

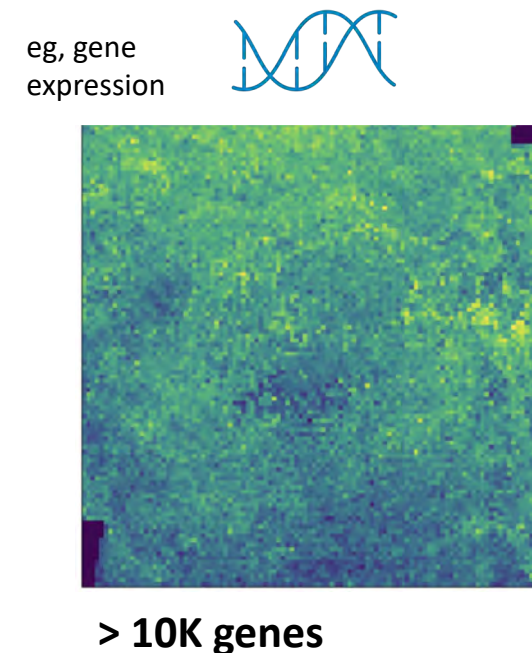
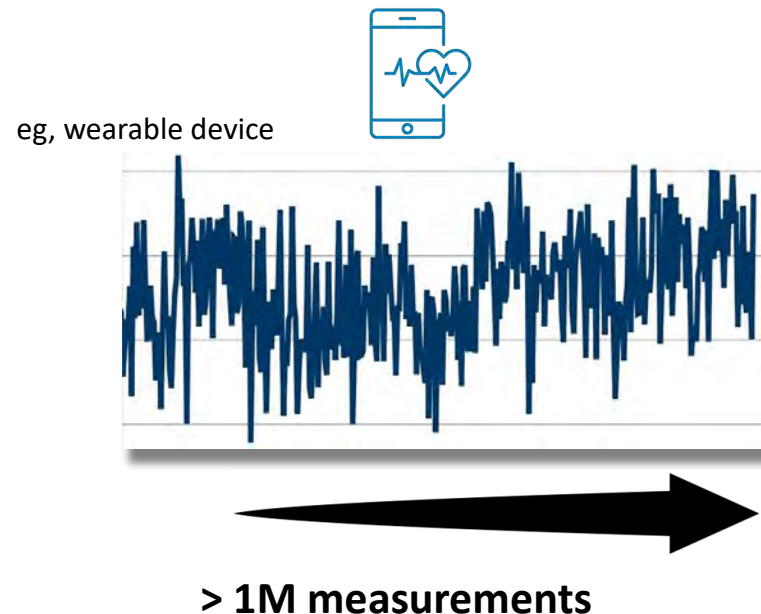
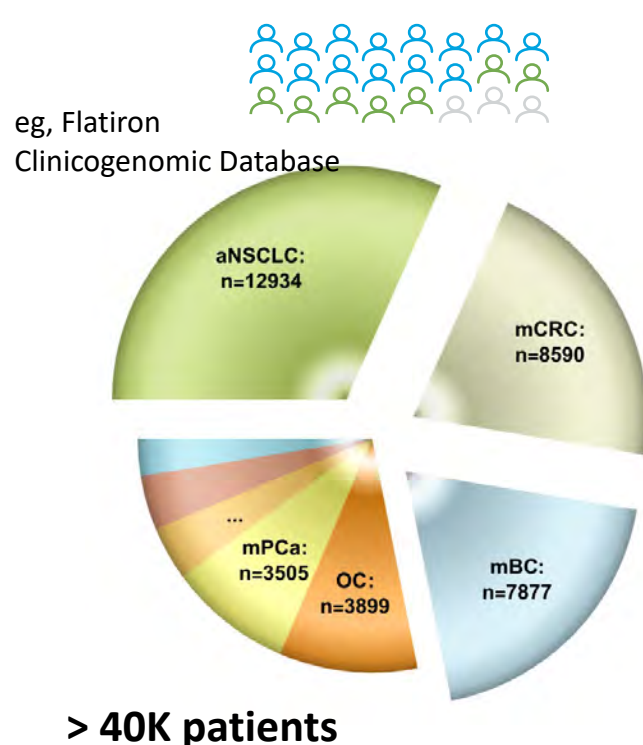
Neural-PK/PD as Pharmacology-Informed Deep Learning Architecture

James Lu, Genentech

IQ Workshop on Machine Intelligence for Quantitative Modeling in Drug Discovery & Development Applications
15-16 September 2022

Motivation: the Data Challenge in the Digital Age

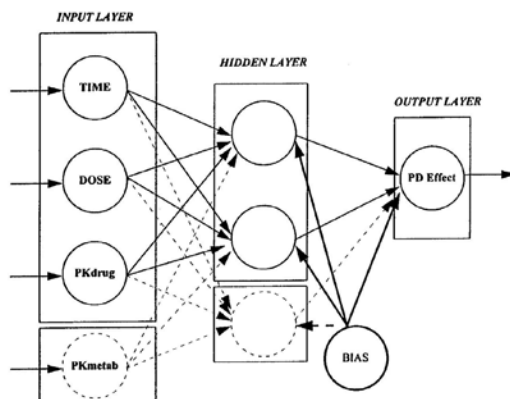
- Current trends in technology & digitization generate growth across:
 - the number of patients
 - the number of longitudinal measurements
 - the dimensionality of data



Cancer Genome Atlas Network, Nature (2012)

A Brief History of Neural Networks for PK & PK/PD Modeling

Feed-Forward Network

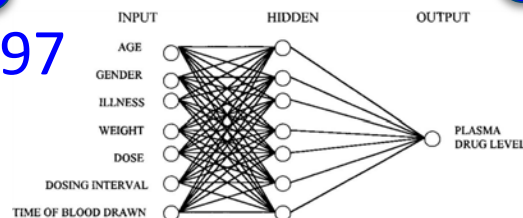


Artificial Neural Networks As a Novel Approach to Integrated Pharmacokinetic-Pharmacodynamic Analysis

JOGARAO V. S. GOBBURU¹* AND EMILE P. CHEN²

Received October 11, 1995, from the ¹Department of Pharmaceutical Sciences, North Dakota State University, Fargo, ND 58105, and ²Department of Drug Metabolism and Pharmacokinetics, Hoffmann-La Roche, Nutley, NJ 07110. Accepted for publication February 19, 1996[®]

