

A Physiologically Based Pharmacokinetic Framework for Predicting Drug Exposure in the Female Reproductive Tract

Sophie Fischer-Holzhausen (1) , Nina Nauwelaerts (1) , Leonie Lautz (1) , Stephan Schaller (1) , Marco Siccardi (1)

(1) esqLABS GmbH, Saterland, Germany



Presenter:
Sophie Fischer

sophie.fischer-holzhausen@esqlabs.com

Intro

Healthcare is advancing, and a deeper understanding of complex physiological systems is crucial for improving treatments. The female reproductive tract (FRT) - affected by conditions such as infections, endometriosis, and cancer - plays a key role in women's health. Nevertheless, the FRT is typically neglected in physiologically-based pharmacokinetic modeling (PBPK). To enhance exploration of drug exposure in the FRT following both local (e.g., vaginal or uterine) and systemic administration routes, we developed a PBPK model extension for the FRT.

Methods

A PBPK extension module (Figure 1) was developed in MoBi® V12. It integrates physiological data for the FRT organs [1] and allows for the simulation of different local drug administration routes. The extension module was validated for **levonorgestrel** and **metronidazole** by comparing predicted plasma, tissue, and fluid concentrations to clinical data, using goodness-of-fit plots and pharmacokinetic parameters. In total, we included 4 local (intrauterine device, vaginal gel/tablet/pessary) and two systemic (IV and oral) routes of administration. Release descriptions for the different formulations were parametrized using clinical data.

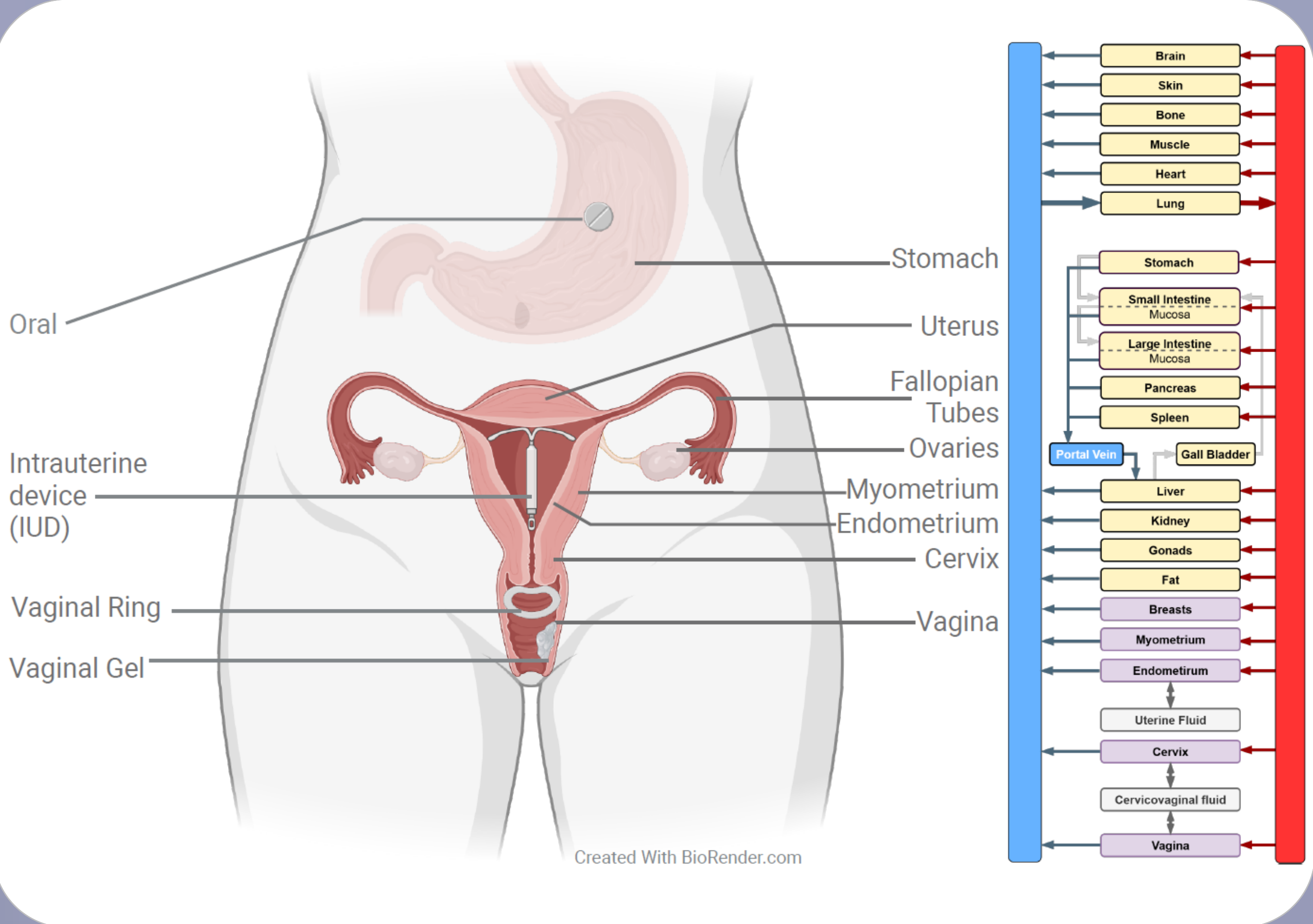


Figure 1: Female reproductive tract structure and representation of drug administrations. (A) Presentation of the anatomical structure of the female reproductive tract organs and the considered local routes for drug administration. (B) Schematic representation of the PBPK human base model structure in PKSim extended by the organs (pink) and fluids (white) of the female reproductive tract.

