

Open-Source and Open-Science to Progress the Integration of Pharmacometrics and Systems Pharmacology

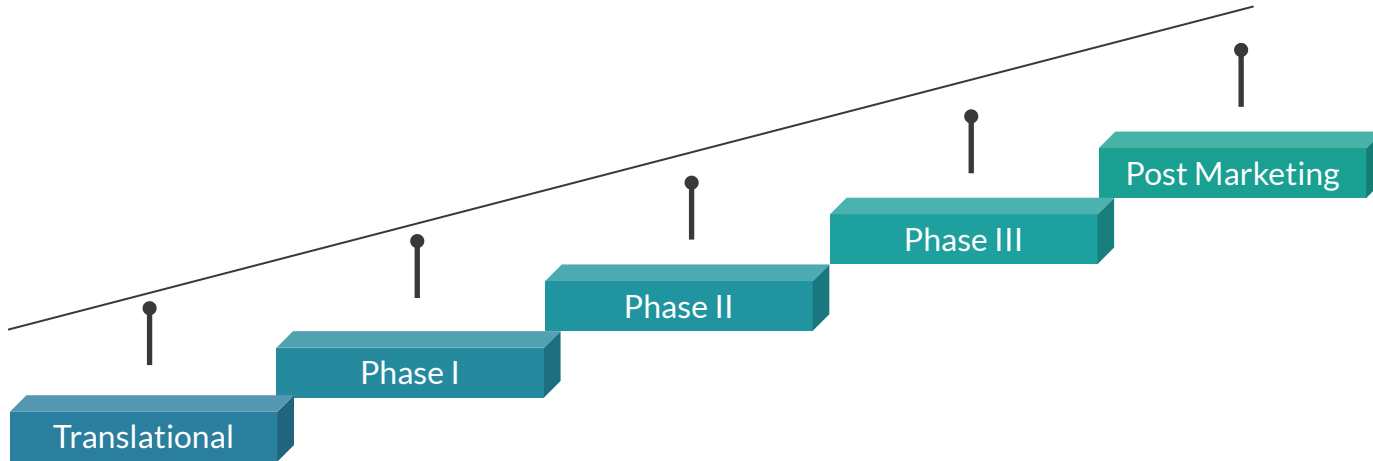
Matthew Riggs, Ph.D.
Chief Science Officer

Metrum Research Group

Monday, 26 August 2019

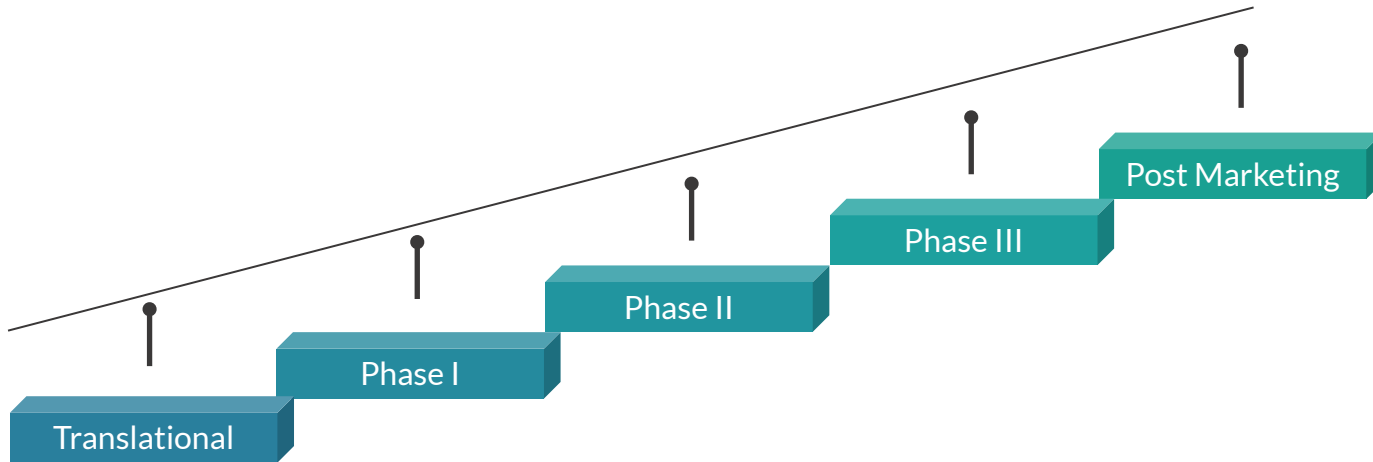
Drug Development

Starts as an idea ... becomes a medicine (if it deserves to be)



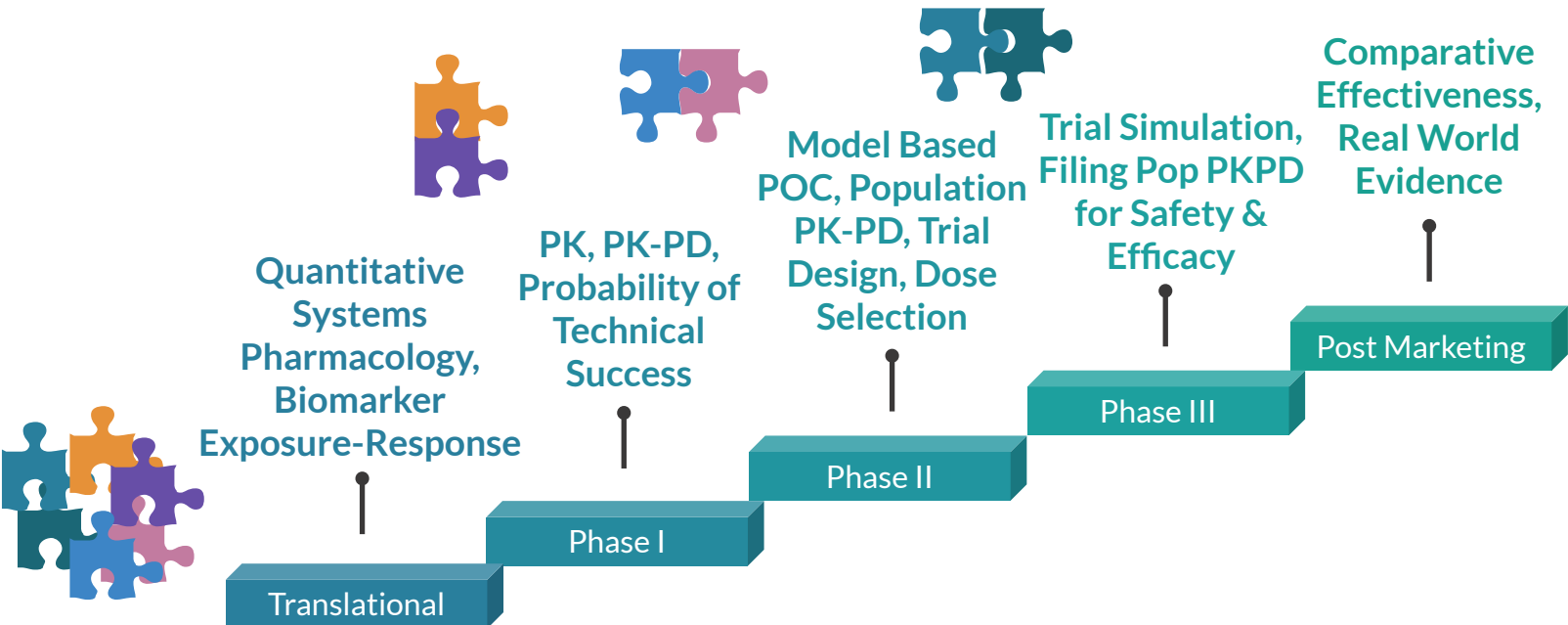
Drug Development

The idea though can be a puzzle at the start



Model-Informed Drug Development

Modeling and Simulation: inform decision support as pieces are fit together

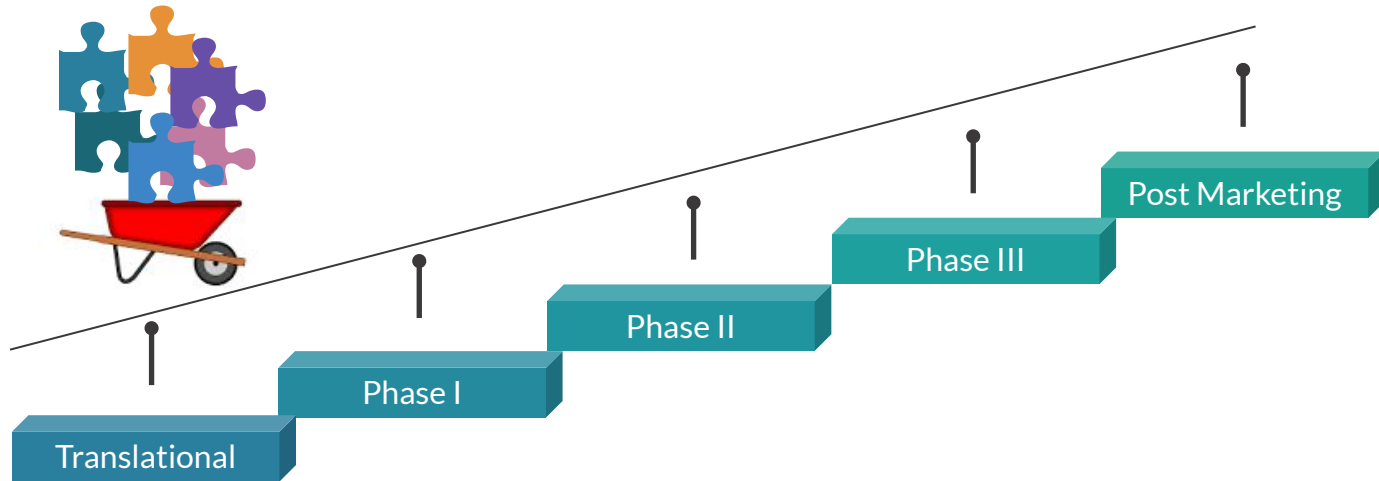


Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

Pretend

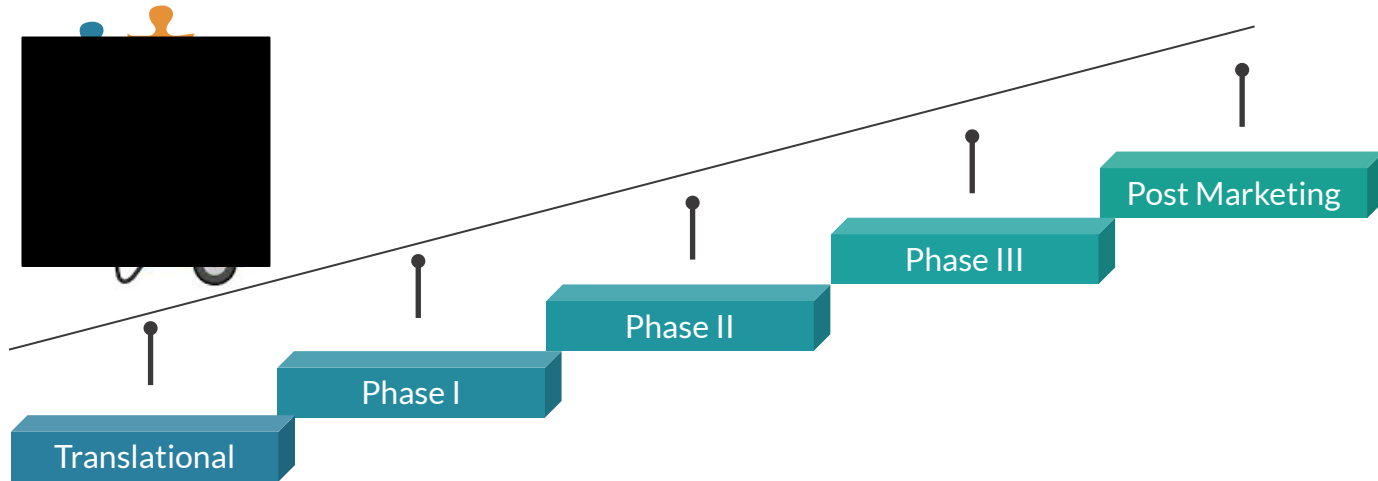
Scenario 1: *Company A* develops a model that helps to carry the pieces more easily through development. They called this a “one-compartment barrow”, and boast at ACoP10 that it accelerated development into Phase 1 ...



Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

Scenario 1: ... but they hide the model. Because how else can Company A make money off the drug?



Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

Scenario 1: ... the repercussions: Company B, C, D, ... Z see the fancy presentation at ACoP10 and want a “one-compartment barrow”, too.

From bootlegged photos, they go to work...



“It looked like it had a sled arch on the bottom with some kind of hook; a labmate did a model like that in grad school. I can work with my adviser and we’ll have it recreated in no time; shouldn’t take more than 2 weeks” Company X

“Let’s form an internal working group...” Company H

“There’s this private-public consortium, we give them all of our data, then they develop a model that they’ll sell back later as a black box” Companies O, M, Y

“Pfft, I’ll use machine learning; it can solve anything, even without data...” Company G

Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

Scenario 1: ... 6 months later....

We need to show management some progress!



“I know but with the new semester about to start and 3 more NIH grant applications, it’s slow but they are making progress (I think?)...” Company X

“Our working group has reached out and engaged key stakeholders, leveraged synergies, and is ensuring with resource utilization that the modeling will be impactful ...” Company H

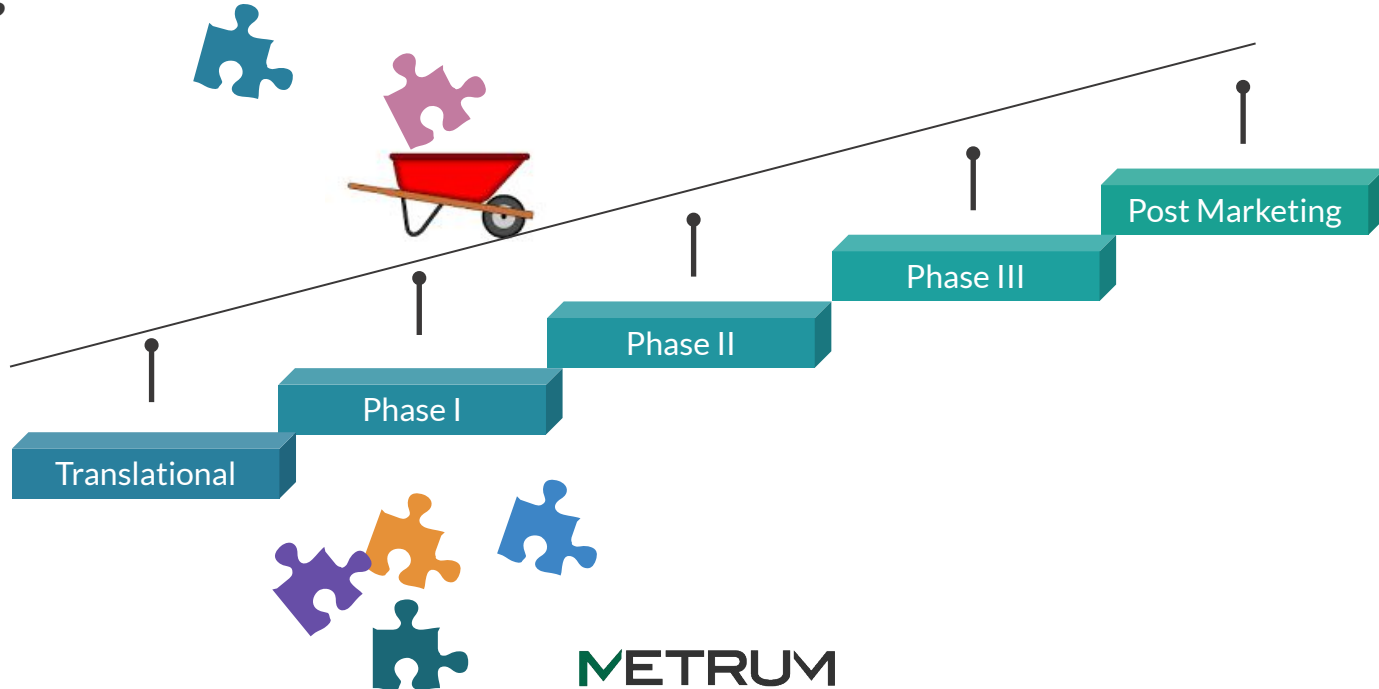
“Legal only let us share summary level placebo data and the goals alignment is, well, complicated... but, we’ve trademarked the term ‘OC barrow’ and assured management that we’re streamlining efficiencies.” Companies O, M, Y

“I’ve trained my AI algorithm to perfectly fit both data points...” Company G

Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

Scenario 1: ... meanwhile, back at Company A... “well, we didn’t see that (AE) coming”



Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

Scenario 1: ... Company A... “we’re exiting this therapeutic area, close everything up and send it to archive...”

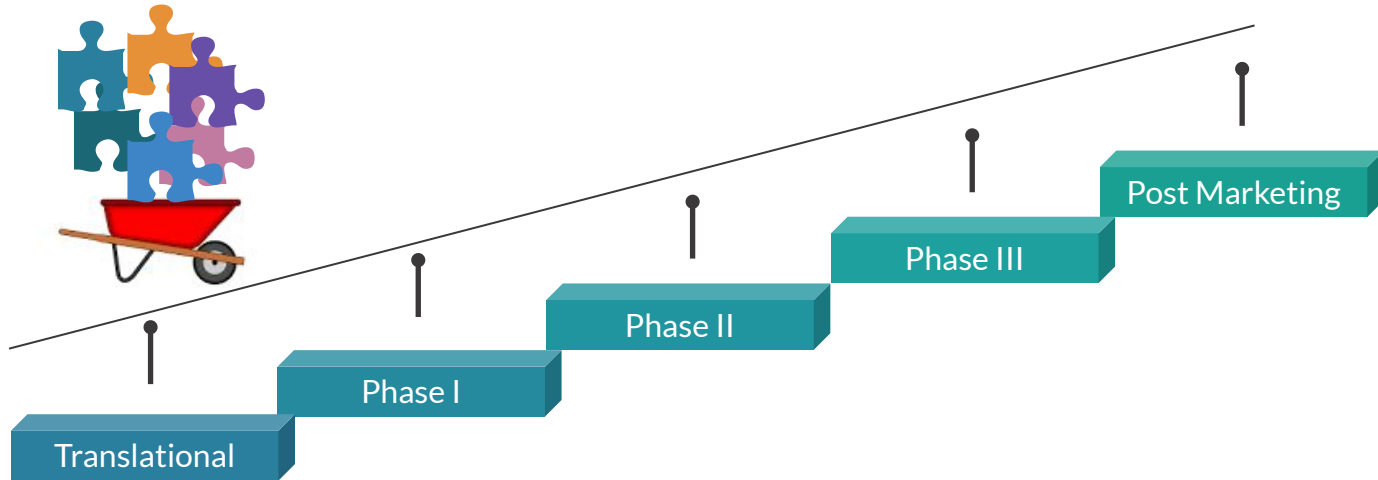


Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

Pretend

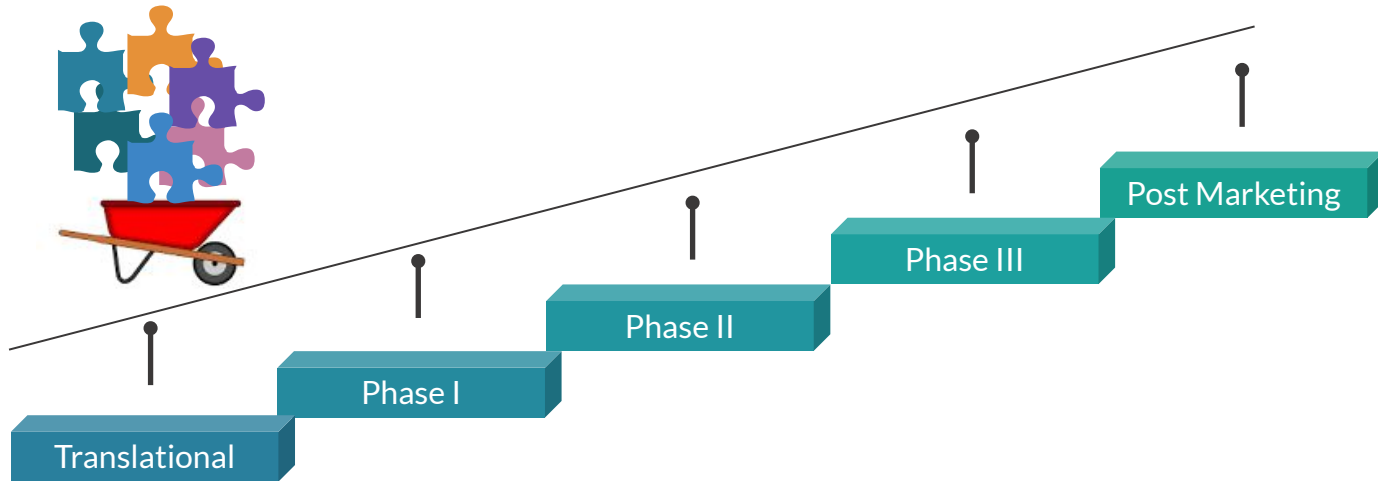
Scenario 2: *Company A* develops a model that helps to carry the pieces more easily through development. They called this a “one-compartment barrow”, and boast at ACoP10 that it accelerated development into Phase 1 ...



Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

Scenario 2: ... and they know the model could be made even better, so they share it on (DDMoRe, github, biomodels, ...). Because how better can *Company A* multiply their progress from program to program?



Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

Scenario 2: ... the advancement: Company B, C, D, ... Z see the fancy presentation at ACoP10 and want a “one-compartment barrow”, too.

From the open-source code, they go to work...



“If we add a second wheel up front it will be less likely to tip over...” Company X

“Let’s build and add a Shiny app to help visualize the results and path forward...”
Company H

“You know, if we share ideas: what do you think about 4 wheels and an engine...”
Company O, M, Y

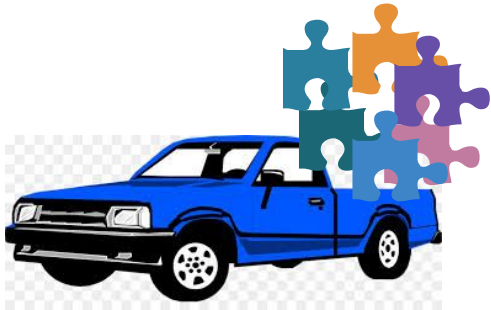
“And we could add an algorithm for better fitting and then use those new verification and validation standards to show its cross-application performance...” Company G

Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

Scenario 2: ... 6 months later....

We can't wait to show management the progress!