1997

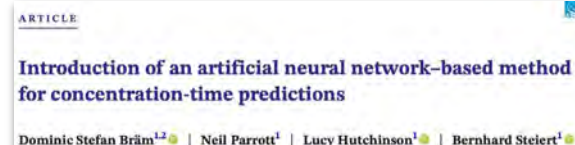


Application of Neural Networks to Population Pharmacokinetic Data Analysis

HSIAO-HUI CHOW¹*, KRISTIN M. TOLLE², DENISE J. ROE³, VICTOR ELSBERRY⁴, AND HSINCHUN CHEN²

Received September 24, 1996, from the ¹Department of Pharmacy Practice and Science, ²Department of Management Information Systems, and ³Arizona Cancer Center, University of Arizona, Tucson, AZ 85721. Accepted for publication March 24, 1997[®]

Recurrent Neural Network



Deep Learning
revolution

'05

Mapping the Dose-Effect Relationship of Orbofiban from Sparse Data with an Artificial Neural Network

DONALD E. MAGER,¹ JASON D. SHIREY,¹ DERMOT COX,² DESMOND J. FITZGERALD,³ DARRELL R. ABERNETHY¹

¹Laboratory of Clinical Investigation, National Institute on Aging, Gerontology Research Center, Baltimore, Maryland

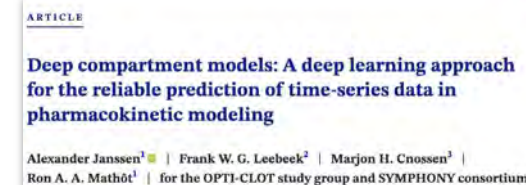
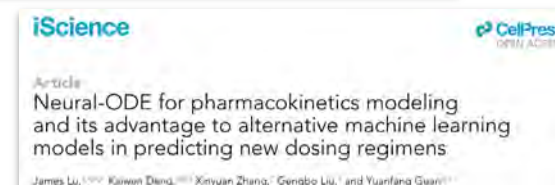
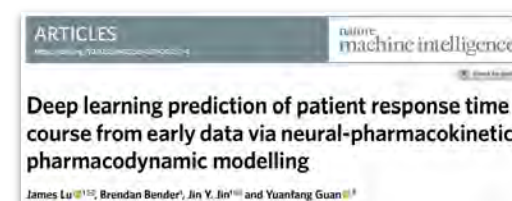
²Department of Clinical Pharmacology, Royal College of Surgeons in Ireland, Dublin, Ireland

³University College Dublin, Dublin, Ireland

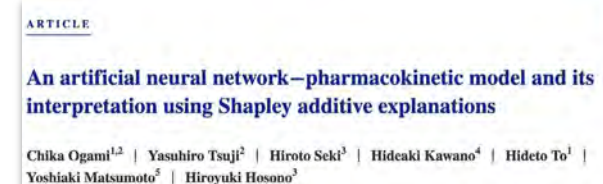
Population pharmacokinetic and pharmacodynamic models of remifentanyl in healthy volunteers using artificial neural network analysis

S. H. Kang, M. R. Poynton, J. M. Kim, H. Lee, D. H. Kim, S. H. Lee, K. S. Bae, O. Linarez, S. E. Kern, & G. J. Noh¹

Neural-ODE Chen et al, NeurIPS (2018)



'21



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PK and PK/PD Modeling from the Dynamical Systems Perspective

- Two distinct types of Dynamical Systems

- Autonomous:**

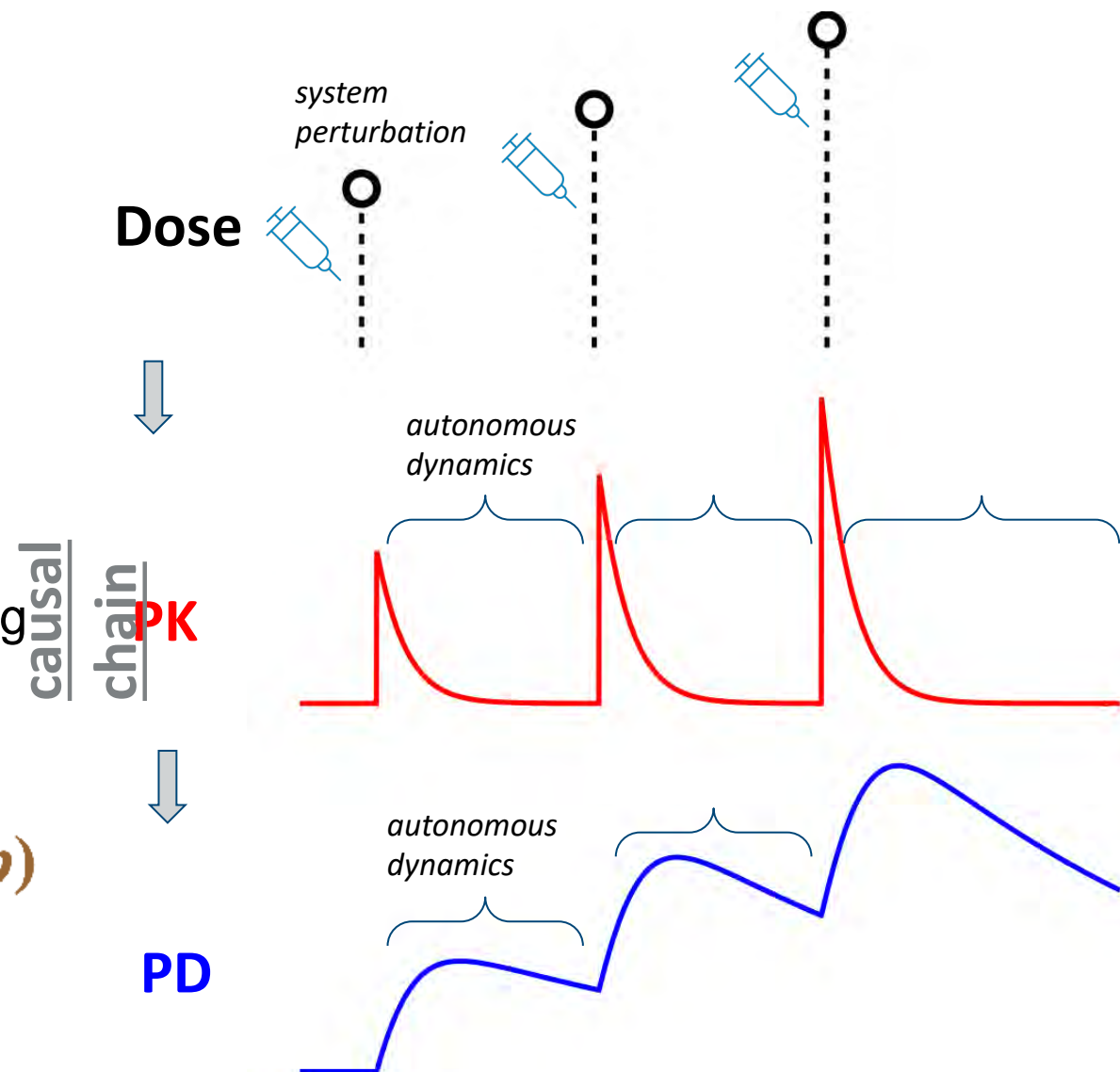
$$y'(t) = f(y(t), p)$$

- Non-autonomous:**

$$y'(t) = f(y(t), t, p)$$

- PK and PK/PD models are piecewise autonomous systems with dosing introducing time-dependent perturbations:

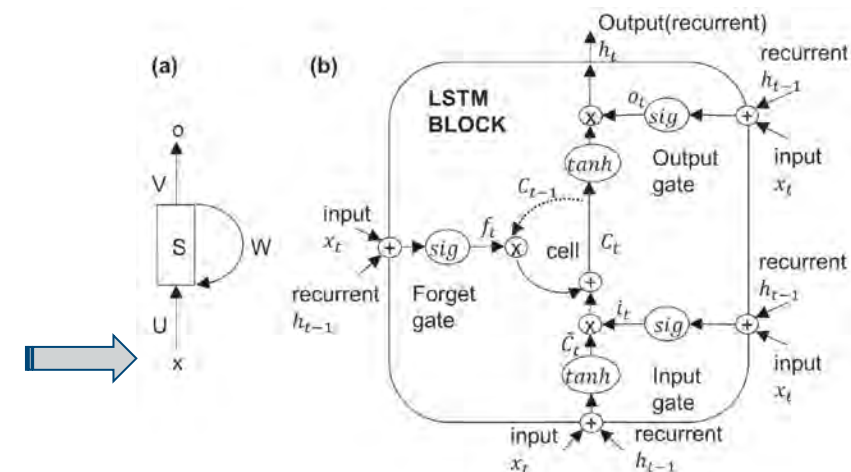
$$y'(t) = \sum_{i=1}^n \text{dose}(i) \delta(t - T_i) + f(y(t), p)$$



Distinctions with alternative Deep Learning Models for Time Series Data

Standard DL architectures for Time Series data (such as LSTM) do not explicitly encode causality relationships between **Dose**, **PK** and **PD**

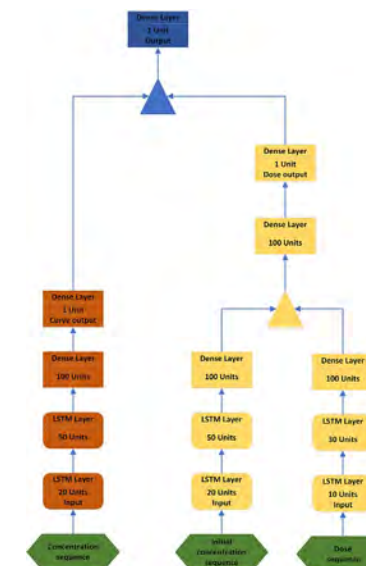
- All input data (x) enters the neural net at an equal footing
- Potentially challenging to perform counterfactual (“what-if”) simulations



Liu et al, IJCPT (2021)

Necessitated the development of architectures that separates out:

- Concentration sequence data
- Dose sequence data



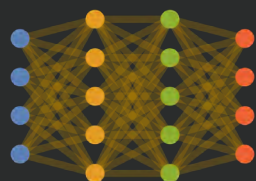
Bram et al, CPT:PSP (2022)

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The Merging of Deep Learning with Dynamical Systems

Deep Learning

- Excels in approximating high dimensional/nonlinear functions
- Learn to improve model as data increases



$$\frac{dy(t)}{dt} = \sum_{i=1}^n \text{dose}(i) \delta(t - T_i) + \left(\text{Neural Network} \right) (y(t), p)$$

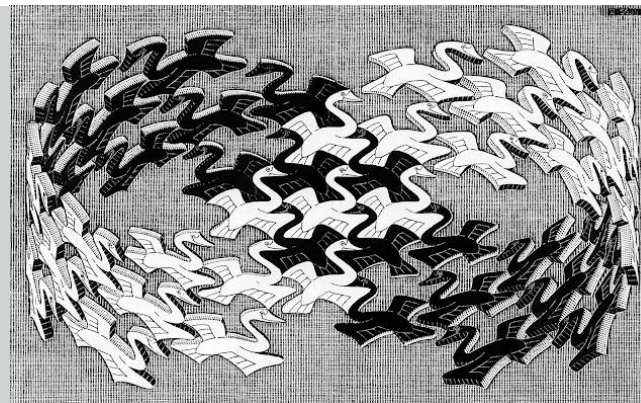
Dynamical Systems

- Enables assumption of causal dynamical relationships
- Enables dosing implementation as perturbation in the system state:

$$\frac{dy(t)}{dt} = \sum_{i=1}^n \text{dose}(i) \delta(t - T_i) + f(y(t), p)$$

Neur

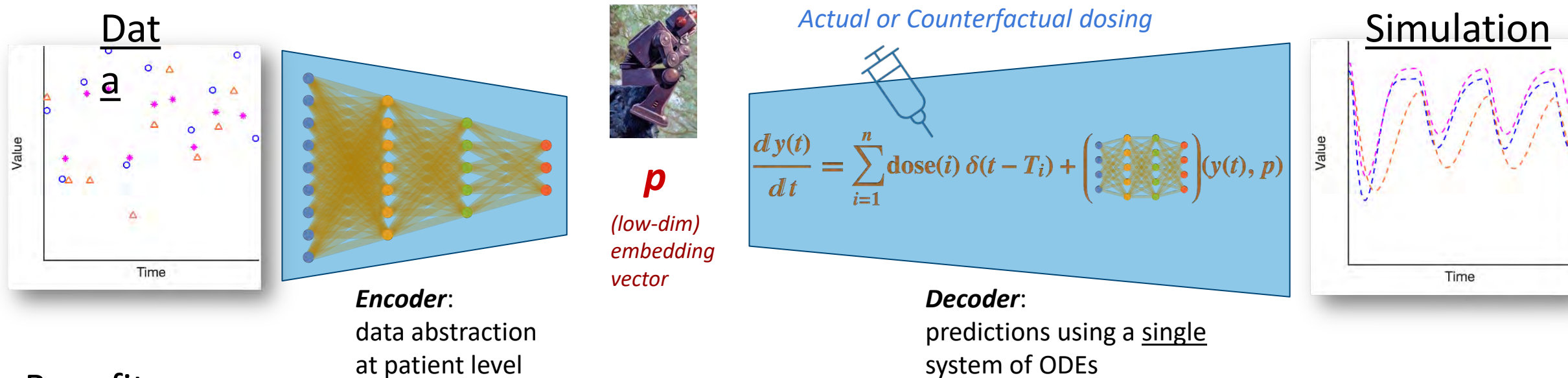
PK/PD



A pharmacology-informed dynamical system that learns from data and enables counterfactual dosing simulations

The Architecture of Neural-PK/PD enhances Interpretability

Pharmacology-Informed Encoder-Decoder Architecture



Benefits:

- Explicit control on the parameter dimension for characterizing *inter-individual variability*
- Explicit control on the state dimension for reproducing observed dynamics
- Explicit dosing port enables counterfactual simulations under different dosing regimens

Formulations of Population-PK/PD vs Neural-PK/PD

Human
Generated
Model

Equations based on human experience & data fit

$$DADT(1)=K_{21} \cdot A(2)-K_{12} \cdot A(1)-K_{10} \cdot A(1)$$

$$DADT(2)=K_{12} \cdot A(1)-K_{21} \cdot A(2)$$

$$DADT(3)=K_{EO} \cdot C_P - K_{EO} \cdot A(3)$$

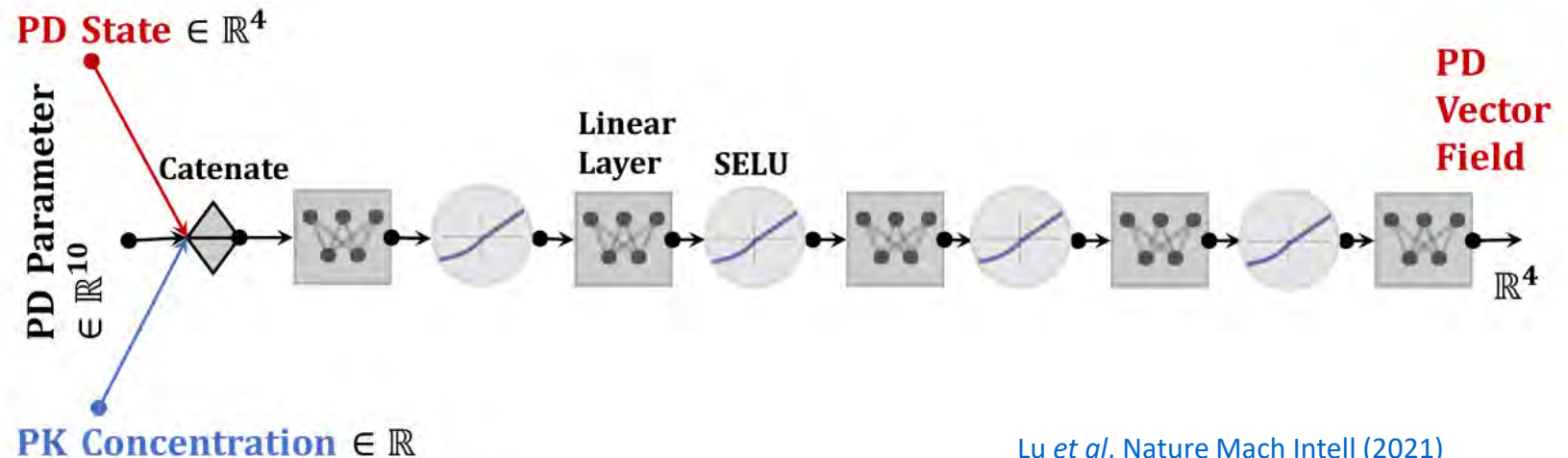
$$DADT(4)=-K_{TR} \cdot A(4)+K_{TR} \cdot A(4) \cdot (1-EFF_{PLT}) \cdot (BT/A(8))^{**}GAM$$

$$DADT(5)=-K_{TR} \cdot A(5)+K_{TR} \cdot A(4)$$

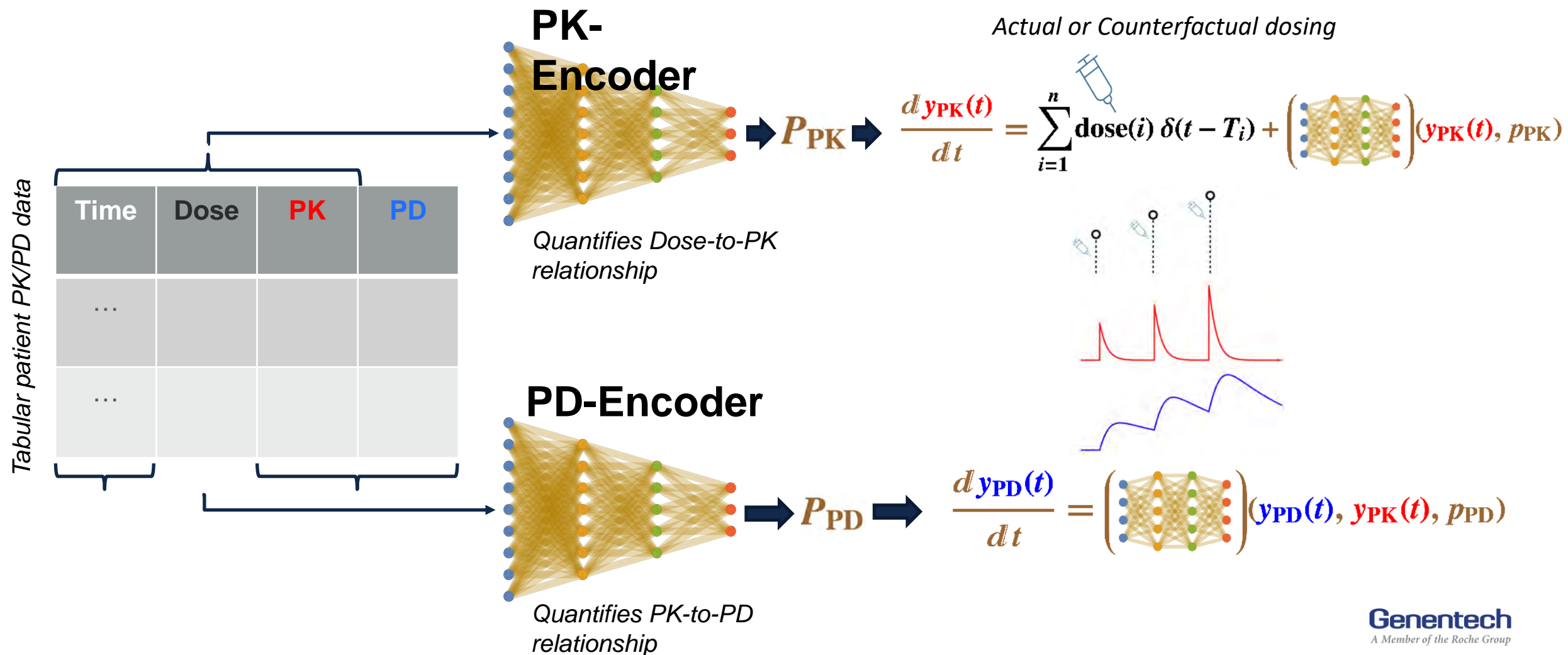
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Neural
Network
Generated
Model