Results

For both compounds, the PBPK models capture the plasma, tissue, and fluid exposure following different routes of drug administration well, as indicated in Figure 2. For metronidazole, time series data were available, allowing the calculation of AUC and C_{max} values. 81.25% of the derived ratios were in the range of 0.85 – 1.15. For levonorgestrel, mainly single-time point measurements were available; therefore, the model predictions were evaluated using the prediction error. 62.5% of prediction errors were in the range of $\pm 50\%$, and all prediction errors were within the range $\pm 100\%$.

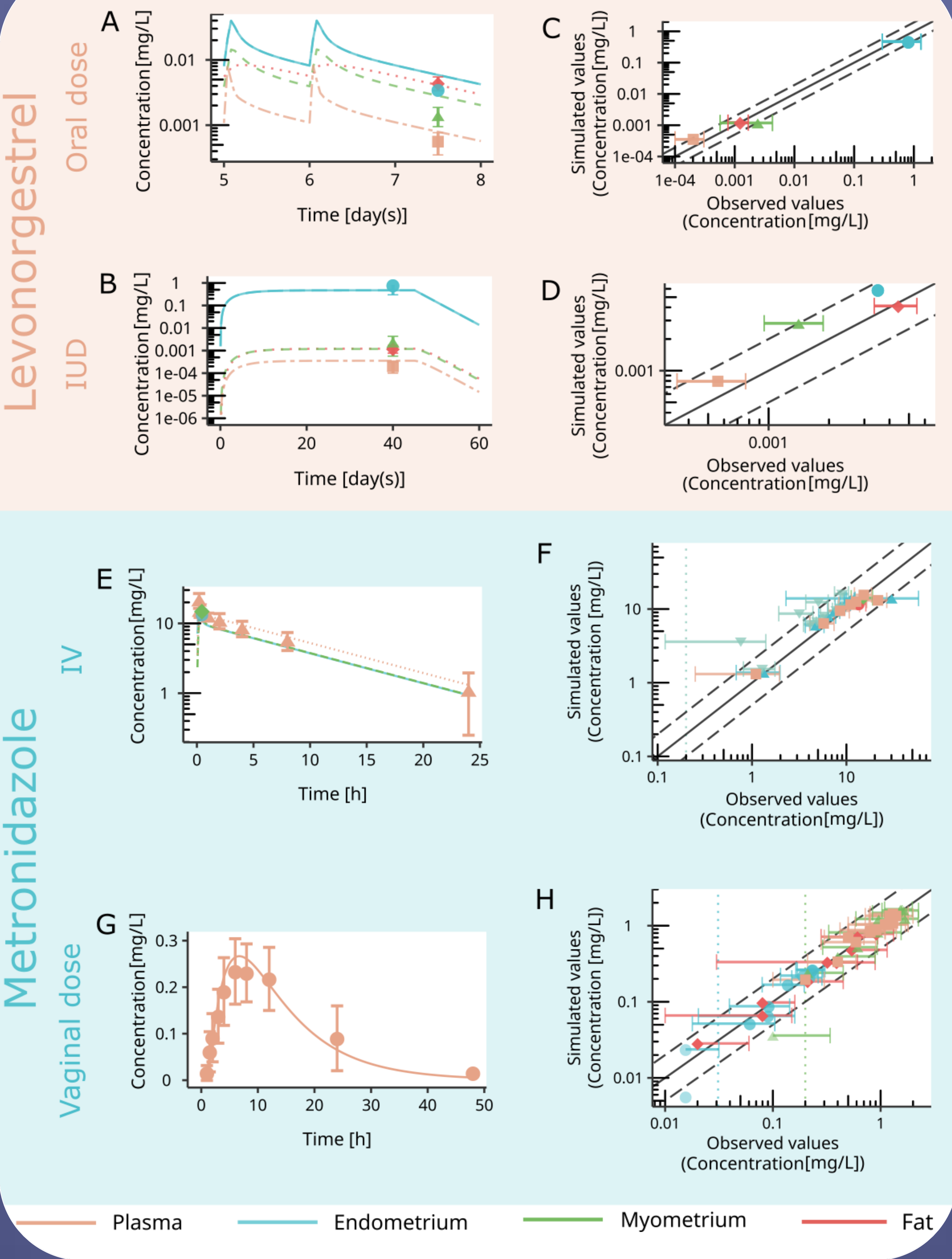
Conclusion

We developed a FRT extension module in MoBi V12 that is openly available on GitHub. The module has been validated for two compounds and six routes of administration. Overall, only minor parameter adjustments were needed to capture clinical data well. Due to the model modularization concept introduced in MoBi V12, the FRT module can easily extend any PBPK base model available in the Open System Pharmacology suite. Hence, this FRT module has significant potential for drug development and safety assessment in female-specific health areas, including infections, cancer, and reproductive toxicity.

Module on GitHub



Explore drug exposure in the organs of the female reproductive tract with our Shiny app



References

[1] Fischer-Holzhausen S., et al. Enhancing PB(P)K Models for the Female Reproductive Tract: A Framework for Local and Systemic Drug Kinetics. PAGE, Greece, Thessaloniki, 2025
[2] Cicali, B., et al. "Quantitative assessment of levonorgestrel binding partner interplay and drug-drug interactions using physiologically based pharmacokinetic modeling." CPT: Pharmacometrics & Systems Pharmacology 10.1 (2021): 48-58.

This work was supported by the Gates Foundation (Grant Number INV-079149).



Modeling



Scaling



Coding

Supporting the open-source development of:

