# Usefulness of physiology-based PK to mitigate risk in early clinical drug development?

May 18th 2017

An Van den Bergh



# Strategic questions as foundation for early PBPK

- Will the predicted PK profile meet the clinical needs?
- Likelihood or understanding of dose (dis) proportionality?
- Accumulation during repeated dosing as expected?
- Drug-Drug Interactions in Patients?
- Among a set of potential candidates, which one has the most favourable PK profile?
- Can we anticipate a PK difference between healthy volunteers and patients?



Understand components governing PK drug behaviour

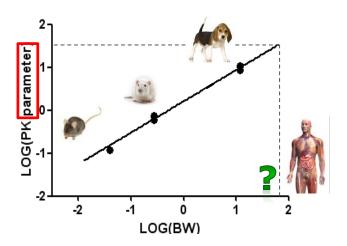






# FIH modeling approaches

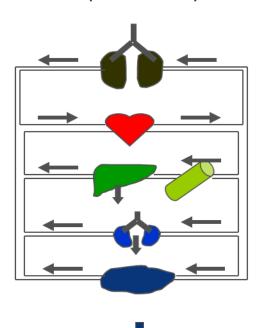
Allometry (empirical)





Extrapolation of isolated PK parameters (CL & Vd)

Physiology Based PK (PBPK) (mechanistic)

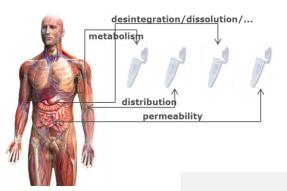






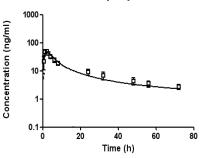
# **PBPK** modeling

Integration of multiple sources of individual data into the physiological context

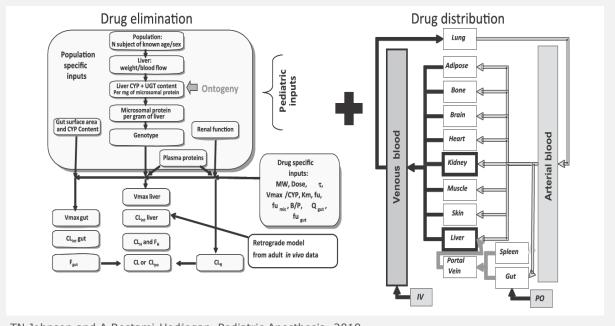




# Predicting human PK .. and have a prepared mind



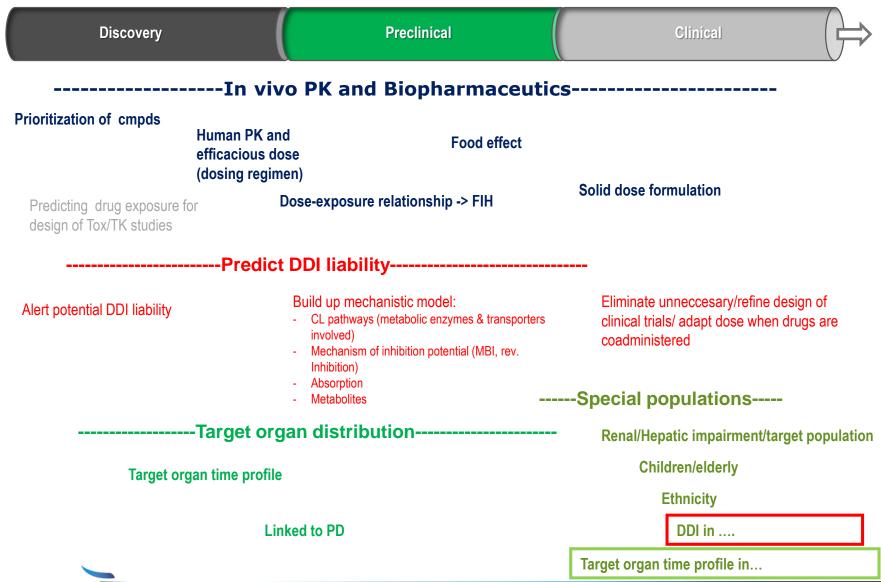
-virtual physiology -compound specific parameters -differential equations



TN Johnson and A Rostami-Hodjegan, Pediatric Anesthesia, 2010

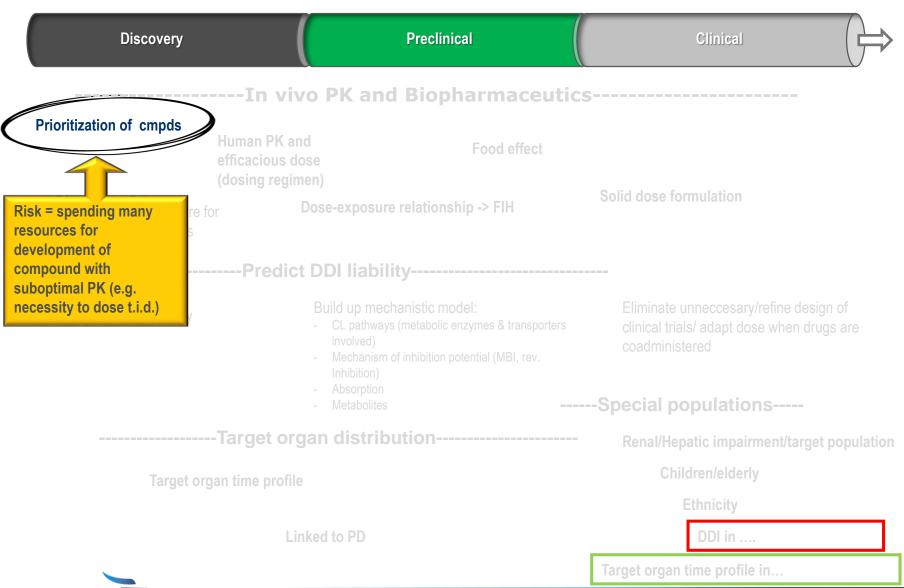


# PBPK modeling throughout development





# PBPK modeling and mitigating risks

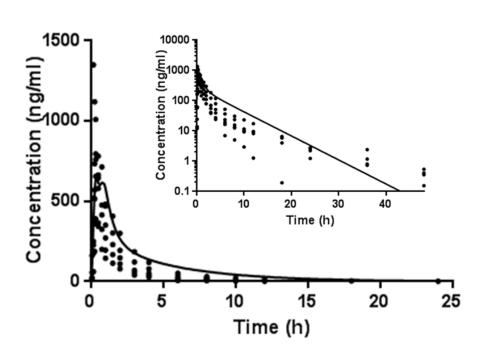




Background: PK of lead cmpd is not optimal (half life short; high fluctuation peak trough)

→ evaluate potential of 2 back-ups, integrating preclinical ADME and human microdose study data prior to SAD study without need for extensive TOX support

	JNJ-X
MW	$\checkmark$
Lipophilicity (logP)	<b>/</b>
Ionization constant (pKa)	$\checkmark$
Solubility (mg/mL)	
Human effective permeability (cm/s)	$\checkmark$
Blood to plasma ratio	
Plasma unbound fraction	$\checkmark$
Microsomal unbound fraction	
Intrinsic clearance hepatocytes	<b>/</b>
Volume of distribution (tissue partitioning)	<b>V</b>





All necessary information available for lead compound



Background: PK of lead cmpd is not optimal (half life short; high fluctuation peak trough)

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	JNJ-X	JNJ-Y	JNJ-Z
MW	$\checkmark$	<b>4</b>	<b>V</b>
Lipophilicity (logP)	<b>/</b>	<b>4</b>	<b>V</b>
Ionization constant (pKa)	<b>/</b>	<b>4</b>	$\checkmark$
Solubility (mg/mL)	<b>/</b>	<b>~</b>	X
Human effective permeability (cm/s)	<b>4</b>	X	X
Blood to plasma ratio	<b>4</b>	<b>~</b>	X
Plasma unbound fraction	$\checkmark$	<b>/</b>	$\checkmark$
Microsomal unbound fraction	<b>/</b>	X	X
Intrinsic clearance hepatocytes	<b>V</b>	<b>VX</b>	VX
Volume of distribution (tissue partitioning)	<b>V</b>	<b>V</b>	X



Limited information available for back-up compounds

→ assumptions to be taken

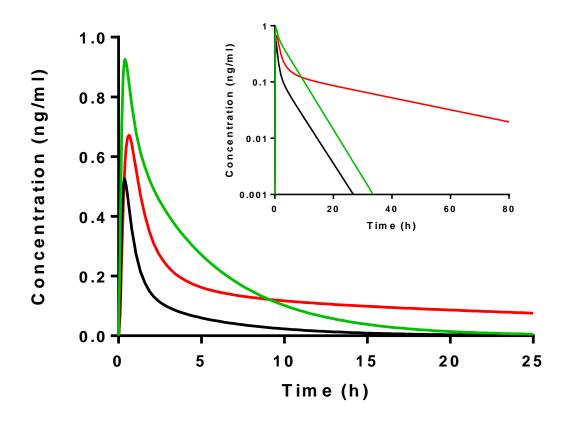


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#### Results microdose study

