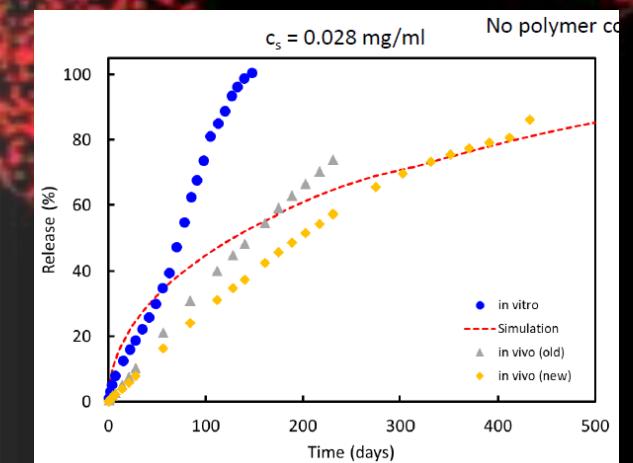
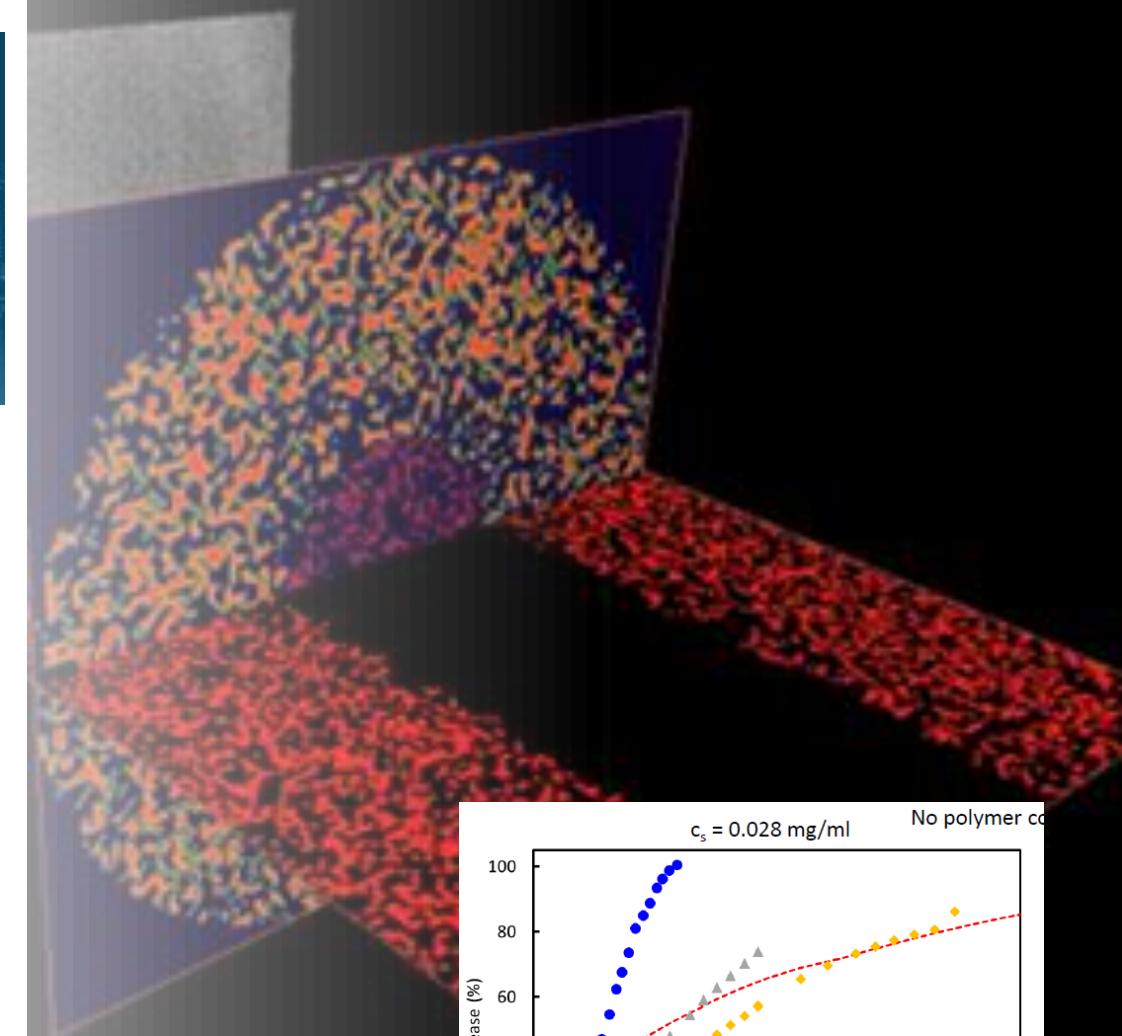