Neural network learned from PK & PD data

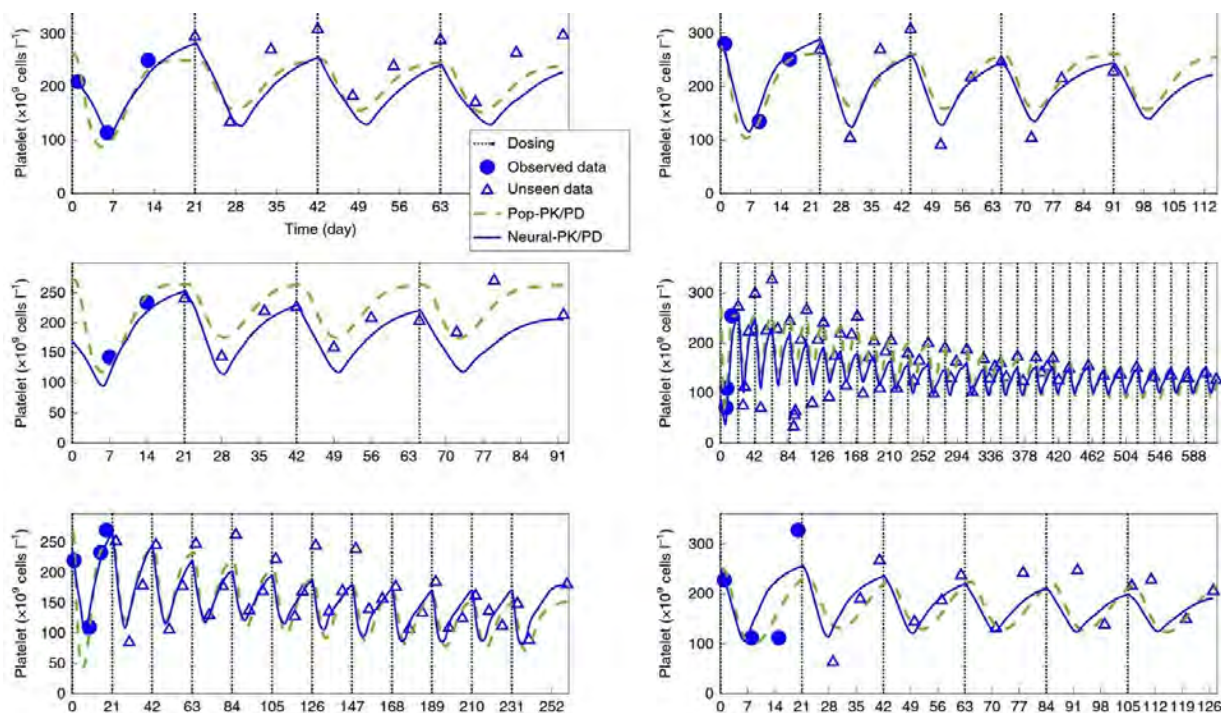


Encoding Causality Assumption into Neural-PK/PD Architecture



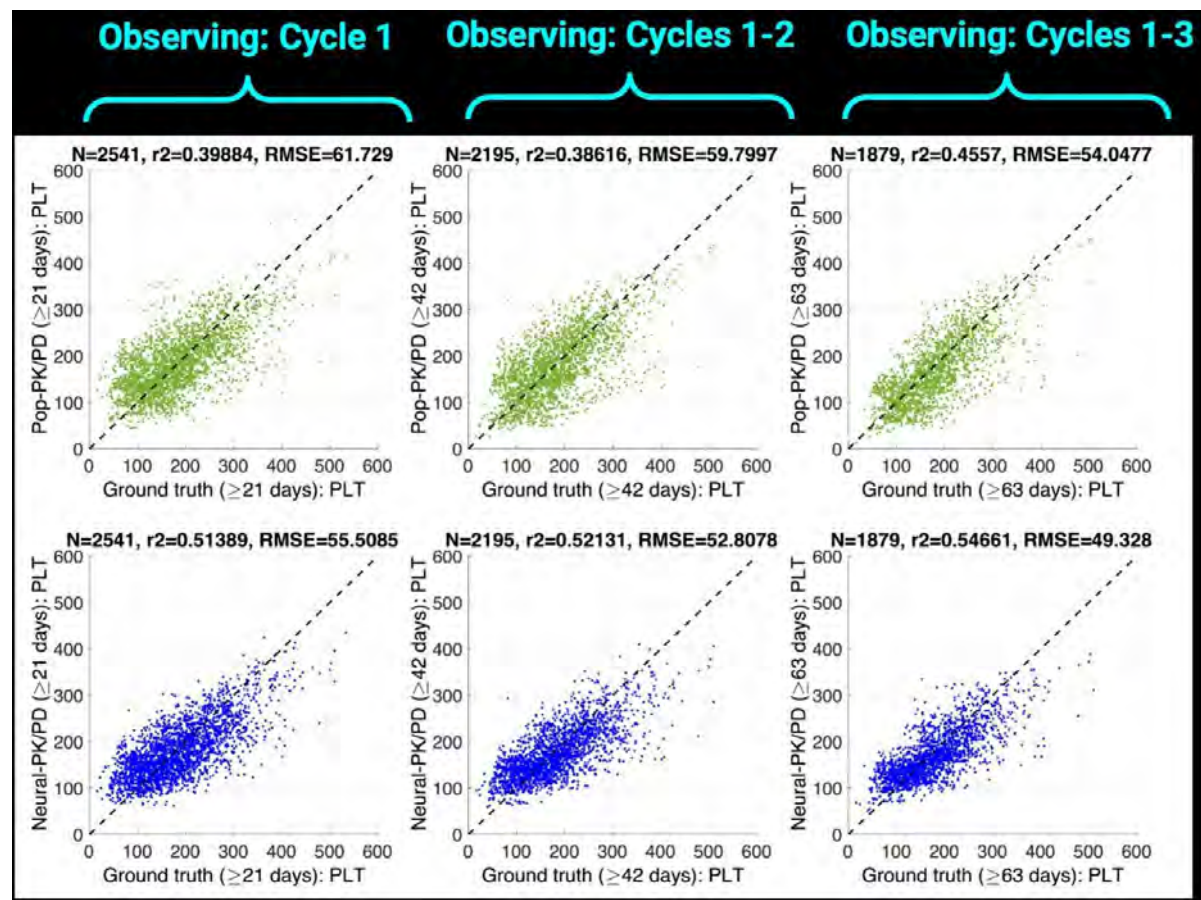
Neural-PK/PD Enables Accurate Predictions in Time