“This is great, everyone on the team is playing around with the model. That steering wheel Shiny app that H added was great ...” Company X

“We’ve identified other therapeutic areas that this will work for, too ...”
Company H

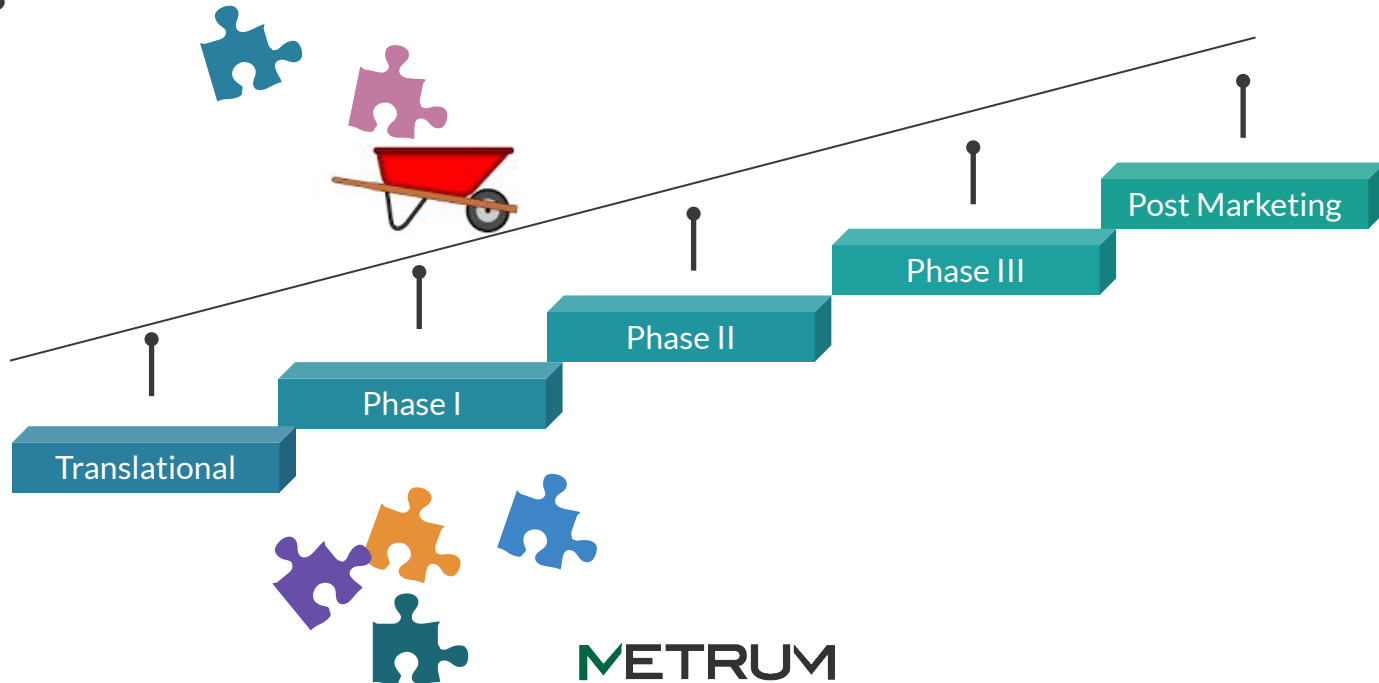
“That VVUQ stuff sounded too complicated but now that the whole community is using the model, we’re more confident that (investors, regulators, payers) will be asking for the results, too ...” Company O, M, Y

“Oh, that “enhanced barrow”, as you’re calling it now, is going to like a clunky old pickup truck when you see what we’re about to add next ...” Company G

Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

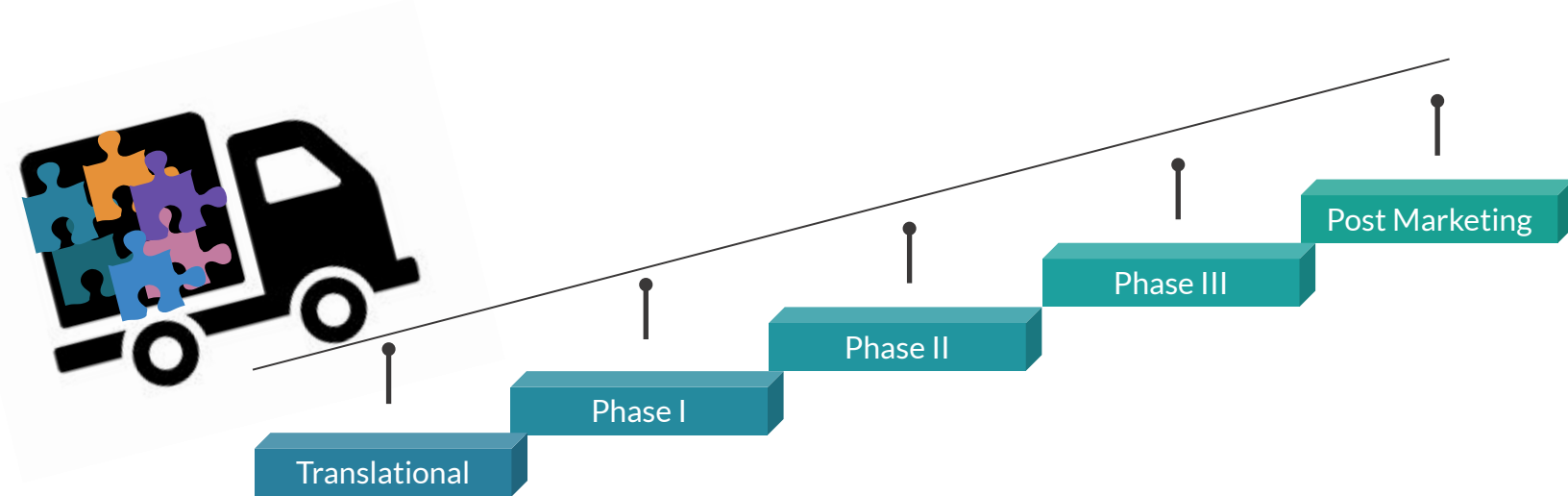
Scenario 2: ... meanwhile, back at Company A... “well, we didn’t see that (AE) coming”



Model-Informed Drug Development

Modeling and Simulation: how do we make our models all the better?

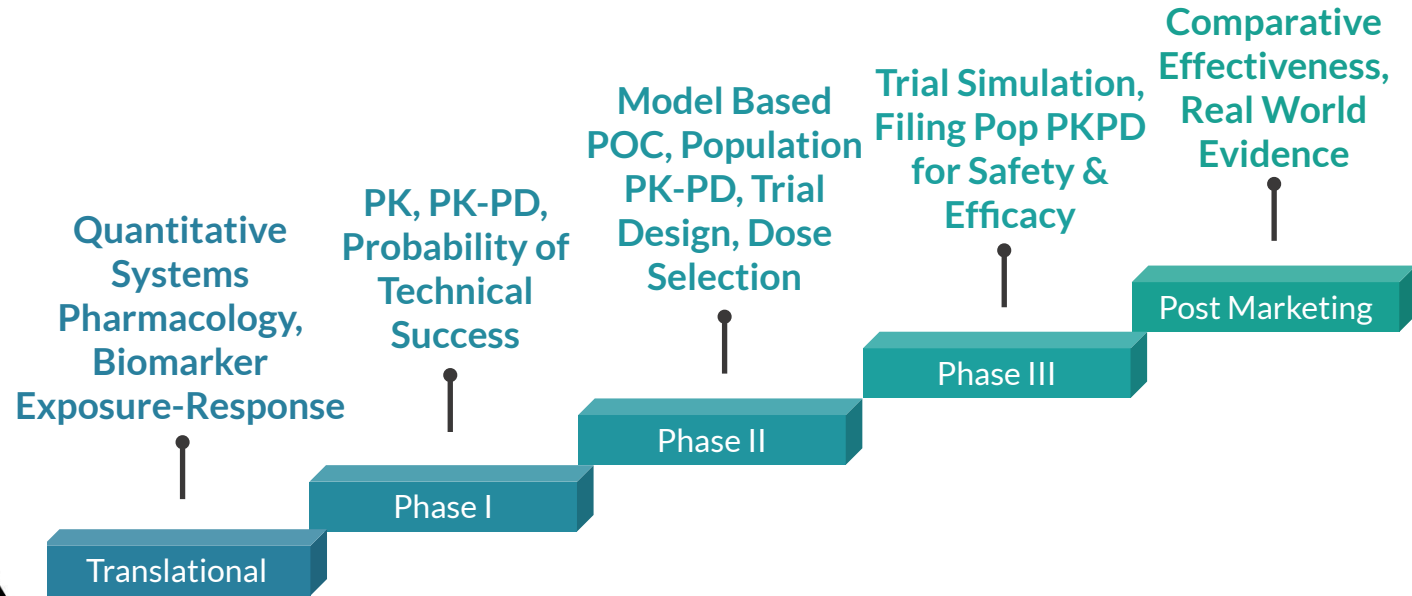
Scenario 2:... Companies A-Z... “we may be exiting this therapeutic, though with improved model-informed decision support we’re so much more confident in these next development programs...”



Open Model-Informed Drug Development

Ok sure, it's idealistic, though ... how could it really work?

Open-Source and Open-Science to Progress the Integration of Pharmacometrics and Systems Pharmacology



Open Model-Informed Drug Development

Examples: integrated Pharmacometrics and Systems Pharmacology (iPSP)

COMMENTARY

Mineral and Bone
Health Model

Cardio-Renal
Model

ERK Model

OSPSuite

When Learn/Confirm Leads to Expand/Understand: The Expanding Role of Quantitative Systems Pharmacology in the Betterment of Therapeutics Development

Matthew M. Riggs¹

Our earnest ambitions to advance therapeutics for patients worldwide often begin by asking two questions: "Is the treatment safe?" and "Is the treatment efficacious?" The next question, however, is can we do even better? Moving beyond "Is the treatment ...?" to "Why is the treatment ...?" and "How can we use that information to make the treatment even better?" We will examine how learn/confirm leads to expand/understand with the expanding role of quantitative systems pharmacology.

Our earnest ambitions to advance therapeutics for patients worldwide often begin by asking two questions: "Is the treatment safe?" and "Is the treatment efficacious?" We design programs to answer those questions that anticipate and answer challenges that are often far more complicated than those seemingly simple questions may imply. Development designs are streamlined for efficiency while gathering as much data as possible to support approval of the new medication as safe relative to measurable improvement in patient outcome. Progress from our efforts has been remarkable. We have achieved sustained response for

previously deadly viral infection,¹ provided platforms assessing survival across cancer types,² and offered new hopes toward curing rare diseases,³ to name a few.

The next question, however, is can we do even better? Should we continue challenging ourselves beyond "Is the treatment ...?" to "Why is the treatment ...?" and "How can we use that information to make the treatment even better?" We are given a resounding yes to the question of "Can we do better?" in the report by Kharana *et al.*⁴ on the "Use of a Systems Pharmacology Model Based Approach Towards Dose Optimization of Parathyroid Hormone

Therapy in Hypoparathyroidism." That report provides a teaching example of how to conduct and position quantitative systems pharmacology (QSP) modeling that resonates whether in discovery or regulatory review, namely: (i) understand the unanswered question(s), (ii) use a validated-for-purpose model to challenge alternative hypotheses, and (iii) take those expanded learnings to progress therapeutics.

To begin, the authors extend an existing, open-source, multicalc QSP model⁵ as their foundation. The model was first extended to include dosing of exogenously administered parathyroid hormone (PTH) and a compartment capturing urinary calcium output. The model was then validated for purpose using sponsor phase I data. The investigators used this *in silico* platform to explore the mechanisms by which the intended treatment provides effectiveness, while seeking an optimum kinetic balance of physiologic feedbacks to minimize an untoward effect in the kidneys (hypocalcemia).