100ug JNJ-X 100ug JNJ-Y 100ug JNJ-Z



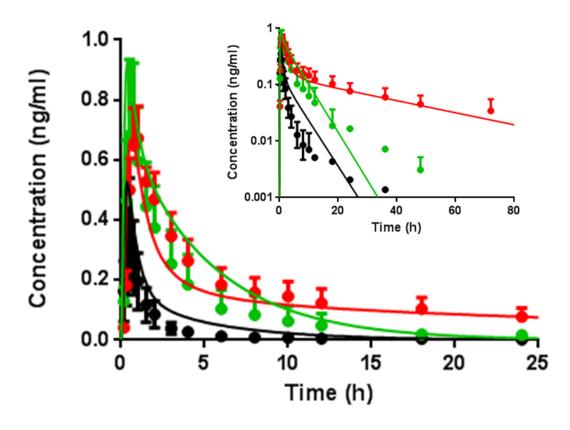


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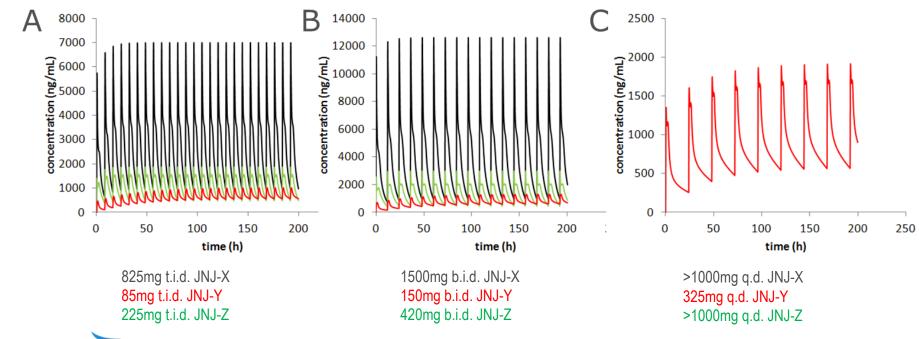
#### Predictions to reach efficacious concentrations

Efficacious concentrations:

1000ng/mL for JNJ-X

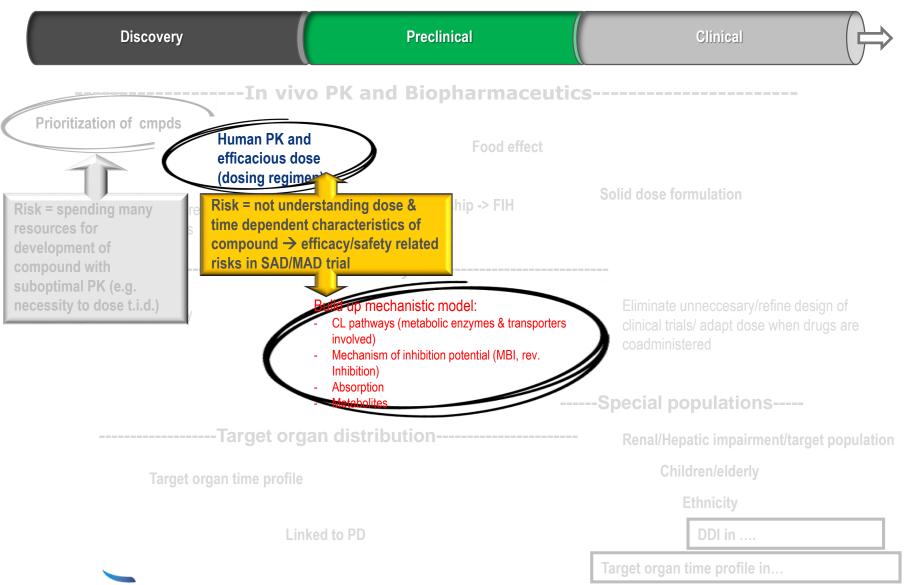
570ng/mL for JNJ-Y

450ng/mL for JNJ-Z





# PBPK modeling and mitigating risks

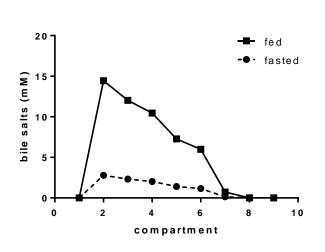


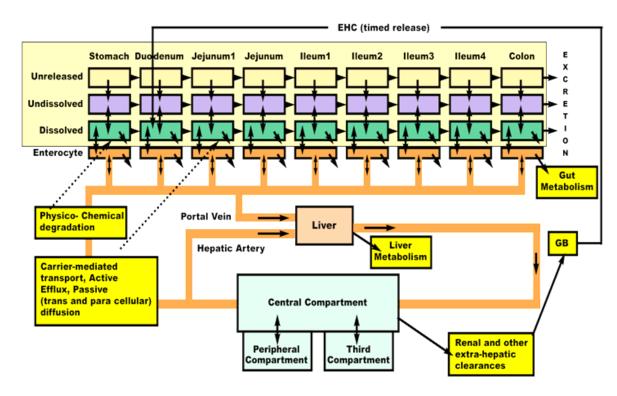


## Anticipate less than dose proportional absorption in SAD

Background: JNJ-X has a high permeability and a low solubility (0.0003mg/mL @ pH7 – note: pKabase<2) Large effect of bile salts due to high lipophilicity (Fassif 0.005 mg/mL; Fessif 0.011mg/mL) Metabolized by CYP3A4 (95%)

→ will enough exposure be reached in SAD and/or from which dose onwards will nonlinearity kick in?



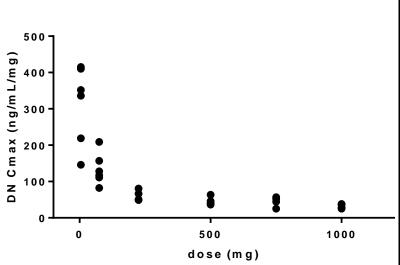




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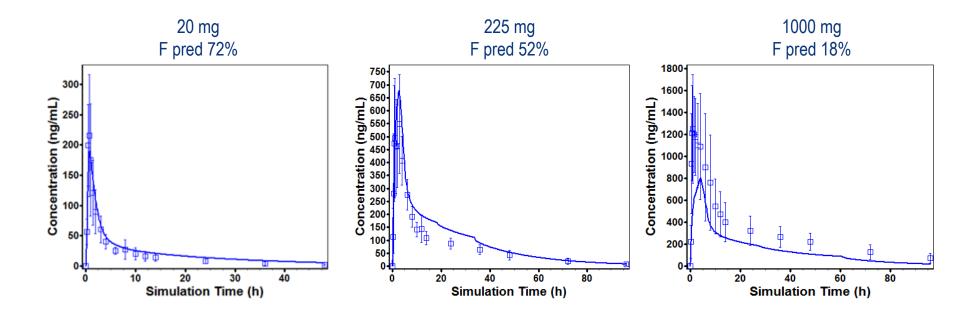
Dose	Cmax	AUC inf	F (bioavailability)
(mg)	(ng/ml)	(ng.hrs/ml)	(%)
5	14-38	36-154	41-73
20	58-102	312-513	37-61
75	105-243	556-1384	22-44
225	311-722	1105-3405	17-33
500	690-916	2140-4188	15-19
1000	1380-1830	4032-7895	14-18
1500	2069-2744	5294-11600	14-18



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Background: JNJ-X is planned to go FIH

Preclinically more than dose proportional PK is observed

What will be the predicted human PK profile? What should be the starting dose? And which dosing steps

should be taken in the SAD trial?

Dog p.o. data		5 mg/kg	
F (%)	(	178	
Cmax (ng/ml)		1980	
Tmax (h)		<b>2</b> -4	
AUC 0-inf (ng.h/ml)		12200	
		<b>↓</b>	

Due to saturation in metabolic pathways? (i.v. study in the dog @ 0.2 mg/kg: moderate clearance)



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F (%)	178	69	61
Cmax (ng/ml)	1980	290	17.1
Tmax (h)	<b>2</b> -4	0.5-2	0.5-2
AUC 0-inf (ng.h/ml)	12200	1420	125
	1		

Due to saturation in metabolic pathways? (i.v. study in the dog @ 0.2 mg/kg: moderate clearance)

Lower doses tested in the dog

→ Saturation kicks in already between 1 and 5mg/kg



Saturation tested in human hepatocytes.
Also there, saturation to be expected from low doses onwards

- Risk of very fast clearance at starting dose, hence very low concentrations
   → detection limit??
- Escalating doses should be carefully chosen

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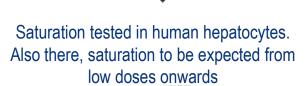
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→ detection limit??