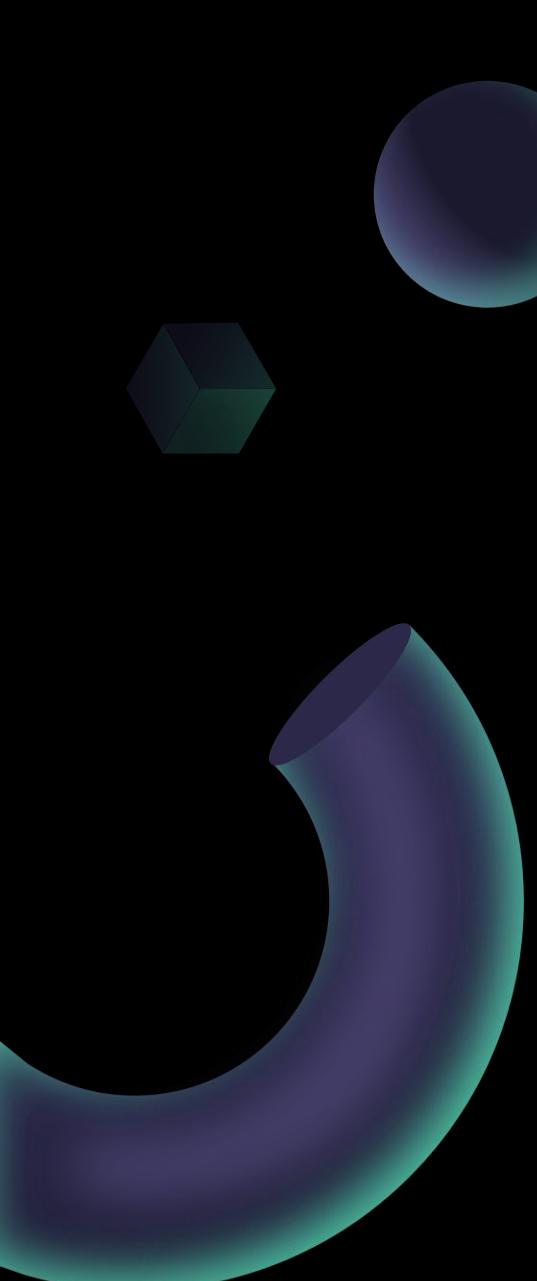


Innovation Showcase

Shortening development cycles
with microstructure-based
prediction of drug release

Dr. Andrew Clark
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DigiM Solution
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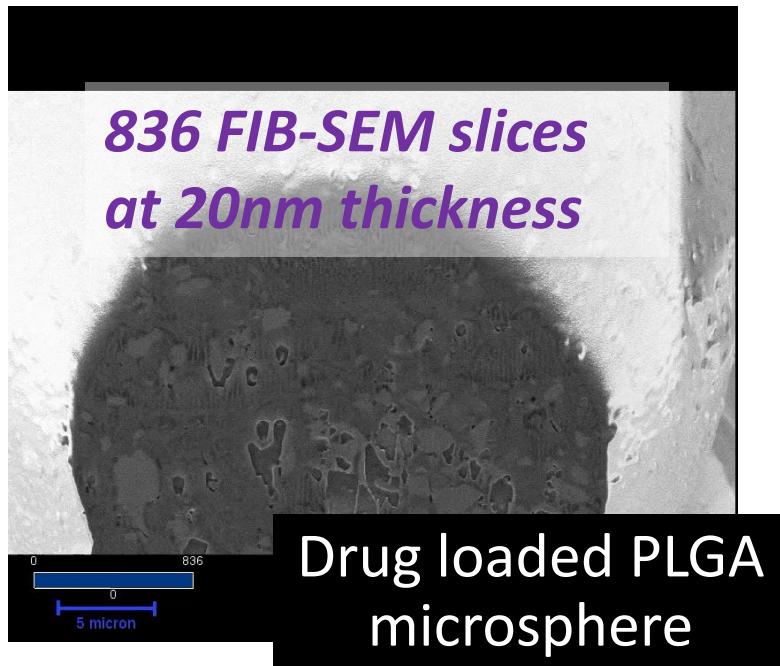