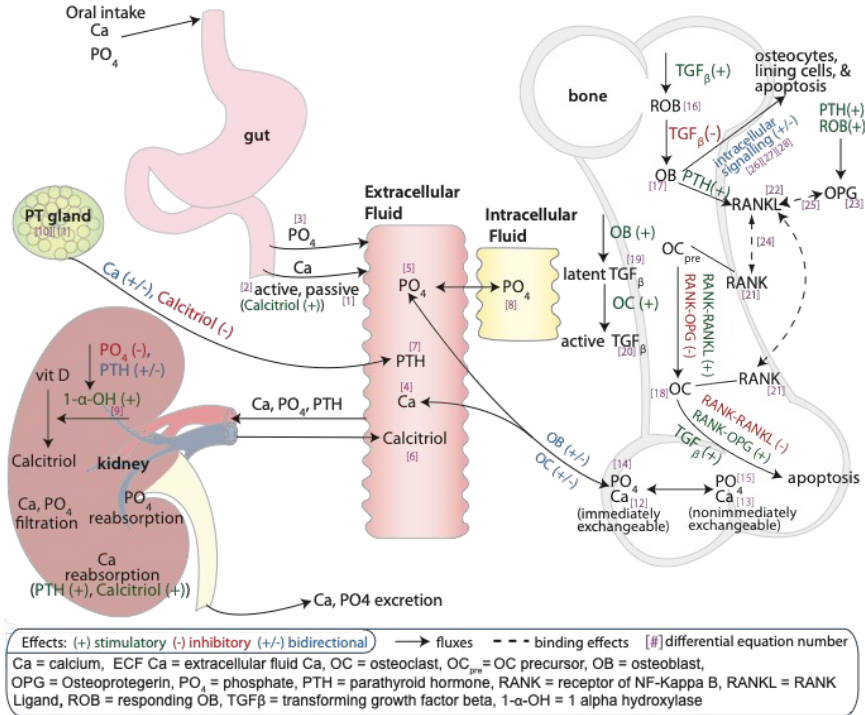
The details of how Kharana *et al.*⁴ progressed to this point are important. First, though, let us further consider the notion of continuing beyond "Is the treatment ...?" to "Can we make it even better?" In the absence of sheer chance, this requires clear understanding of the mechanisms, or what we can consider as physiologic and pharmacologic rules, by which the treatment is operating. In his seminal 1997 commentary, Dr Lewis Sheiner begins with: "George Box" views scientific progress as consisting of, and requiring, alternating steps of induction and deduction: the former being learning from experience, and the latter being confirmation of what has been learned." He continues "The understandable focus of commercial drug development on confirmation, as this immediately

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Received September 27, 2018; accepted November 2, 2018; advance online publication Month 00, 2018. doi:10.1002/cpt.1287



Mineral and Bone Health Model



Peterson and Riggs. Bone 2010

Open Model-Informed Drug Development

Promoting Open-Source, Open Science Community Involvement on GitHub:

<https://github.com/metrumresearchgroup/OpenBoneMin>

Mineral and Bone
Health Model

Code: Open-Source R
with *mrgsolve*

GitHub, Inc. [US] | github.com/metrumresearchgroup/OpenBoneMin 🔍 ☆

About

A multiscale systems model of bone health and mineral homeostasis. Please see the [wiki page](#) for more information on this project.

Community contributions to this project are included [here](#).

Documentation

- Documentation [here](#)

Installation

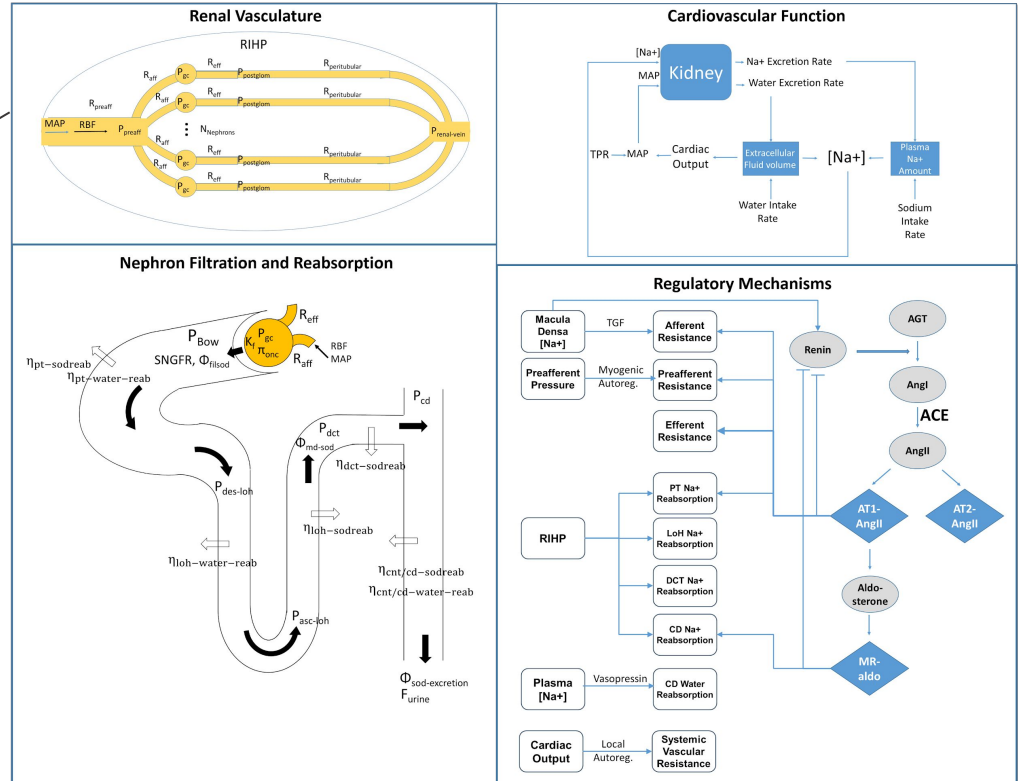
Installation of `OpenBoneMin` requires the `devtools` package

```
if(!require("devtools")) install.packages("devtools")
```

Use the `install_github` function inside `devtools` to install the `OpenBoneMin` package from GitHub to your local machine

```
devtools::install_github("metrumresearchgroup/OpenBoneMin")
```

Cardio-Renal Model



Source: <https://github.com/hallowkm/RenalModel/blob/master/Schematic.png>

Open Model-Informed Drug Development

Promoting Open-Source, Open Science Community Involvement on GitHub:

<https://github.com/hallowkm/RenalModel>

Cardio-Renal Model
Code: Open-Source R with RxODE/nlmixR
Shiny app:
<http://qsp.engr.uga.edu:3838/CVRmod/>

hallowkm / RenalModel

Code Issues 1 Pull requests 1 Projects 0 Wiki Security Insights

A quantitative systems physiology model of renal function and blood pressure regulation

2 commits 2 branches 0 releases 1

Branch: master New pull request Create new file Upload files Find File

hallowkm added schematic Latest commit a5

README.md	Initial commit
Schematic.png	added schematic

README.md

RenalModel

A quantitative systems physiology model of renal function and blood pressure regulation

Cardio-Renal Model

Integrated PK/PD
model for direct renin
inhibitor: imarikiren

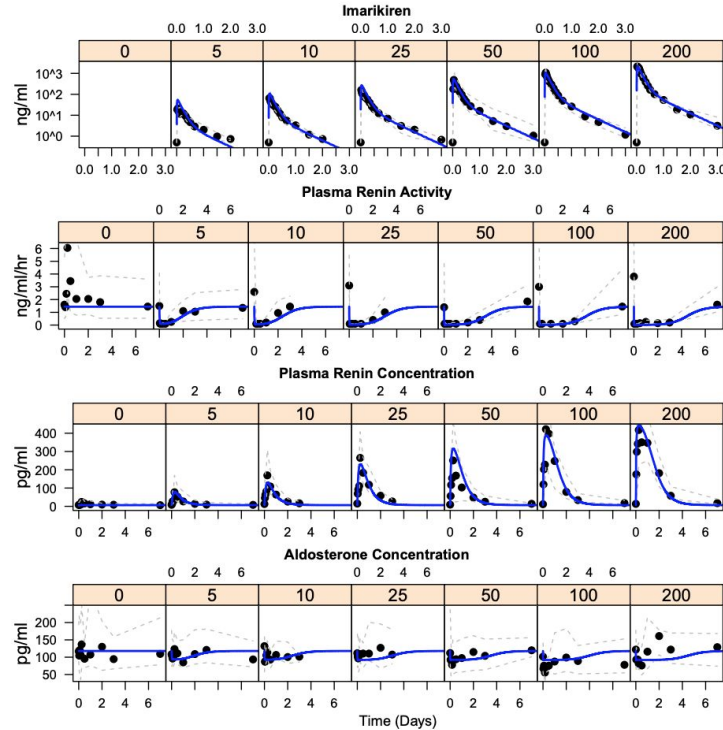
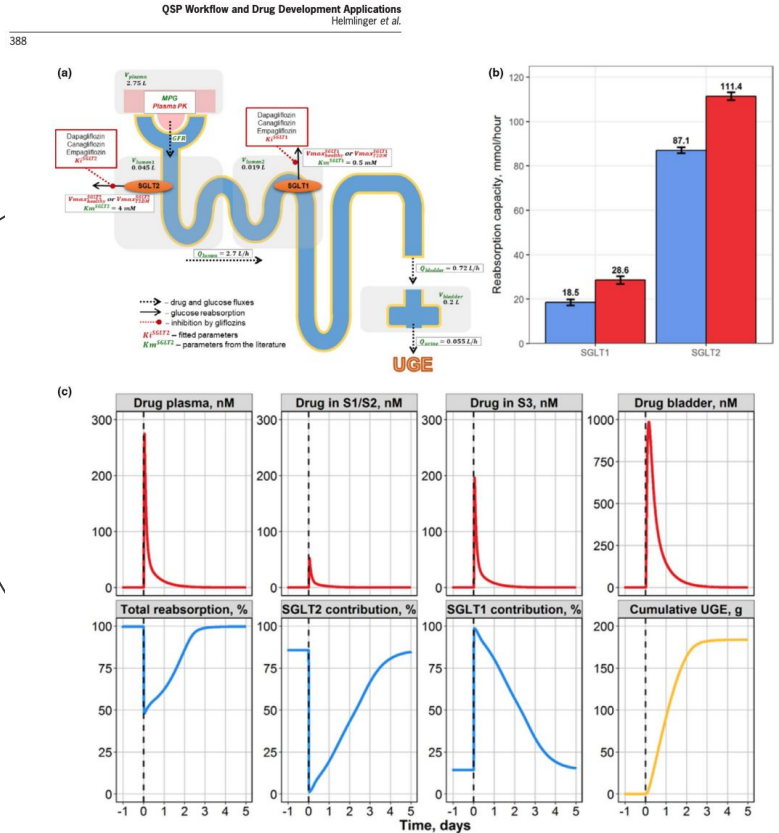


Fig. 2 Plasma concentration–time profile of imarikiren for doses ranging from 5 to 200 mg (top row). Pharmacodynamics response of renin inhibition as shown by PRA (second row), PRC (third row), and aldosterone (fourth row). Panel labels indicate the dose in mg. Solid

blue curves represent model fit for the typical individual and filled circles represent clinical data. The dashed gray curves in the figure represent the 5 and 95% quantiles of the clinical data (Color figure online)

Gebremichael, Y., Lahu, G., Vakilynejad, M., & Hallow, K. M. (2019). J Pharmacokinet Pharmacodyn, 46(1), 15–25.