Escalating doses should be carefully chosen

PBPK model can guide those decisions



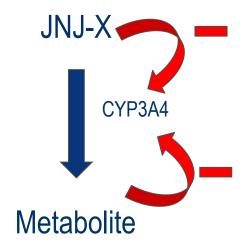
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### Anticipating exposure during MD? Auto-inhibition case

Background: JNJ-X is a high clearance compound; mainly metabolized by CYP3A4

It is also a reversible and time-dependent inhibitor of CYP3A4

- → it may inhibit its own metabolism
- → from which concentration onwards will nonlinearity kick in in the MAD study?

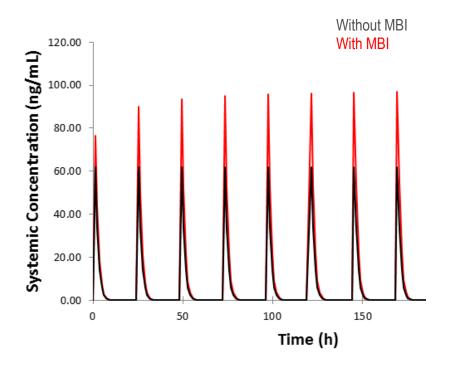




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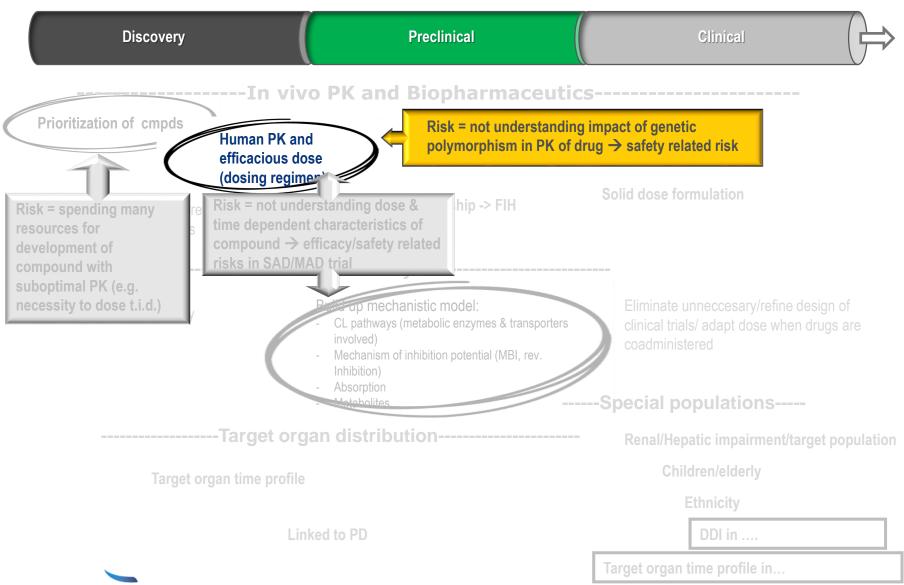
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# PBPK modeling and mitigating risks





#### Anticipate impact of genetic polymorphism in PK profile

Background: Clopidogrel is a prodrug that is metabolized by an esterase-dependent pathway leading to an inactive metabolite and a CYP450 dependent pathway leading to its active metabolite mediated mainly by CYP2C19 Polymorphisms of CYP2C19 affect the PK (and PD) of clopidogrel

Can impact of different CYP2C19 activ. (poor, intermediate, extensive, ultrarapid metabolizers) be anticipated?

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http://dx.doi.org/10.1124/dmd.114.062596 Drug Metab Dispos 43:510-522, April 2015

# Physiologically Based Pharmacokinetic Modeling for Sequential Metabolism: Effect of CYP2C19 Genetic Polymorphism on Clopidogrel and Clopidogrel Active Metabolite Pharmacokinetics

Nassim Djebli, David Fabre, Xavier Boulenc, Gérard Fabre, Eric Sultan, and Fabrice Hurbin

Sanofi R&D, Drug Disposition, Disposition Safety and Animal Research, Montpellier, France

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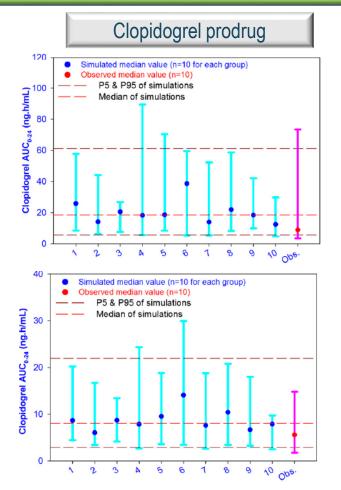
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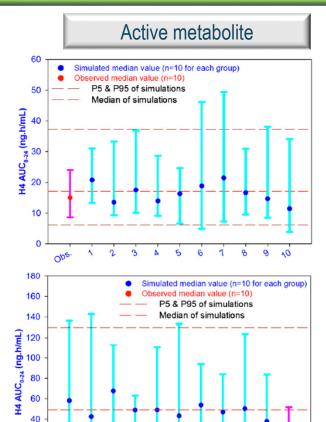
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Can impact of different CYP2C19 activ. (poor, intermediate, extensive, ultrarapid metabolizers) be anticipated?

Poor metabolizers

> Ultrarapid metabolizers

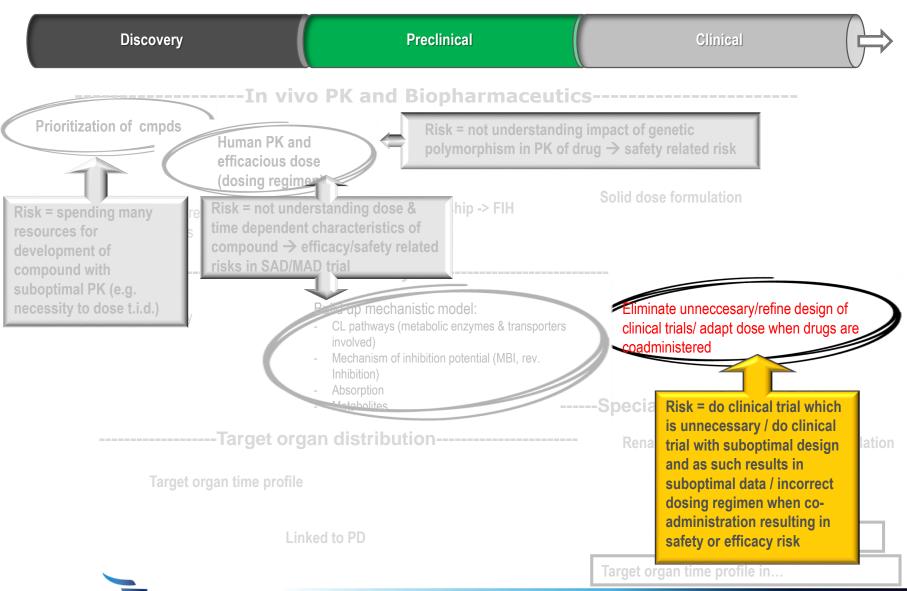






PHARMACEUTICAL COMPANIES
OF Johnson Johnson

# PBPK modeling and mitigating risks

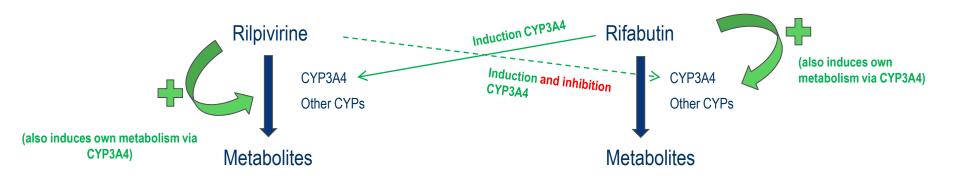


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## Adapting dose in case of induction by rifabutin?

Background: Rifabutin lowers the concentrations of rilpivirine.

- → dose of rilpivirine x 2 to overcome the decrease in exposure upon coadministration
- → what to do after stopping rifabutin? (when rifabutin is stopped an ongoing inducing is expected for a few more days)