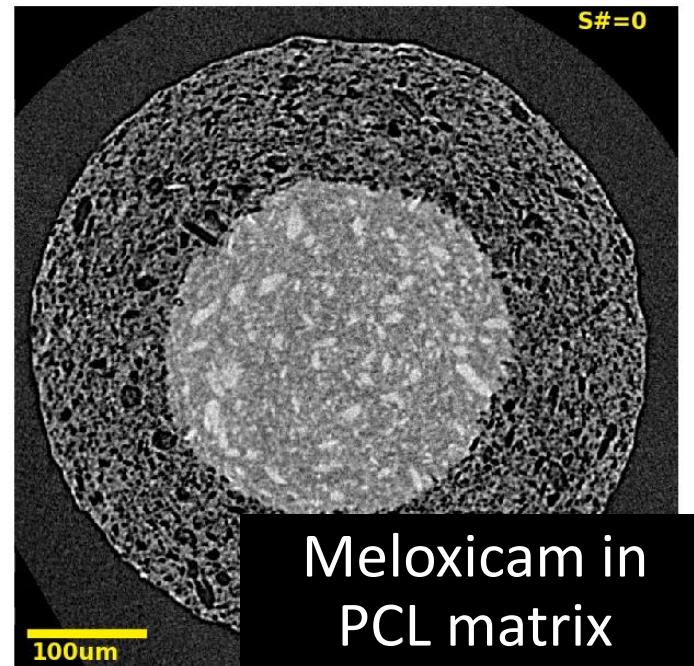
What if we could simulate drug release profiles, based on the actual structures driving release?

We can! It all starts with imaging the structure in 3D...

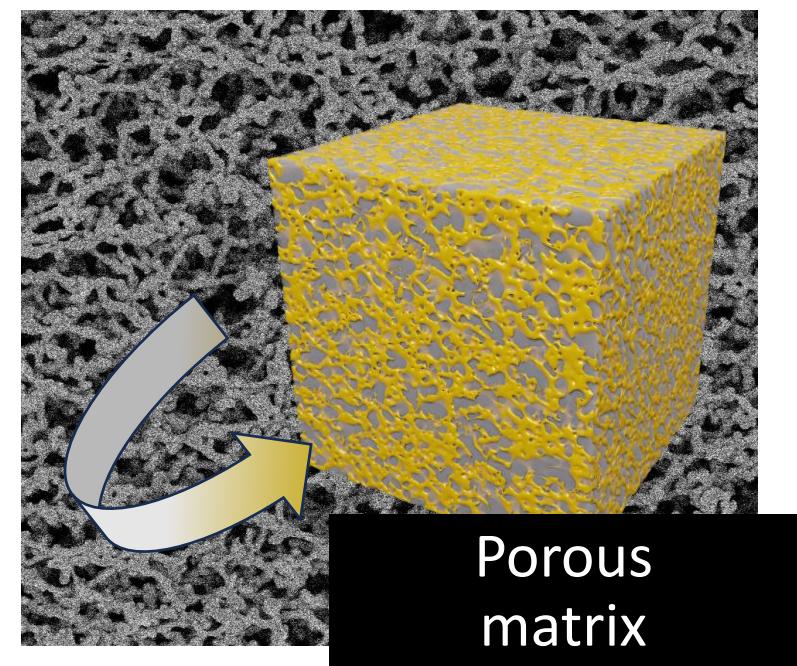
3D FIB-SEM



3D X-ray Microscopy
(X-ray micro-CT)

