AAPS Forum to Connect Predictive Modelers MAY 6–7, 2019 • BOSTON, MA

Systems Pharmacology A day in the life

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METRUM

RESEARCH GROUP





Session Description and Objectives

 Description: Six pharmaceutical predictive modelers will share the day-to-day methods, challenges, teams, and applications specific to their respective disciplines.

- Introduce and foster an appreciation of what Predictive Modelers in other disciplines contribute to pharmaceutical R&D;
- Develop an understanding of common and unique approaches across PM disciplines; comparisons and contrasts will be carried into the following breakout sessions

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What is Systems Pharmacology?

Tagline

If I push this button (i.e. - block this pathway), why does THAT happen?

What it feels like sometimes Whack-a-mole



https://gph.is/2IVWWY8



What is Systems Pharmacology?



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Example of a Systems Model

"When he first presented his mathematical model of cardiovascular function ... in **1968**... responses ... (2)... reflected a tone of disbelief and even sarcasm. Dr. Guyton's systems analysis had **predicted a dominant role for the renal pressure natriuresis mechanism in long-term blood pressure regulation**, a concept that seemed <u>heretical</u> to most investigators at that time." http://www.the-aps.org/membership/obituaries/arthur_guyton.htm

2. Guyton AC, Coleman TG. Quantitative analysis of the pathophysiology of hypertension. Circ. Res. 1969, 24 (Suppl I): I1-I19.



Guyton AC, Coleman TG, Granger HJ 1972. Circulation: overall regulation. Annu Rev Physiol 34:13-46.



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Types of questions our models inform

Efficient Impact on Critical Decisions

Identification

Viable target/pathways involved in disease for drug or combination of therapies? Effects of perturbing one or more, simultaneously or timedfor purpose

Dosing support

Patient responder identification, biomarker identification and development, study design (what to measure when in whom)

Knowledge / Assumption Testing

What is the interplay Pharmacology - Biology -Pathophysiology -- where are the gaps, how much do those gaps affect predictability

Expand / Understand

How can one drug, disease, and/or mechanism inform another?

Teams/partners interface

How does it work... teamwork!



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You must remember this. An ODE is just an ode, a sine is just a sign. The fundamental things apply. Louis Armstrong, paraphrase



to describe pathways, cytokines, tissues, organ functions (and structure)

From integrated proteomic subtypes to drug sensitivity prediction

Martin Frejno et al. Mol Syst Biol 2017;13:951 © as stated in the article, figure or figure legend

Proteomics

molecular

systems biology

Wang, Z., Birch, C. M., & Deisboeck, T. S. (2008). Cross-scale sensitivity analysis of a non-small cell lung cancer model: linking molecular signaling properties to cellular behavior. *Bio Systems*, 92(3), 249–258.

https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2430419/figure/F1/?report=objectonly

Figure 1

Phenotypic decision process for a cancer cell between two time steps.

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Cosgrove, J., Butler, J., Alden, K., Read, M., Kumar, V., Cucurull-Sanchez, L., ... Coles, M. (2015). Agent-Based Modeling in Systems Pharmacology. *CPT: Pharmacometrics & Systems Pharmacology*, *4*(11), 615–629.

Agent-Based Modeling in Systems Pharmacology

CPT: Pharmacometrics & Systems Pharmacology, Volume: 4, Issue: 11, Pages: 615-629, First published: 12 August 2015, DOI: (10.1002/psp4.12018)

You must remember this. An ODE is just an ode, a sine is just a sign. The fundamental things apply. Louis Armstrong, paraphrase

Cross-Functionality is Captured Mathematically

$$\begin{split} d\,/\,dt\; A(21) &= ((k_{21D}\text{*}A(21)_0 + k_{21-24}\text{*}A(21)_0\text{*}A(22)_0 - k_{24-21}\text{*}\\ A(24)_0)\,/\,A(20)_0^{120,21} \text{*}A(20)^{120,21} - k_{21D}\text{*}A(21) \end{split}$$

 $d/dt A(22) = k_{225} - k_{220} * A(22) - k_{21-24} * (A(23) * A(22))$

 $-\mathbf{k}_{21-24} \cdot \mathbf{A}(21) \cdot \mathbf{A}(22) + \mathbf{k}_{24-21} \cdot \mathbf{A}(24)$

$$\begin{split} &+ A(21)^*A(22)) + k_{24-21}*(A(24) + A(25)) \\ &k_{224} = k_{220}*A(22)_0*(A(17)/A(17)_0)^{Y(7,22)}*\alpha_{7,22}*(A(7)/V_{value}) \end{split}$$

 $/(\delta_{7,22}^{*}(A(17)/A(17)_{0})^{\gamma_{17,22}} + (A(7)/V_{max}))$

 $d/dt A(7) = H_{4,10-7}^{-} *(A(10)/0.5)*A(11) - k_{70}*A(7)$

Intracellular phosphate

Kidney

 $d/dtA(8) = v_{3-4} - v_{8-5}$

 $d / dt A(9) = k_{05} * H_{2,9} * H_{3,9} - k_{00} * A(9)$

H_{5.9} = 1 for A(5)<A(5)₀

You must remember this. An ODE is just an ode, a sine is just a sign. The fundamental things apply. Louis Armstrong, paraphrase

Trame, M. N., Riggs, M., Biliouris, K., Marathe, D., Mettetal, J., Post, T. M., ... Musante, C. J. (2018). A Perspective on the State of Pharmacometrics and Systems Pharmacology Integration. *CPT: Pharmacometrics & Systems Pharmacology*. https://doi.org/10.1002/psp4.12313

Methods behind our models

What makes them tick?

