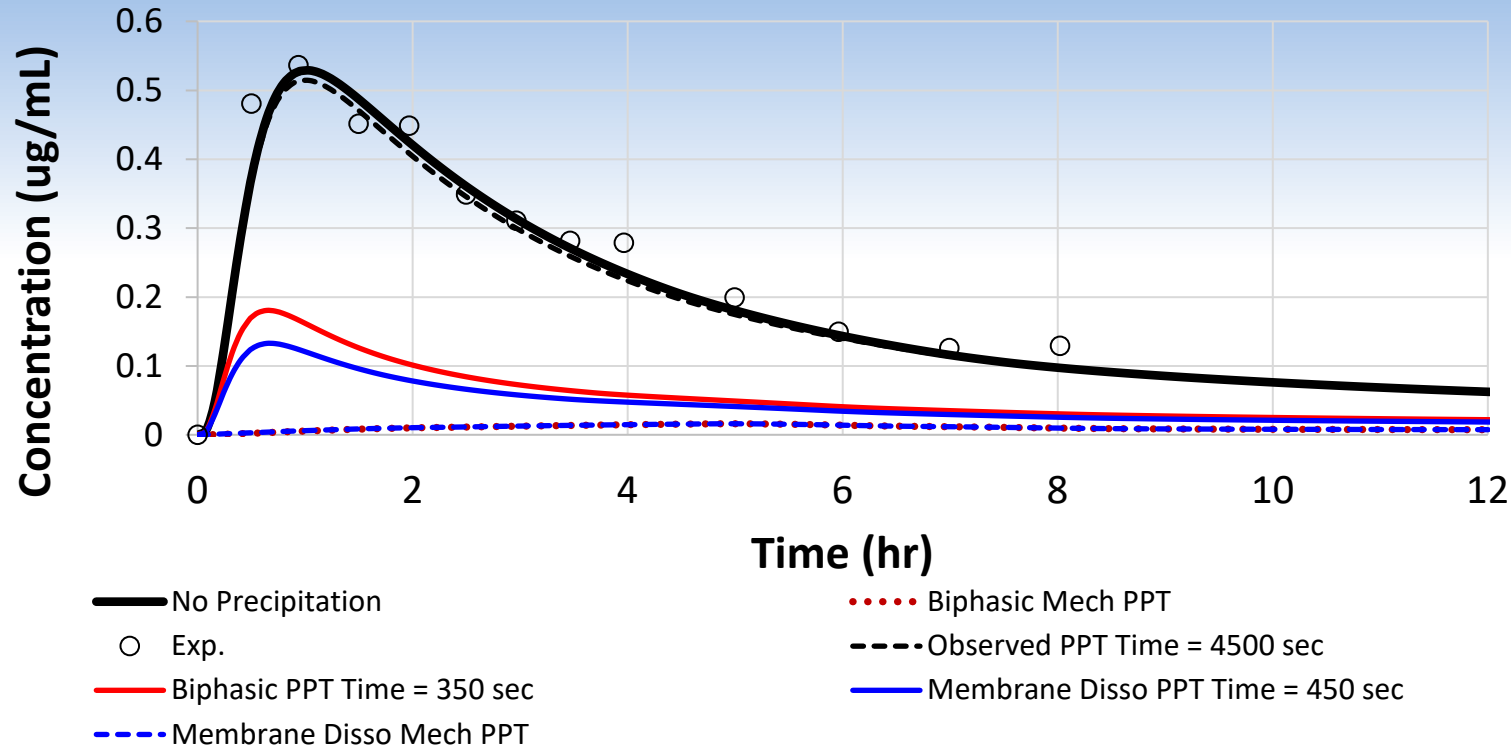
S+Rbp (human) = 0.67 (AP 9.5)

Exp Rbp = 0.58 (Kato-PharmRes-25-8-1891-2008)

Metabolism CYP3A4: Km = 0.00275 mg/L, Vmax (gut) = 0.0764 nmol/min/mg Prot.