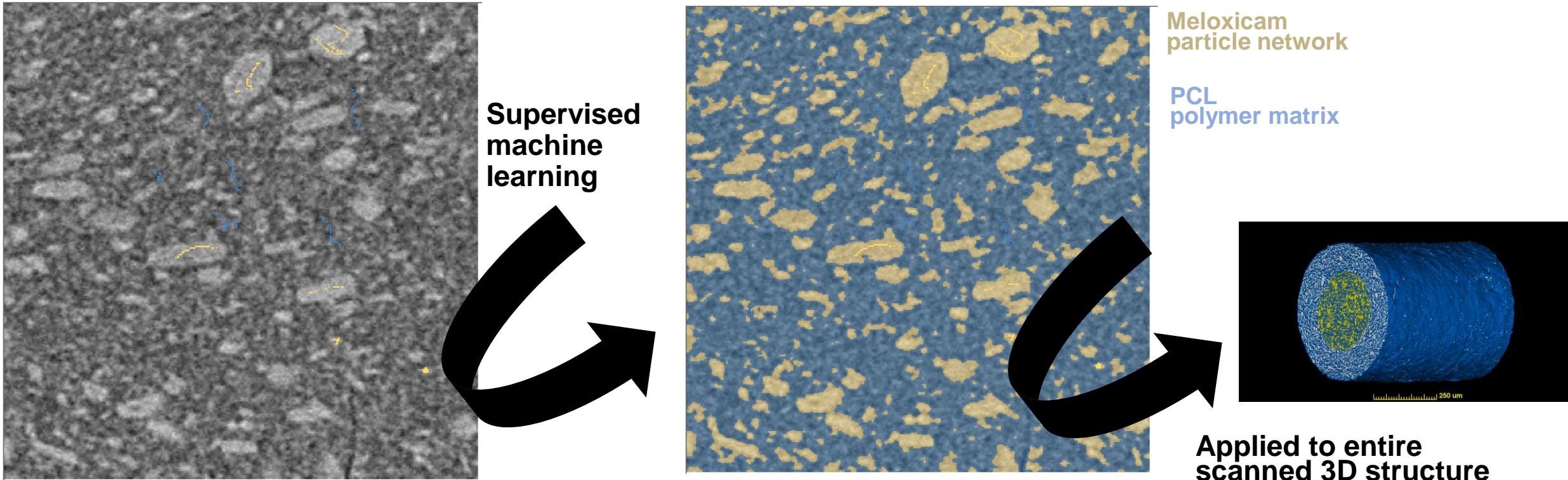


2D to 3D with Generative AI



Obtaining the structure provides a real input without assumptions about porosity/particle arrangement

To simulate from the imaged structure we need to classify each material domain

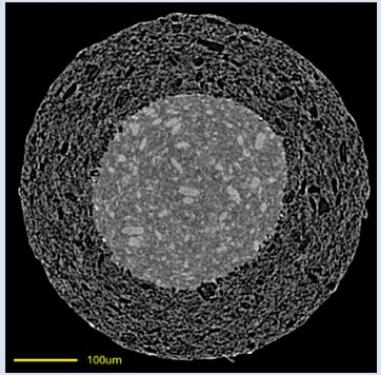


With Purdue and Eli Lilly
<https://doi.org/10.1016/j.jconrel.2023.05.049>

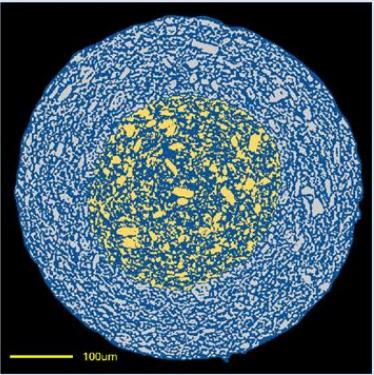
After classification, a library of structural properties can be computed (PSD, spatial distribution, surface area, connectivity)