Persistence over Parsimony

Models to describe the "system" often contain many, many, many parameters: fix, tune or estimate

Parameter Values

Sensitivity analysis, likelihood profiling, optimization algorithms, verification and validation

"Virtual" Patients

With this method, in reality, we don't differentiate lack of identifiability from real variability in these "patients"

One Lump or Two?

Scale reduction techniques: linearization

Bayesian Estimation

Balance known/unknown, existing info with incoming information, understand variability accordingly

Duffull, S. B. (2016). A Philosophical Framework for Integrating Systems Pharmacology Models Into Pharmacometrics. *CPT: PSP*, *5*(12), 649–655. Rieger et al. (2018). Improving the generation and selection of virtual populations in quantitative systems pharmacology models. *Progress in Biophysics and Molecular Biology*, 139, 15–22.

Ribba B, Grimm HP, Agoram B, et al. Methodologies for Quantitative Systems Pharmacology (QSP) Models: Design and Estimation. *CPT PSP*. 2017;6(8):496–498

A. Dokoumetzidis and L. Aarons. Proper lumping in systems biology models. IET Syst Biol, 3(1):40, 2009 Jan. Plan EL, Baron KT, Gastonguay MR, French JL, Gillespie WR, Riggs MM. Bayesian Joint Modeling of Bone Mineral Density And Repeated Time-To-Fracture Event For Multiscale Bone Systems Model Extension. PAGE Meeting. 2012.

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Software Programs

Pro: many options / Con: Lack of uniformity / tranferability in our tools

Pro: Many options

Con: Lack of uniformity / interoperability

Trade-offs: Speed, versatility (complexity, code) vs ease of use (GUI, visual)

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Literature vs Internal vs 3rd Party

Data Sources

With so many sources, how / when is it appropriate to integrate.

Gulati A and Tannenbaum S. Using Visualization to Address the Reliability of Sources of Initial Parameter Values in a Quantitative Systems Pharmacology (QSP) Model. AAPS PharmSci 360 2018.

Model Verification and Validation

Reproducible research

Model Verification and Validation

Reproducible research

Cucurull-Sanchez et al. (2019). Best Practices to Maximize the Use and Reuse of Quantitative and Systems Pharmacology Models: Recommendations From the UK Q&SP Network. *CPT: PSP.*

Kirouac, D. C. (2018). How Do We "Validate" a QSP Model? CPT: PSP, 7(9), 547–548.

Friedrich, C. M. (2016). A model qualification method for mechanistic physiological QSP models to support model-informed drug development. CPT: PSP, 5(2), 43–53.

Agoram, B. (2014). Evaluating systems pharmacology models is different from evaluating standard pharmacokinetic-pharmacodynamic models. CPT: PSP, 3, e101.

Kirouac, D. C., Cicali, B., & Schmidt, S. (2019). Reproducibility of Quantitative Systems Pharmacology Models: Current Challenges and Future Opportunities. CPT: PSP, 8(4), 205–210.

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Challenges Faced

Don't cry to give up, cry to keep going. ... Get a reward from it E. Thomas

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Example of a Systems Model

"In 2004, denosumab was under development for treating osteoporosis, ... Questions existed that could not be practically addressed with clinical studies due to the protracted dosing interval (q6M) and required trial duration, nor by traditional PKPD models. ... " Peterson and Riggs. CPT:PSP, 4(3), 2015.

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If I push this button (i.e. - block this pathway), why does THAT happen?

As reported in: M.R. McClung, E.M. Levencki, S. B. Cohen, M.A. Biolognese, G. C. Wootson, A. H. Moffet, M. Pascosk, P. D. Millor, S. N. Lederman, C. H. Chesnul, D. Lain, A. J. Kivlar, B. L. Holoway, C. Zh, M. C. Peterson, P. J. Bester, and AMD 142 Bone Loss Study Group, Denounnal in postmenopausa women-with tow tone ministra density. N Engl J Mod, 35(4):121-31, Feb 2006.

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Denosumab: RANKL inhibition \rightarrow Bone Markers \rightarrow BMD Change

Matthew M. Riggs, Kyle T. Baron, Elodie L. Plan, Marc R. Gastonguay. Qualification of a Physiologically-Based Model for Predicted Bone Marker and Bone Mineral Density Changes Associated with Denosumab Treatment. Presented at American Society of Bone Mineral Research (ASBMR) Annual Meeting, Minneapolis, MN; October 14, 2012 (Abstract# SU0363). Available at: <u>http://metrumrg.com/index.php/publications</u>

Model Development Integrate System, Disease, Drug

Chronic Kidney Disease-Mineral Bone Disorder

Fig. 1, 2; M. M. Riggs, M. C. Peterson, and M. R. Gastonguay. Multiscale physiology-based modeling of mineral bone disorder in patients with impaired kidney function. J Clin Pharmacol, 52(1 Suppl):455–53S, Jan 2012.

- Can these effects be used to describe PTH and Ca response following long-term etalcalcetide treatment?

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Multiscale Model

Riggs MM, Baron KT, Melhem 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.

Model Development Integrate System, Disease, Drug **Chronic Kidney Disease-Mineral Bone Disorder**

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.

-+ Pharmacology **Direct and indirect** effects in PT gland

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Model Development Integrate System, Disease, Drug

Chronic Kidnev 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.

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Acknowledgements

Eudy-Byrne Baron Plan Shibayama French Shimizu Okada Jansen Ocampo-Pelland Melhem vanderGraaf Martin Zhou Sawamura Peterson Gillespie

(Word)*It*Out