Cardio-Renal Model
Integrated PK/PD
model for SGLT
inhibition



Brady, J. A., & Hallow, K. M. (2017). Model-Based Evaluation of Proximal Sodium Reabsorption Through SGLT2 in Health and Diabetes and the Effect of Inhibition With Canagliflozin. *J Clin Pharmacol*.

Hallow, K. M., Greasley, P. J., Helmlinger, G., Chu, L., Heerspink, H. J., & Boulton, D. W. (2018). Evaluation of renal and cardiovascular protection mechanisms of SGLT2 inhibitors: model-based analysis of clinical data. *Am J Phys Ren Phys*, 315(5), F1295–F1306.

Hallow, K. M., Helmlinger, G., Greasley, P. J., McMurray, J. J. V., & Boulton, D. W. (2018). Why do SGLT2 inhibitors reduce heart failure hospitalization? A differential volume regulation hypothesis. *Diabetes, Obesity & Metabolism*, 20(3), 479–487.

Helmlinger, G., Sokolov, V., Peskov, K., Hallow, K. M., Kosinsky, Y., Voronova, V., ... Penland, R. C. (2019). Quantitative Systems Pharmacology: An Exemplar Model-Building Workflow With Applications in Cardiovascular, Metabolic, and Oncology Drug Development. *CPT: PSP*, 8(6), 380–395.





PUBLISHED
20 August 2019

Cardio-Renal Model

Integrated PK/PD
model for SGLT
inhibition

This announcement contains inside information

20 August 2019 07:00 BST

DAPA-HF is the first heart failure outcomes trial with an SGLT2 inhibitor in patients with and without type-2 diabetes

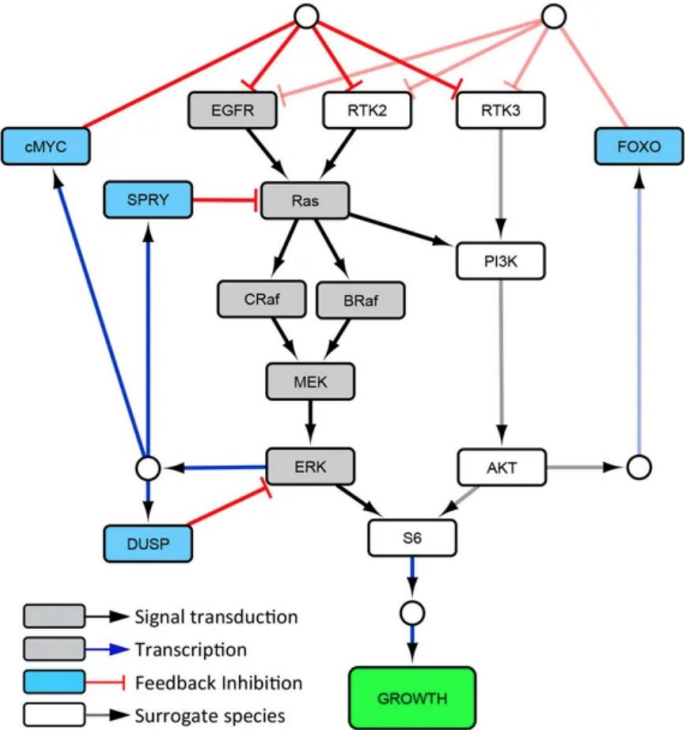
Farxiga significantly reduced the risk of cardiovascular death or worsening of heart failure when added to standard of care

AstraZeneca today announced positive results from the landmark Phase III DAPA-HF trial which showed that *Farxiga* (dapagliflozin) met the primary composite endpoint with a statistically-significant and clinically-meaningful reduction of cardiovascular death or the worsening of heart failure (defined as hospitalisation or an urgent heart failure visit), compared to placebo. The trial was conducted in patients with reduced ejection fraction (HFrEF) on standard of care treatment, including those with and without type-2 diabetes.

The safety profile of *Farxiga* in the DAPA-HF trial was consistent with the well-established safety profile of the medicine.



ERK Model



Kirouac, D. C., Schaefer, G., Chan, J., Merchant, M., Orr, C., Huang, S.-M. A., ... Ramanujan, S. (2017). *NPJ Systems Biology and Applications*, 3, 14.

Source: <https://www.nature.com/articles/s41540-017-0016-1>

Open iPSP-Informed Drug Development

Promoting Open-Source, Open Science through Supplementary Materials (and GitHub)

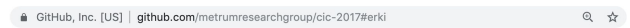
ERK Model

Code: MatLab files in Supplement

Conversion to R/mrgsolve on GitHub



- Figure S1
- Figure S2
- Figure S3
- Figure S4
- Figure S5
- HillEQ.m
- VPop_Simulator_fixed.m
- Supplementary Tables S1-S11
- MAPK_model.sbioproj
- Supplementary Materials
- Supplementary Materials



Clinical responses to ERK inhibition in BRAF(V600E)-mutant colorectal cancer predicted using a computational model

- Daniel C. Kirouac, Gabriele Schaefer, Jocelyn Chan, Mark Merchant, Christine Orr, Shih-Min A. Huang, John Moffat, Lichuan Liu, Kapil Gadkar and Saroja Ramanujan
- *npj Systems Biology and Applications* (2017) 3:14 ; doi:10.1038/s41540-017-0016-1
- Model, Example, R script, Generate figure 6b

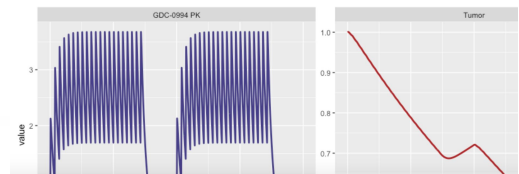
ERK inhibition in CRC (Kirouac et al.) [code]

```
mod <- mread("mapk", "content/model")
vp <- read.csv("content/data/s18vpop.csv", header=TRUE) %>% sLice(1)
mod <- param(mod,vp) %>% inIt(vp)
```

Simulate two cycles of GDC-8994 dosing

```
data <- expand.ev(amt=400, cmt=12, time=c(0,28), i1=1, addl=28) %>% mutate(ID=1)
out <- mrgsim(mod, data=data, end=56, delta=0.1, Req="GDC,TUMOR")
```

Plot (code not shown)

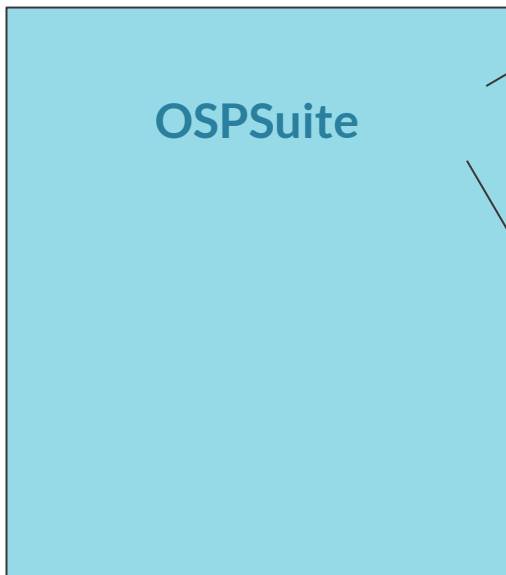


Kirouac, D. C., Schaefer, G., Chan, J., Merchant, M., Orr, C., Huang, S.-M. A., ... Ramanujan, S. (2017). *NPJ Systems Biology and Applications*, 3, 14.

Open Model-Informed Drug Development

Promoting Open-Source, Open Science Community Involvement on GitHub:

<https://github.com/Open-Systems-Pharmacology>



Open Systems Pharmacology
Latest suite release can be found here:
<http://setup.open-systems-pharmacology.org>

Repositories 116 Packages People 46 Teams 3 Projects 3

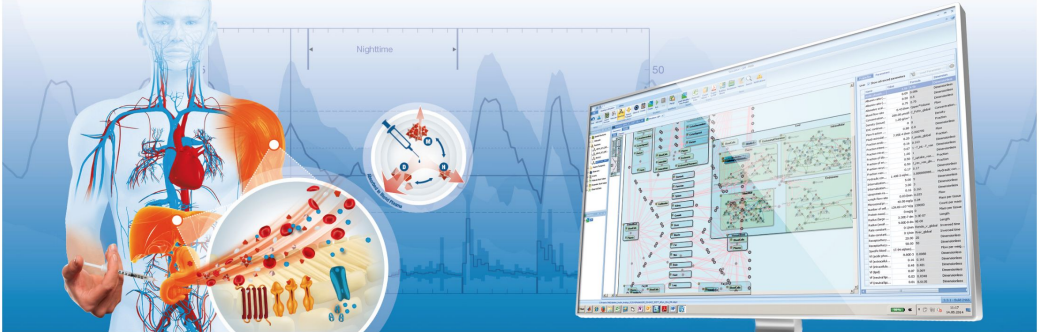
Pinned repositories

- Forum**
Discussion forum for the Open Systems Pharmacology Project
★ 24 🍴 6
- Suite**
Open Systems Pharmacology Suite Setup
Rich Text Format ★ 50 🍴 15
- PK-Sim**
PK-Sim® is a comprehensive software tool for whole-body physiologically based pharmacokinetic modeling
● C# ★ 33 🍴 22
- MoBi**
MoBi® is a software tool for multiscale physiological modeling and simulation
● C# ★ 13 🍴 4
- ACoP-10-PBPK-Workshop-2019**
PBPK Workshop at ACoP 10 2019
- Vision-Mission**
Vision & Mission of Open Systems Pharmacology
★ 2 🍴 1

OSPSuite
PBPK Models
(PK-Sim)
Bio Models (MoBi)

GitHub, Inc. [US] | github.com/Open-Systems-Pharmacology/Glucose-Insulin-Model

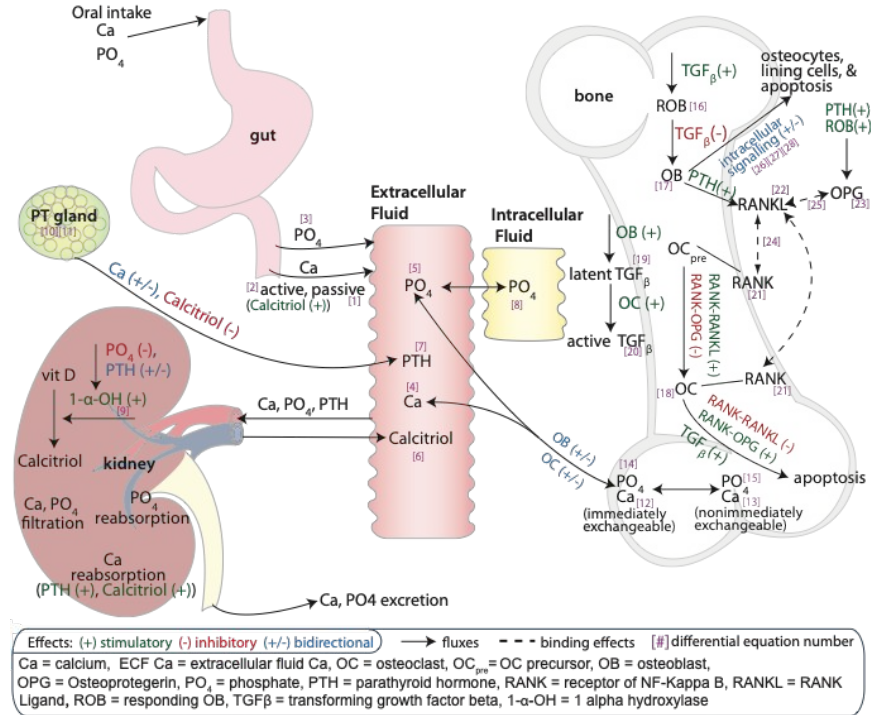
Welcome to the Glucose Insulin model



The physiologically-based whole-body model of the glucose-insulin-glucagon regulatory system

Within this repository, we distribute the physiologically-based whole-body model of glucose-insulin-glucagon regulation based

Taking a Deeper Dive
 How did this model come to be?
 How has it expanded?
 How did it come back years later (with extra wheels) to help Sponsor again?