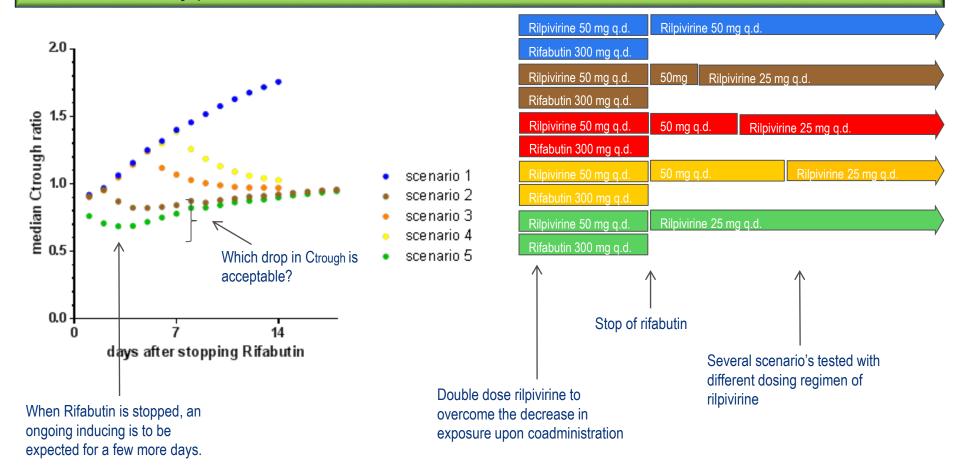




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# **Summary**

#### Value of PBPK in Risk Mitigation in Early Studies.

- Predict time profile of drug concentrations based upon in vitro ADME data and plausible biology
- Confirm mechanisms governing the (nonlinear) PK during dose escalation in the first human SAD & MAD studies
- Anticipate impact of genetic polymorphism in PK profile
- Anticipate drug interactions in First-in-Patient studies
- Differentiate drug candidates

"The Prepared Mind" (Dr. Paul Janssen, 1983)



# **Acknowledgments**

- Achiel Van Peer
- Loeckie De Zwart
- Jan Snoeys
- Marieke Voets
- Mario Monshouwer

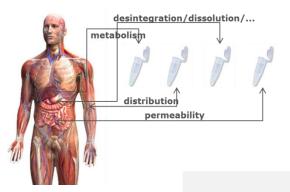


# **Back-ups**



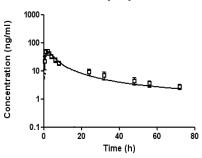
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Integration of multiple sources of individual data into the physiological context

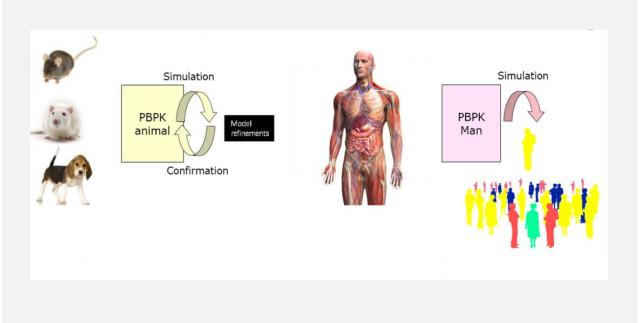




Predicting human PK
.. and have a prepared mind



-virtual physiology -compound specific parameters -differential equations

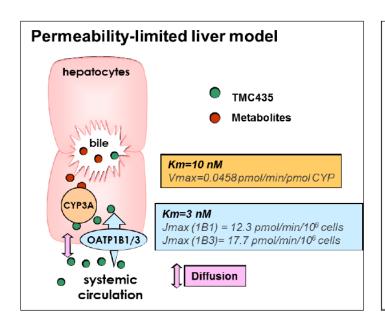


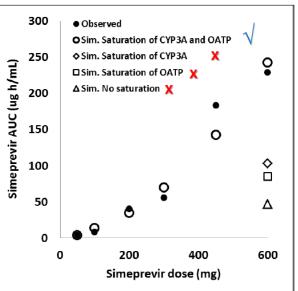
**Background:** 



#### Case Study: FDA Review of Simeprevir

Can saturation mechanisms explain nonlinear pharmacokinetics of simeprevir?





http://www.accessdata.fda.gov/drugsatfda\_docs/nda/2013/205123Orig1s000ClinPharmR.pdf

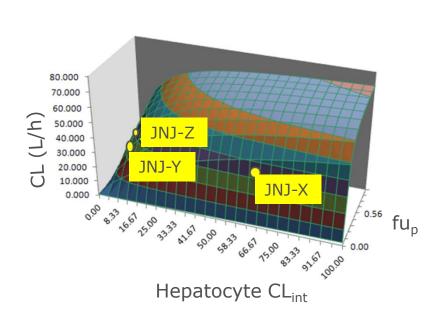


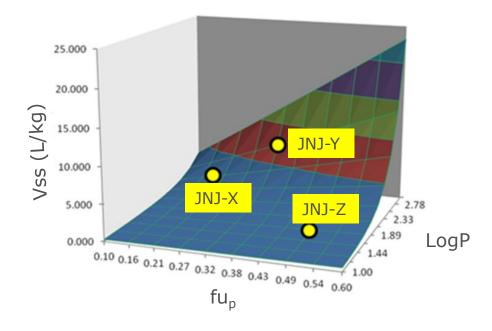
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# Discovery: candidate differentiation