Learned model demonstrates qualitatively similar dynamics but more precise predictions as compared to the state-of-the-art pop-PK/PD model for platelet dynamics



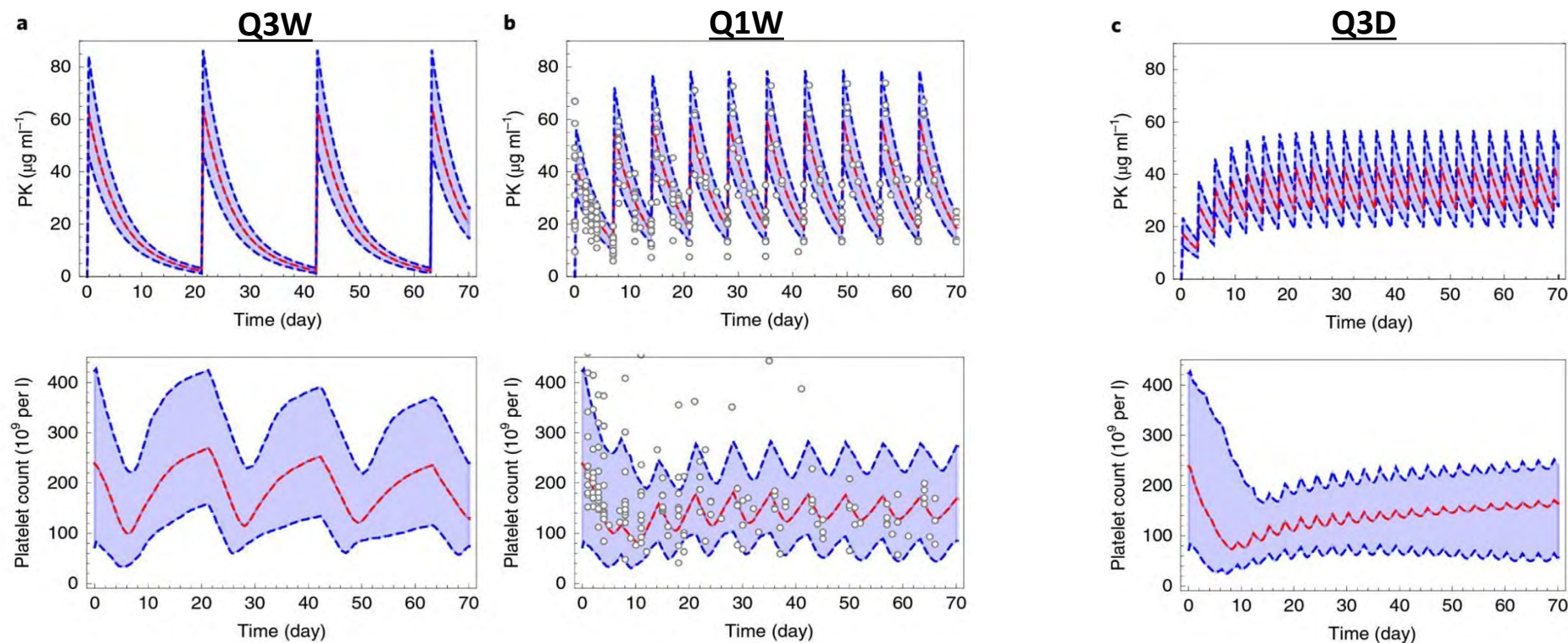
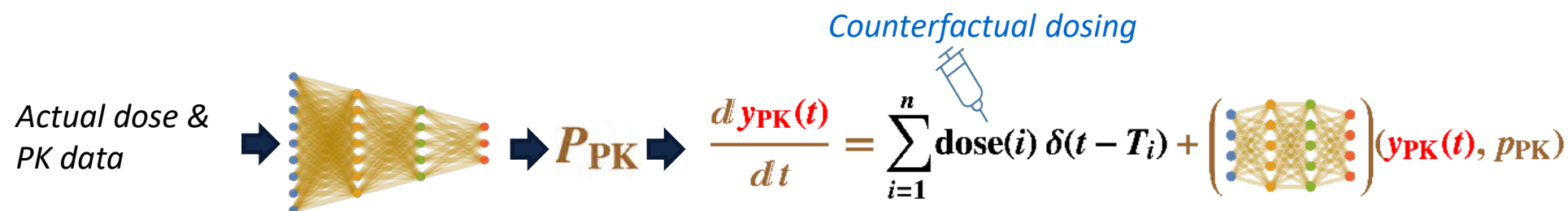
Lu et al, Nature Mach Intell (2021)

Model demonstrates superior predictive performance



Neural-PK/PD Enables Predictions for Alternate Dosing Regimens

Simulations of counterfactual dosing regimens shown to be consistent with data



Lu et al, Nature Mach
Intell (2021)

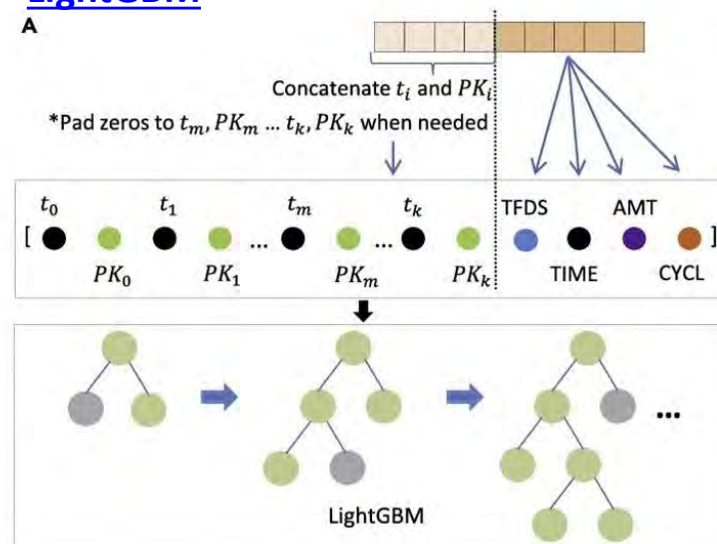
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The Importance of the Architecture Choice in Generalizability

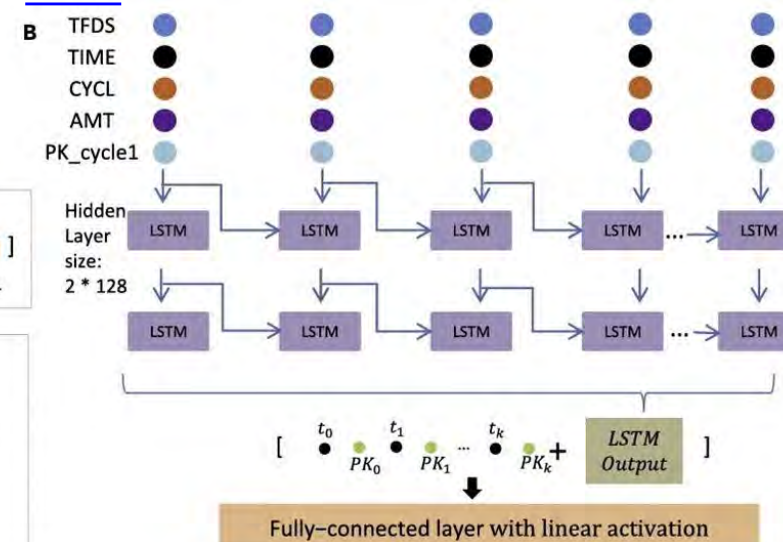
- Comparison of Neural-ODE based architecture vs alternate models:



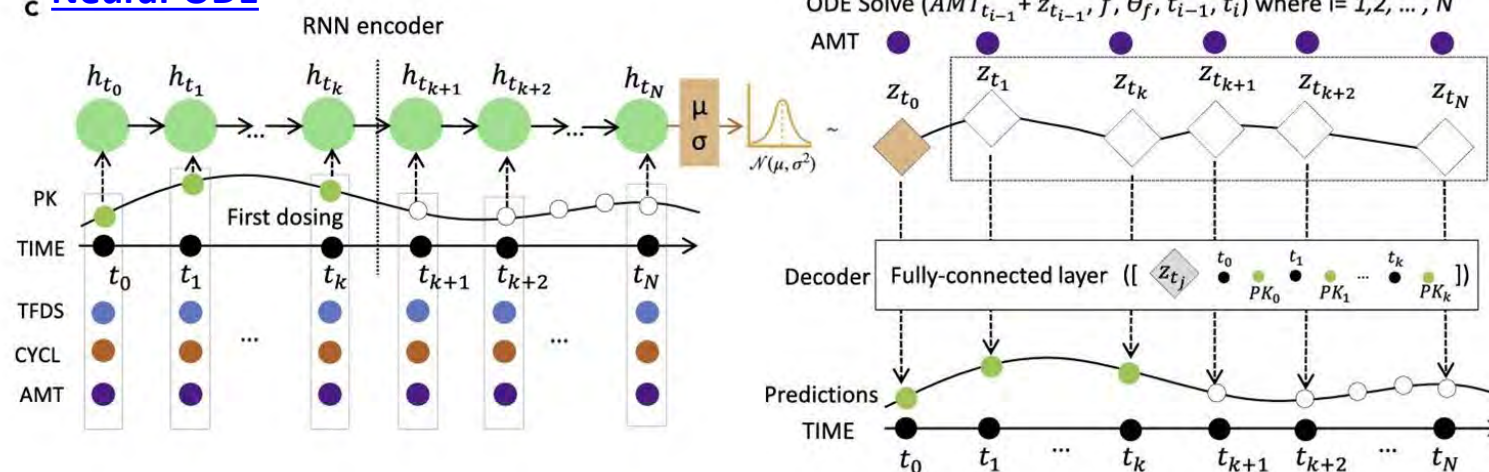
LightGBM



LSTM

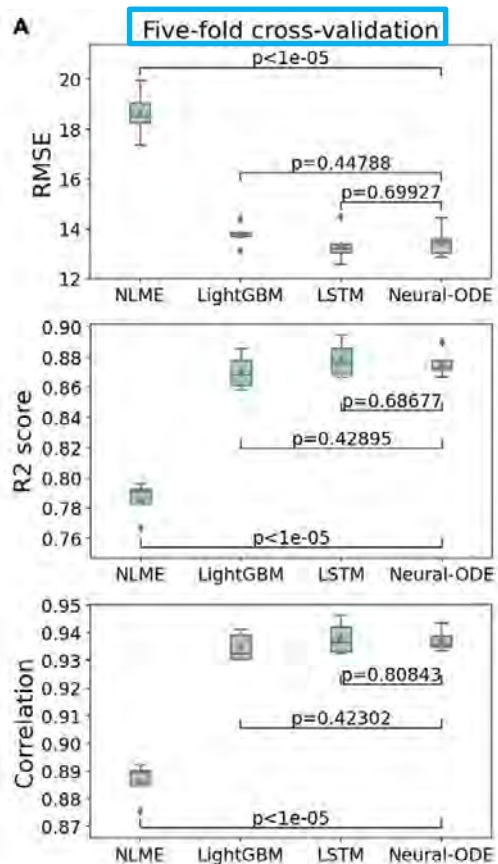


Neural-ODE

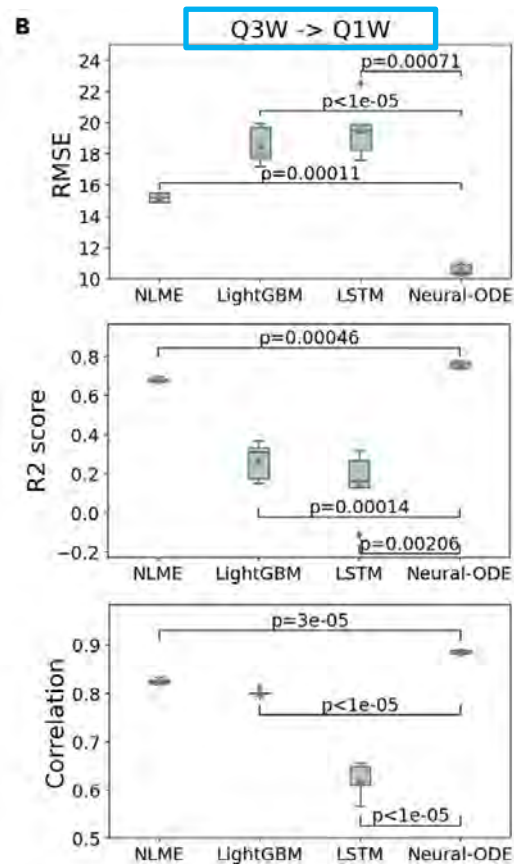


The Importance of the Architecture Choice in Generalizability

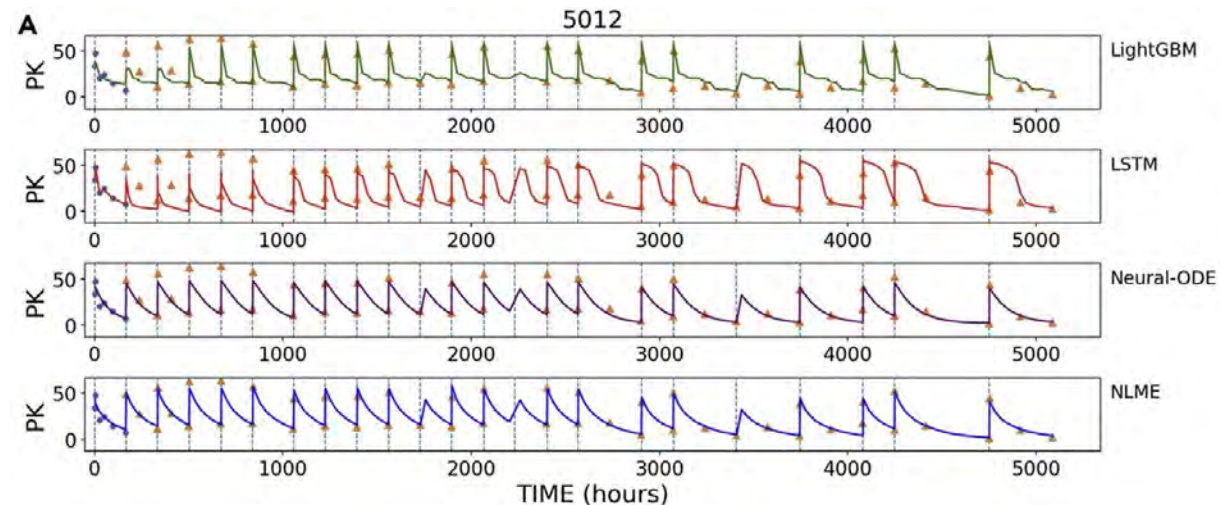
ML/DL models exhibit similar performance when train and test sets are from the same distribution



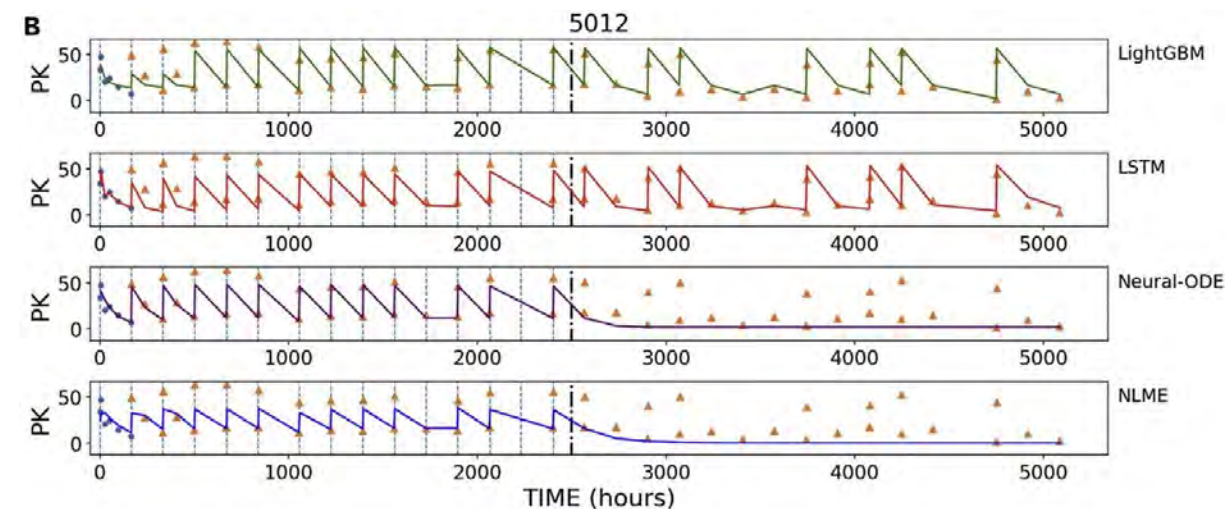
Models such as LightGBM and LSTM do not generalize well across dosing regimens



Comparison of model predictions for a patient



Comparison of model predictions with dosing stopped @ t=2500 hour



Conclusion

- The rise of complex, high volume PK/PD data in the Digital Age necessitates the development of AI methodologies
- Neural-PK/PD is a pharmacology-informed DL model that encapsulates *dose-concentration-effect* relationship within an encoder-decoder architecture:
 - leverages dynamical systems concepts
 - encoder bottleneck may improve better generalization from limited data
 - decoder enables counterfactual simulations

Acknowledgement

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University of Michigan/Ann Arbor Algorithms

- Kaiwen Deng, Xinyuan Zhang, Yuanfang Guan

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