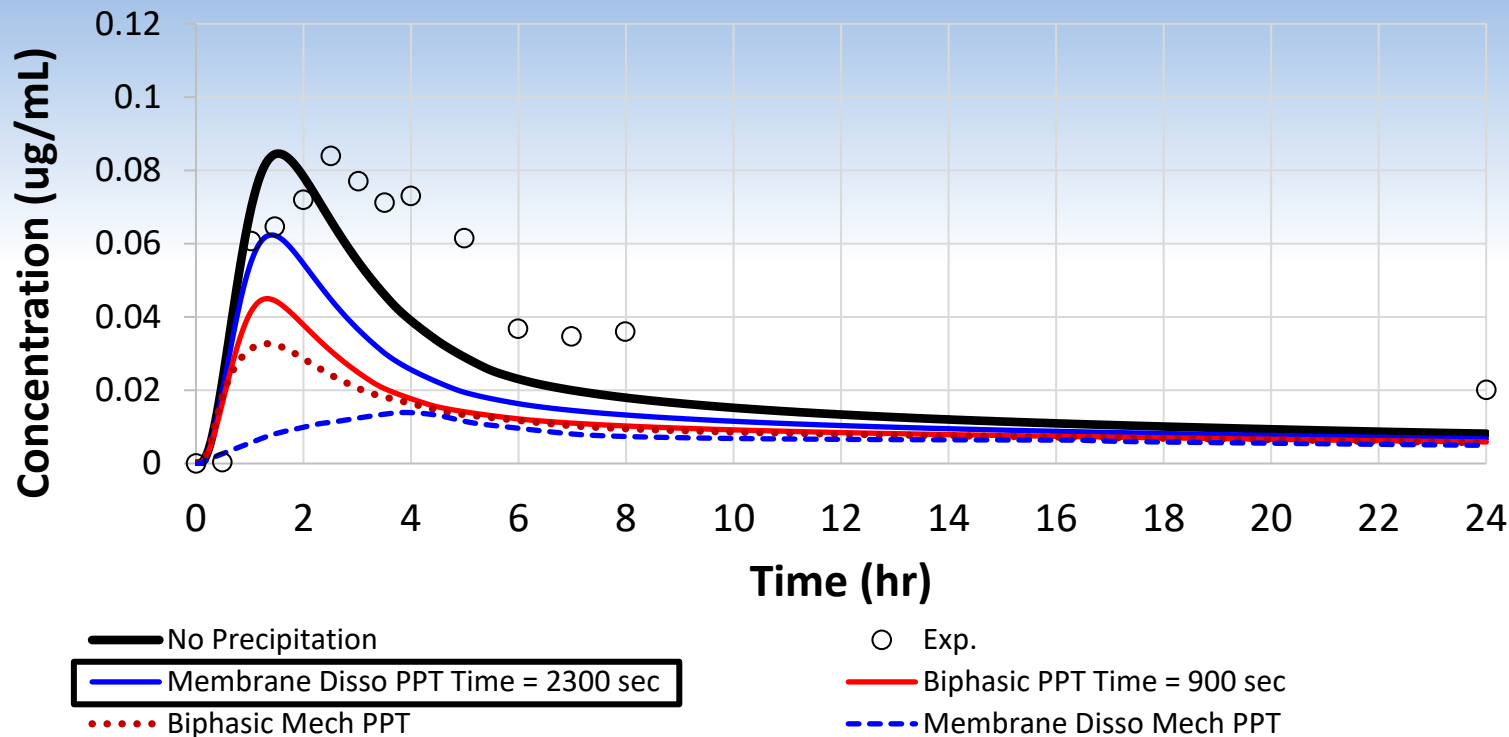
Vmax (pbpk) = 0.00022

Itraconazole IVIVE Precipitation 200 mg IR Solution



Itraconazole IVIVE Precipitation

200 mg IR Capsule



Conclusion

- DDDPlus provides complex models to handle:
 - Membrane dissolution
 - Biphasic dissolution
 - Artificial Stomach Duodenum test
- IVIVE is challenging for all complex *in vitro* tests
 - Precipitation is best optimized to experimental PK data in our current understanding
 - However, *in vitro* tests provide valuable information on propensity to crystallize and formulation solubility