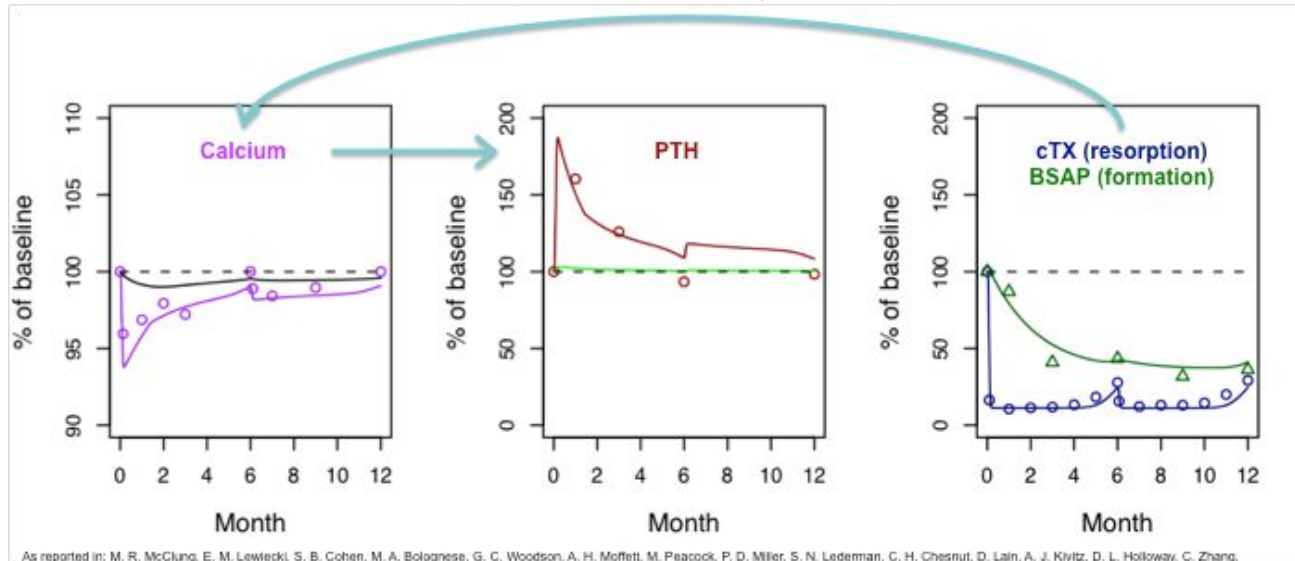


Peterson and Riggs. Bone 2010

The question: Can we better understand (and predict) the inter-rational effects of dmab and the time-courses of on/off t effects?

Denosumab: RANKL inhibition

- ↓ Bone Resorption
- Can we better understand the other changes (e.g., bone formation marker, serum calcium and PTH)? Should we be concerned?

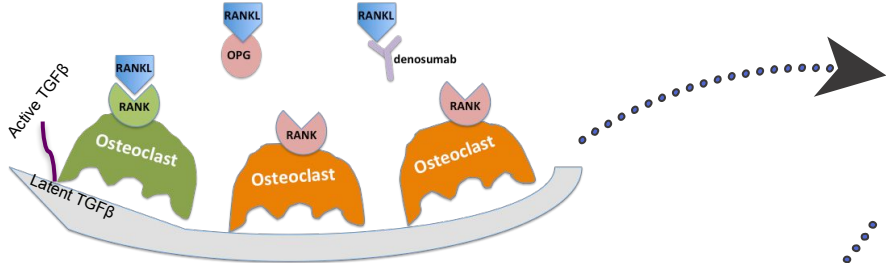


As reported in: M. R. McClung, E. M. Lewiecki, S. B. Cohen, M. A. Bolognese, G. C. Woodson, A. H. Moffett, M. Peacock, P. D. Miller, S. N. Lederman, C. H. Chesnut, D. Laine, A. J. Kivitz, D. L. Holloway, C. Zhang, M. C. Peterson, P. J. Bekker, and AMG 162 Bone Loss Study Group. Denosumab in postmenopausal women with low bone mineral density. *N Engl J Med*, 354(8):821-31, Feb 2006.

iPSP: A Case Study

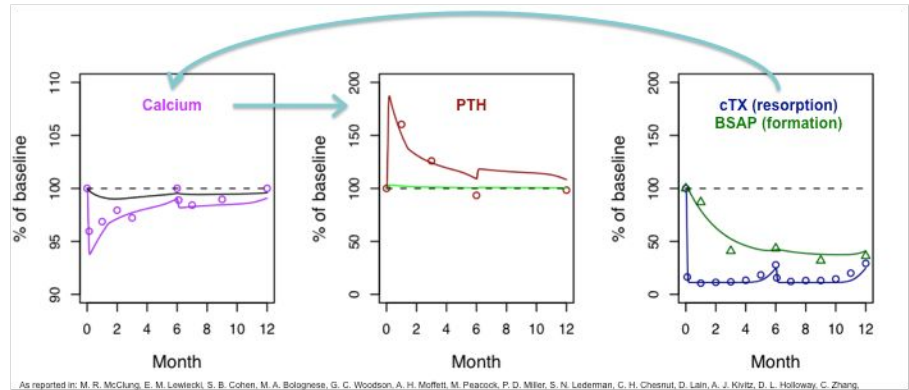
The question: Can we better understand (and predict) the inter-rational effects of dmab and the time-courses of on/off t effects?

Denosumab: RANKL inhibition



- ↓ available RANKL
- ↓ RANK--RANKL interaction
- ↓ Osteoclast activity (sCTx)
- ↓ Activation of TGF-β
- ↓ Osteoblast activity (BSAP)
- ↑ bone mineral density (BMD)

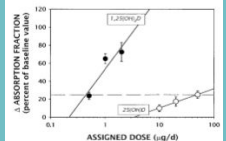
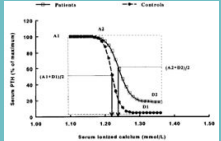
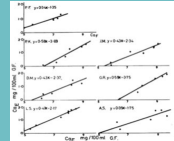
- ↓ Calcium release from bone
- ↓ Serum calcium
- ↓ Ca sensing in PT gland
- ↑ PTH release (calcium-sparing)



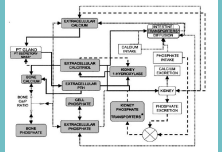
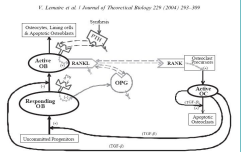
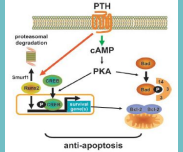
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iPSP: Integrate Existing Data & Models

Literature and in-house information is often available to inform model parameter and disease state effects

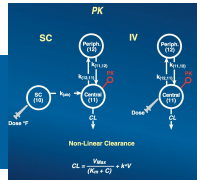
Calcium Absorption	PTH Secretion	Calcium Excretion	Bone Therapeutics	Disease States
			Anabolic (<i>teriparatide, 2004</i>) Catabolic (<i>denosumab, 2006</i>)	Hyper- and hypo-PTH CKD-MBD (<i>Rix et al. 1999</i>)
e.g., Heaney et al. 1997	e.g., Ramirez et al. 1993	e.g., Peacock and Nordin 1968		



Calcium Homeostasis	Bone Remodeling	Intracellular Signaling
		
e.g., Raposo et al. 2002	e.g., LeMaire et al. 2004	e.g., Bellido et al. 2003

A Population PK/PD Model Describes the Rapid, and Sustained Suppression of Urinary N-telopeptide Following Administration of AMG 162, a Fully Human Monoclonal Antibody Against RANKL, to Healthy Postmenopausal Women

M. C. Peterson¹, B. J. Stouch¹, D. Chen¹, S. Baughman¹, D. L. Holloway², P. J. Bekker², S. W. Martin¹
¹Pharmacokinetics and Drug Metabolism, ²Clinical Research, Amgen Inc., Thousand Oaks, CA, USA



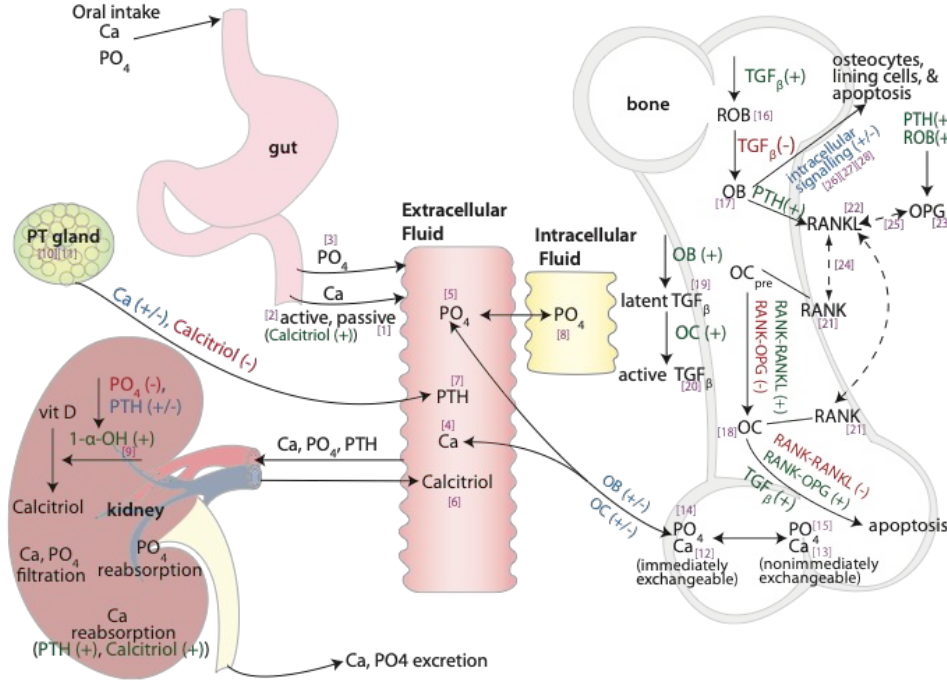
Multiscale Model

Peterson MC and Riggs MM (2010) A physiologically based mathematical model of integrated calcium homeostasis and bone remodeling. *Bone* 46:49-63.



iPSP: A Case Study

The question: Can we better understand (and predict) the inter-rational effects of dmab and the time-courses of on/off effects?