Aim: PK of lead cmpd is not optimal (half life short; high fluctuation peak trough)

→ evaluate potential of 2 back-ups, integrating preclinical ADME and human
microdose study data prior to SAD study without need for extensive TOX support





Even though hepatocyte Clint is 7-8 fold lower for JNJ-Y and JNJ-Z

→ Clearance is only ~2 fold lower for JNJ-Y and JNJ-Z

Higher lipophilicity (logP) of JNJ-Y

→ Higher volume of distribution of JNJ-Y

Terminal half-life reasonably longer for JNJ-Y compared to the 2 other compounds (related to a higher volume of distribution)

 $\rightarrow$  JNJ-Y seemed to be most valuable with regard to a multiple dosing regimen.



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# Clinical: eliminate unnecessary trial

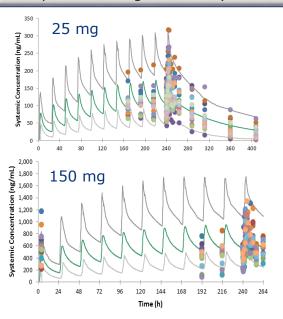
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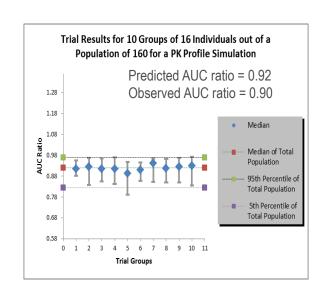
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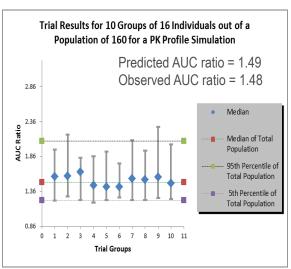
Multiple ascending dose study

50mg sildenafil + 75mg qd rilpivirine

150mg qd rilpivirine+400mg qd keto







#### **VERFICATION RILPIVIRINE MODEL**

Verification of CYP3A4 contribution?

- Clinical study data with CYP3A4 inhibitor ketoconazole
- Multiple ascending dose study (auto-induction)

Verification of inhibition and induction effects on CYP3A4?

- Clinical study data with CYP3A4 substrate sildenafil



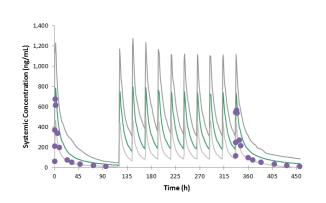
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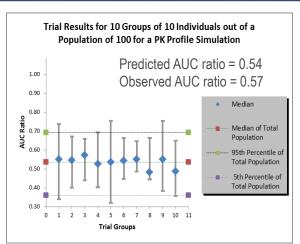
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- → dose of rilpivirine x 2 to overcome the decrease in exposure upon coadministr.
- → what to do after stopping rifabutin? (when rifabutin is stopped an ongoing inducing is expected for a few more days)

Multiple ascending dose study

25mg rilpivirine + 300mg qd rifabutin





#### **VERIFICATION RIFABUTIN MODEL**

Verification of induction effects on CYP3A4?

Clinical study data with rifabutin

Verification of inducing concentrations at steady state?

- Multiple ascending dose study



#### **Abstract**

The transition from preclinical development to clinical studies is a major milestone in a drug development project. During this transition the selection of the most suitable starting dose and dosing regimen of the first-in-man study is one of the primary expectations. The starting dose and dosing regimen should be sufficiently low to avoid toxicity (i.e. safety related risk) and sufficiently high to avoid unnecessary human trials (i.e. efficacy related risk). A consolidation of all available preclinical information is essential to provide estimates for these doses. Several approaches are being used in this regard, from very simple (minimum anticipated biological effect level (MABEL), allometry, Dedrick approach) to more complex (physiology based PK modeling (PBPK)). Unlike the simple approaches, PBPK modelling integrates preclinical in vitro and in vivo data into a mechanistic framework to predict plasma and tissue concentration-time profiles, and to address mechanistic issues that are relevant to pharmacokinetics (e.g. saturable metabolism and/or absorption, interspecies differences, expression levels and activity of metabolizing enzymes and transporters). Such mechanistic models result in a prepared mind once the first human volunteer/patient is being dosed. Several case examples are shown where mechanistic models helped mitigating potential issues (compound selection during drug discovery, setting the dose, the escalation steps, and the regimen of first-in-man studies (thereby taking into consideration saturation and time-dependent processes), and setting the dose for drug-drug interaction studies).