Effects: (+) stimulatory (-) inhibitory (+/-) bidirectional → fluxes --- binding effects [#] differential equation number
 Ca = calcium, ECF Ca = extracellular fluid Ca, OC = osteoclast, OC_{pre} = OC precursor, OB = osteoblast, OPG = Osteoprotegerin, PO₄ = phosphate, PTH = parathyroid hormone, RANK = receptor of NF-Kappa B, RANKL = RANK Ligand, ROB = responding OB, TGF_β = transforming growth factor beta, 1-α-OH = 1 alpha hydroxylase

Multiscale Model

Peterson MC and Riggs MM (2010) A physiologically based mathematical model of integrated calcium homeostasis and bone remodeling. Bone 46:49-63.



iPSP: On/Off Treatment Effects

Denosumab: RANKL inhibition → Bone Marker Changes

Dose-Ranging: 6 → 210 mg, Q3M and Q6M, d/c, re-Tx

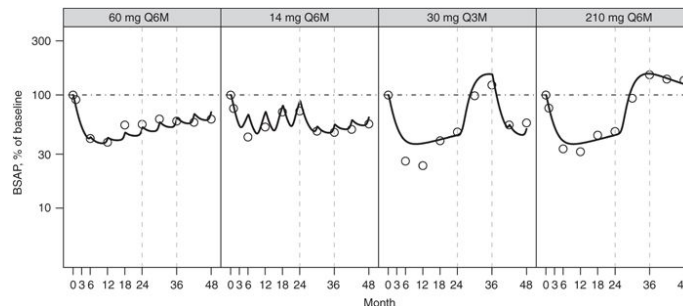
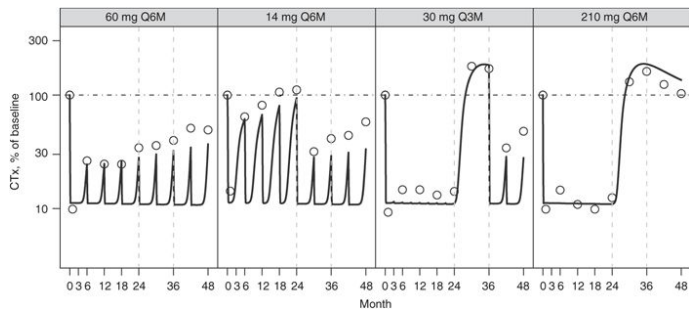
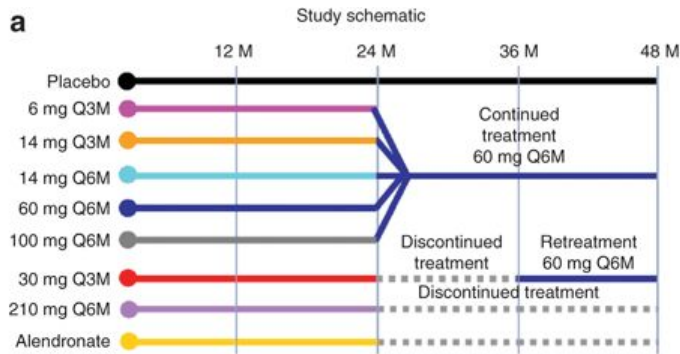


Fig.3 and 4; Peterson MC and Riggs MM. CPT: Pharmacometrics & Systems Pharmacology (2012) 1, e14; doi:10.1038/psp.2012.15

iPSP: Expand to Predict Clinical Outcome

Denosumab: RANKL inhibition → Bone Markers → BMD Change

Dose-Ranging: 6 → 210 mg, Q3M and Q6M, d/c, re-Tx

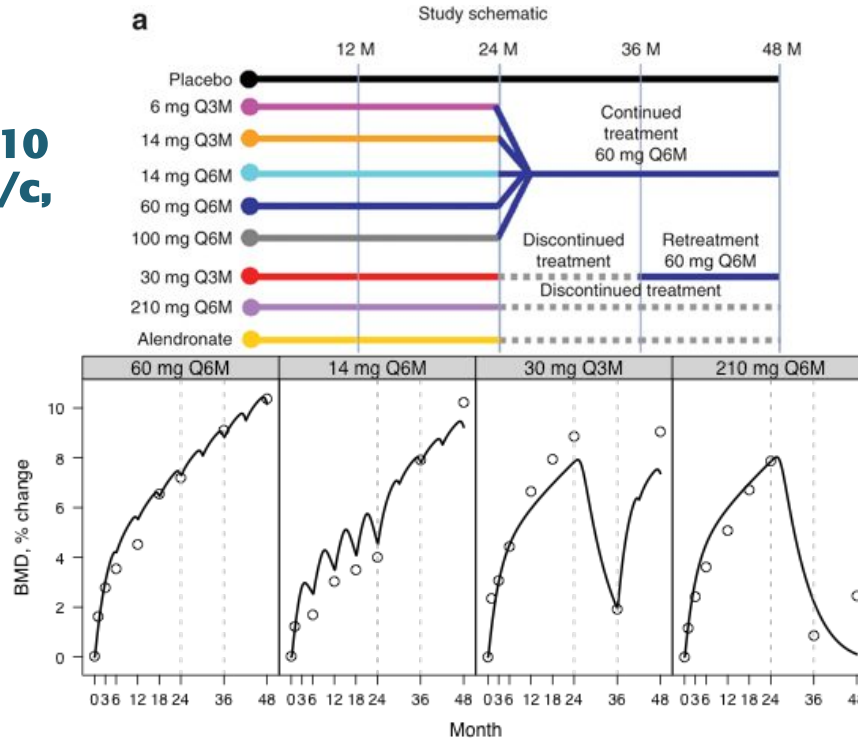


Fig. 5; Peterson MC and Riggs MM, CPT: Pharmacometrics & Systems Pharmacology (2012) 1, e14; doi:10.1038/psp.2012.15

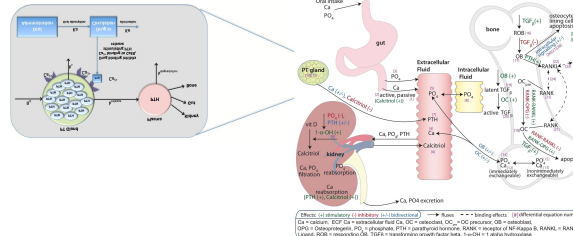
Open iPSP Calcilytic, target for osteoporosis?

Question: Were prior results a “class effect” or was this still a viable target?

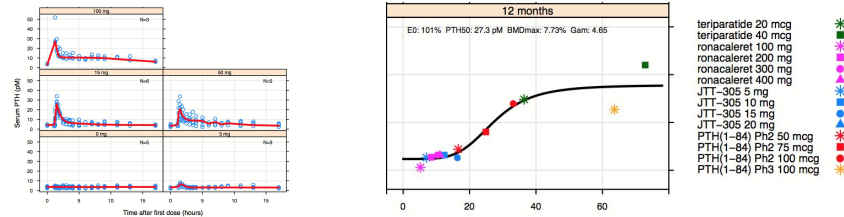


Evaluate prior tests' persistent hyperPTH and minimal BMD changes. Can PK / binding improve target

PT gland capacity limits maximal PTH response



Limited maximal PTH response (confirmed in Ph 1) = would likely limit BMD response



K. Baron, M. Riggs, R. Sawamura, T. Shimizu, F. Okada, J. Zhou, T. Shibayama, and M. Jansen. An evaluation of calcilytic effects on parathyroid hormone and bone mineral density response using a physiologically-based, multiscale systems pharmacology model. *J. Bone Miner. Res.* 28, (Suppl 1, 2013).

Open iPSP GnRH modulation for endometriosis?

Question: Can we design a shorter trial? Which biomarker(s) to target response to optimize efficacy while staying below a threshold for BMD loss?



Updated
Existing
Systems Pcol



Logistic
PK-PD

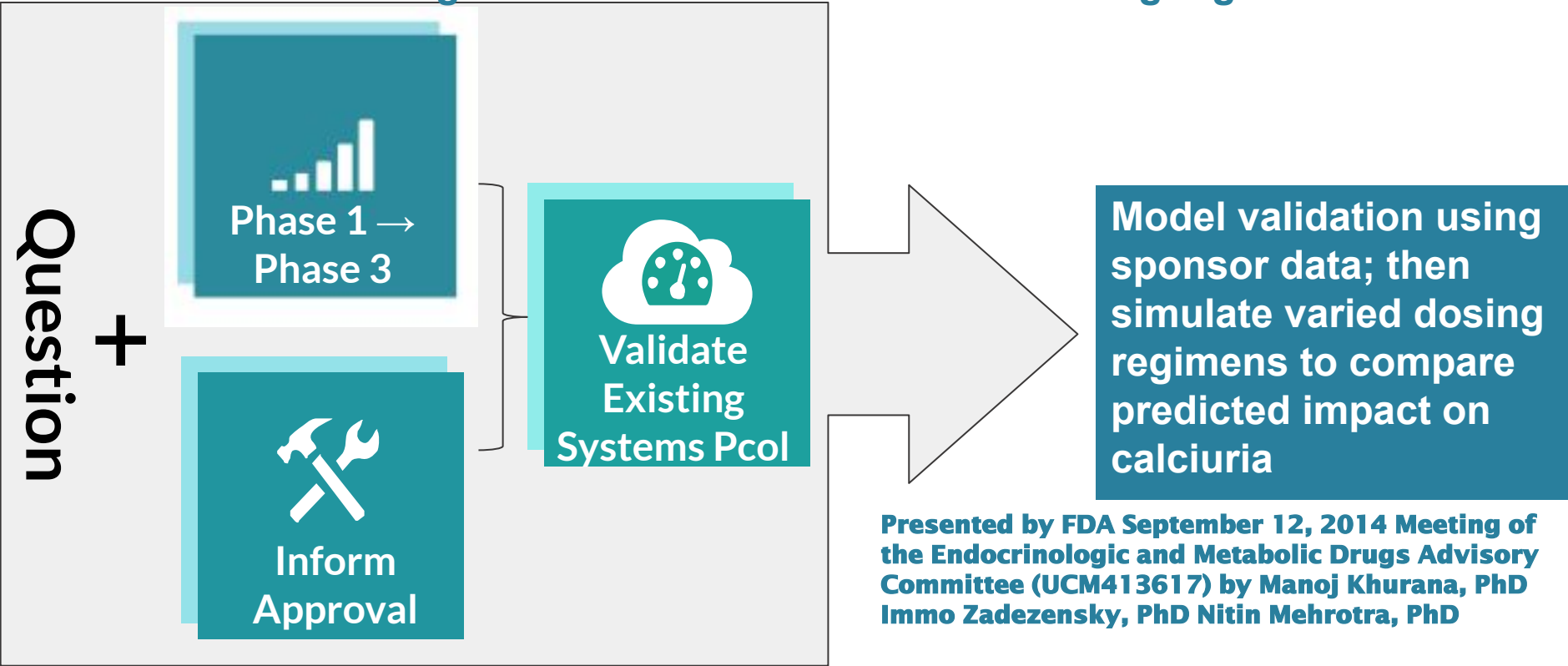
Evaluate time and magnitude of changes in candidate biomarkers. Define therapeutic index.

“ ... targeting the GnRH pathway to achieve the desired range of serum estrogen levels would be difficult to achieve; therefore, **the research program was halted before any compound entered the clinic.**”

P. A. Milligan, M. J. Brown, B. Marchant, S. W. Martin, P. H. van der Graaf, N. Benson, G. Nucci, D. J. Nichols, R. A. Boyd, J. W. Mandema, S. Krishnaswami, S. Zwillich, D. Gruben, R. J. Anziano, T. C. Stock, and R. L. Lalonde. Model-based drug development: a rational approach to efficiently accelerate drug development. Clin Pharmacol Ther, 93(6):502–14, Jun 2013.

Open iPSP PTH, replacement in hypoPTH

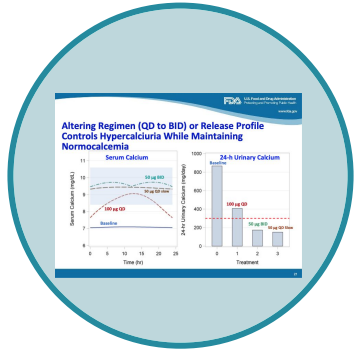
Question: Is QD dosing the safest and most effective dosing regimen?



Presented by FDA September 12, 2014 Meeting of the Endocrinologic and Metabolic Drugs Advisory Committee (UCM413617) by Manoj Khurana, PhD Immo Zadezensky, PhD Nitin Mehrotra, PhD

Open science opens doors

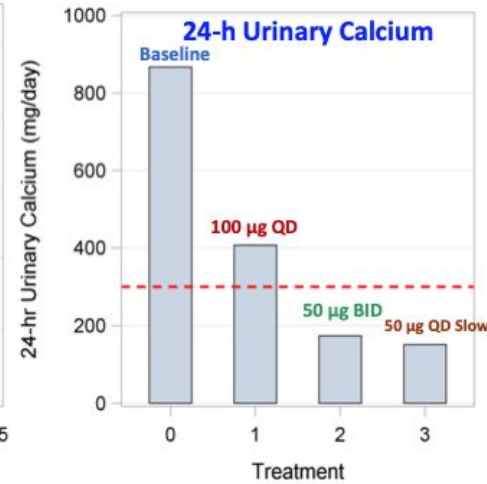
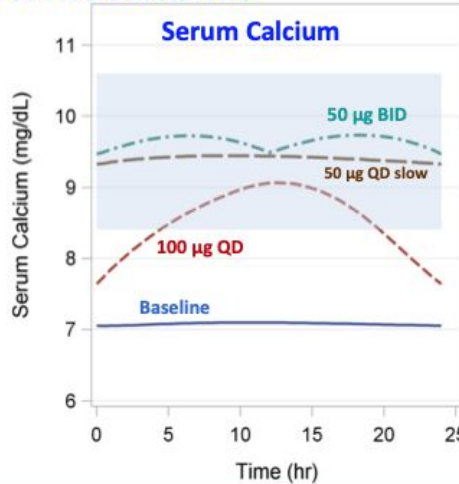
Altering Regimen (QD to BID) or Release Profile Controls Hypercalciuria While Maintaining Normocalcemia



PTH for Hypoparathyroidism

Clinical data

FDA suggested BID or sustained release likely to retain efficacy while minimizing risk of hypercalciuria



21

Presented at FDA September 12, 2014 Meeting of the Endocrinologic and Metabolic Drugs Advisory Committee (UCM413617) by Manoj Khurana, PhD Immo Zadezensky, PhD Nitin Mehrotra, PhD

Open iPSP PTH, replacement in hypoPTH

Question: Is QD dosing the safest and most effective dosing regimen?

11-75 Development of a Quantitative Systems Pharmacology Model to Support Dosing of rhPTH(1-84), a Recombinant Human Parathyroid Hormone, in Adult Patients with Hypoparathyroidism

Thomas Peyret,¹ Benjamin Rich,¹ JF Marier,¹ Nicole Sherry,² Richard Finkelman,² Ivy Song²

¹Certara Strategic Consulting, Princeton, NJ; ²Shire Human Genetic Therapies, Inc., Lexington, MA, USA, a member of the Takeda group of companies

Model Customization #2: Calcium Reabsorption

Various publications have reported a lower renal reabsorption of calcium in patients with hypoPT.[2] The effect is believed to be due to lower number of sites of hormone action in renal tubules, where it promotes calcium reabsorption. Based on this mechanism of action, the QSP model was customized as follows:

$$\text{CaReabsActive} = \frac{\text{CaReabs}_{\text{max}} \times \text{Ca}_{\text{serum}}}{\text{CaReabs}_{50} + \text{Ca}_{\text{serum}}} \times \text{PTHeffect}$$

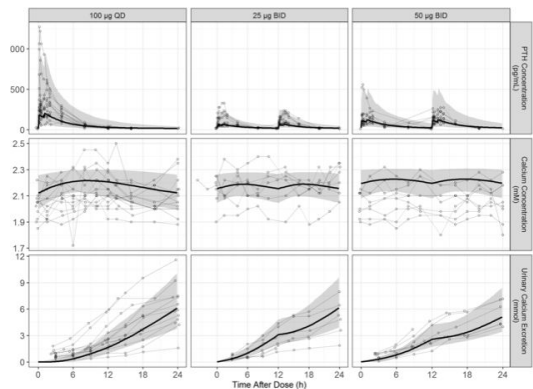
$$\text{PTHeffect} = \frac{\text{PTHeffect}_{\text{max}} \times \text{PTH}_{\text{plasma}}}{\text{PTHeffect}_{50} + \text{PTH}_{\text{plasma}}}$$

Where CaReabs₅₀ = C_{serum} needed to achieve 50% of CaReabs_{max}; CaReabsActive = rate of calcium active renal reabsorption (mmol/h); CaReabs_{max} = maximum CaReabsActive; C_{serum} = Calcium serum concentration (mM) (T16 in model code); PTH_{plasma} = PTH plasma concentration (pM); PTHeffect = effect of PTH on Calcium renal active reabsorption (unitless); PTHeffect₅₀ = PTH_{plasma} needed to achieve 50% of PTHeffect_{max} (T17 in model code); PTHeffect_{max} = maximum PTHeffect.

In the original QSP model, the maximum effect of PTH on calcium renal active reabsorption, PTHeffect_{max}, was fixed to 1.06147 (106% of reference reabsorption) and the PTH concentration needed to achieve 50% PTHeffect_{max} was scaled. In the current QSP model, the following values were used:

$$\text{PTHeffect}_{50} = 3.85 \times \text{T16} - 3.85 = 0.2366595$$

Figure 1. QSP Model Qualification - Study SHP634-101^[3]



PAGE 28 (2019) Abstr 9002

[www.page-meeting.org/?abstract=9002]

CONCLUSIONS

Both QD and BID dosing regimens of rhPTH(1-84) at daily doses from 25 µg to 100 µg markedly reduced urinary calcium excretion and the possibility of hypercalciuria while maintaining serum calcium level in target range as compared to the SOC.

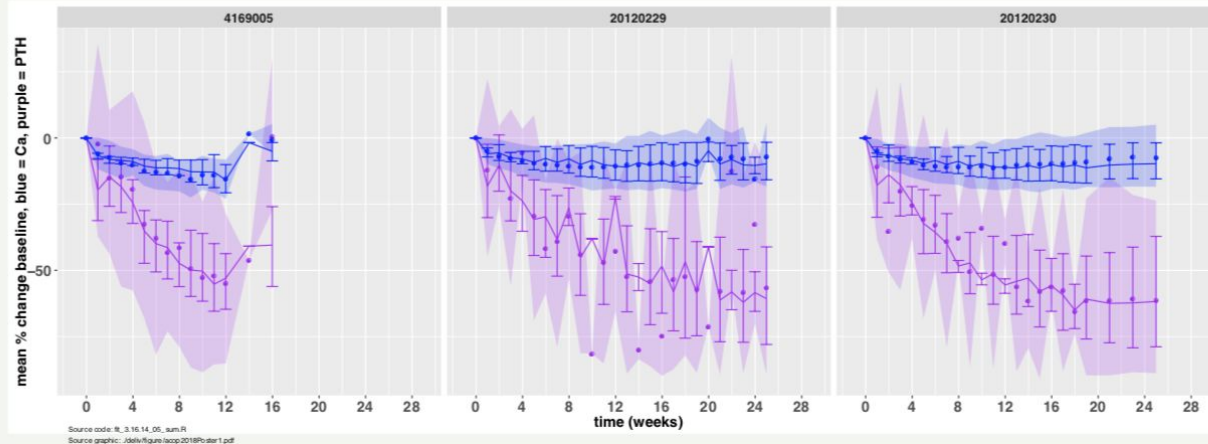
Although these modeling simulations appear to show that rhPTH(1-84) BID dosing regimens may predict a lower likelihood of hypercalciuria than the QD dosing regimens, clinical data are needed, and a clinical study is planned to confirm these findings.

Open iPSP: Integrate System, Disease, Drug

Model comes full circle, applied now to another Amgen-Sponsored Program

Chronic Kidney Disease-Mineral Bone Disorder

Long-Term Predictive Checks

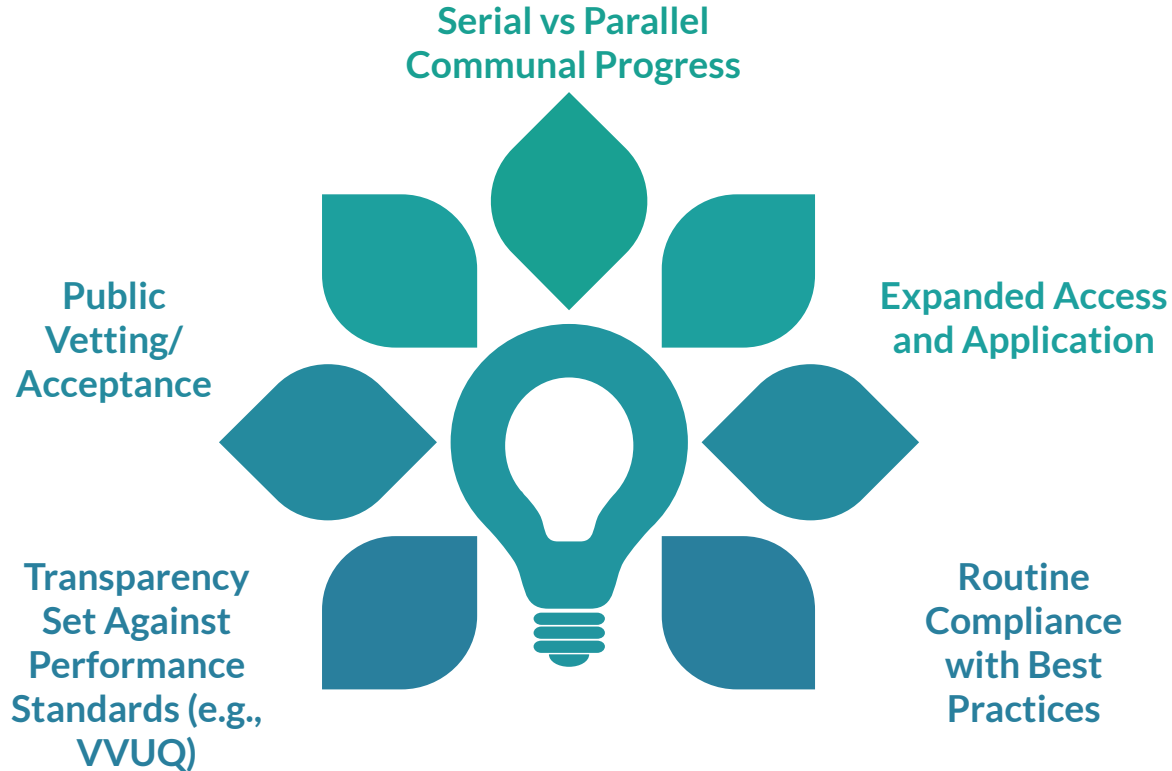


Despite continued decline in PTH (e.g., beyond weeks 4-6), feedback controls lead to leveling and partial rebound in Ca.

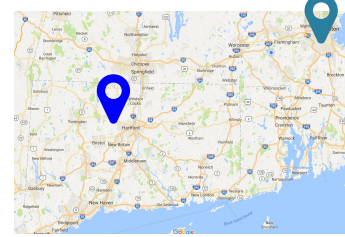
Figure 2: *Predictive check: change from baseline (percentage) for serum calcium (blue) and PTH (purple)*

Phase 3 Study 20120229 was included as external validation. Observed data: solid circle (mean) and 10th - 90th percentile range (shaded region); Simulated data: mean (solid line) and 10th - 90th percentile range (error bars).

Riggs MM, Baron KT, Melhelm M (2018) Multiscale physiology-based modeling of mineral bone disorder in patients with chronic kidney disease and secondary hyperparathyroidism on hemodialysis: application to etelcalcetide treatment effects on calcium homeostasis. ACoP9 Abstract #T-078.



Questions?



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