Lastly, we simulate drug release on the classified pixels

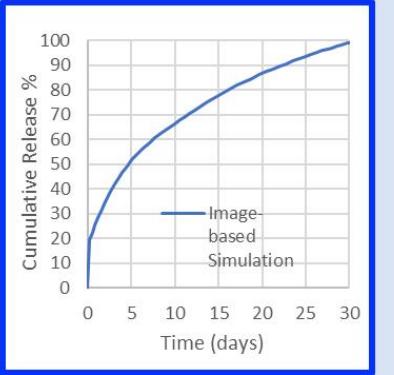
1. XRM Imaging



2. A.I. Image Analytics

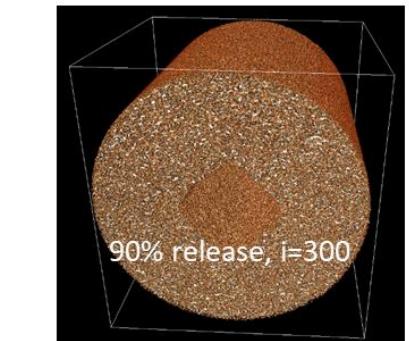


3. Image-based Simulation



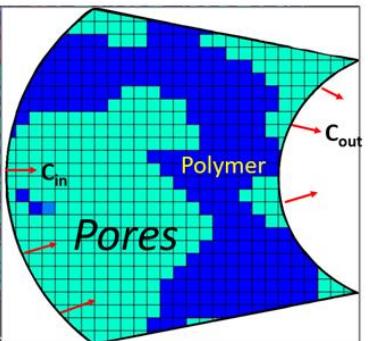
a. Percolation Simulation

Drug Release at Each Step

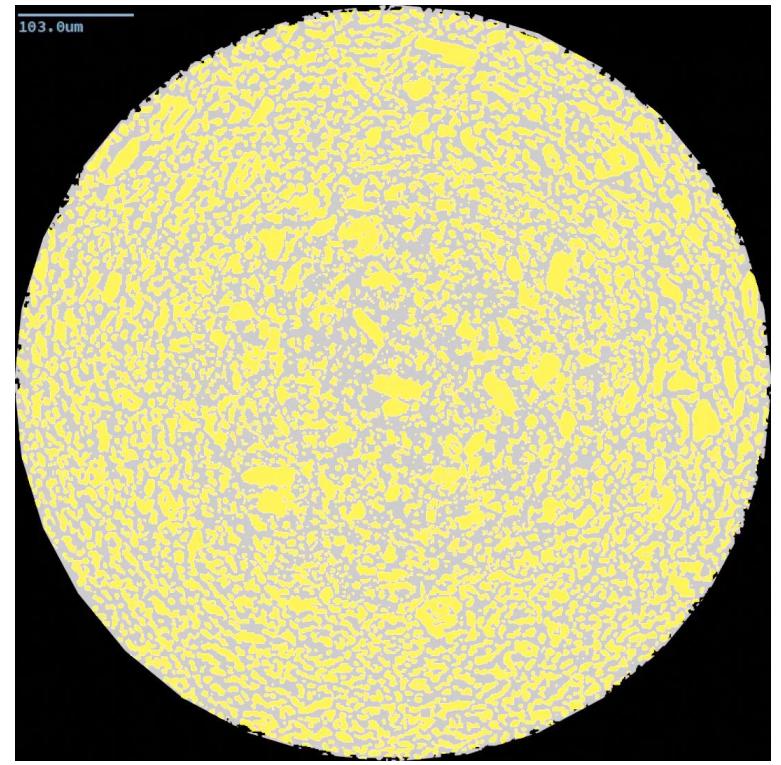
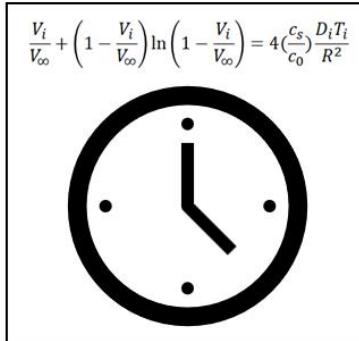


b. Effective Diffusivity

Dynamic Fick's Diffusion



c. Time Conversion

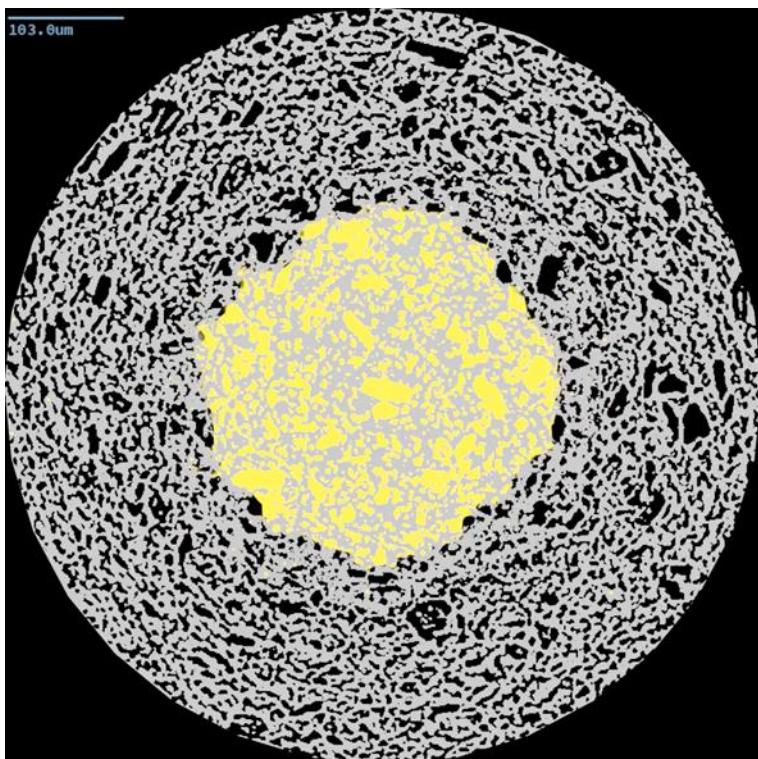


This simulation is on a biostable system → there are also models for biodegradable systems

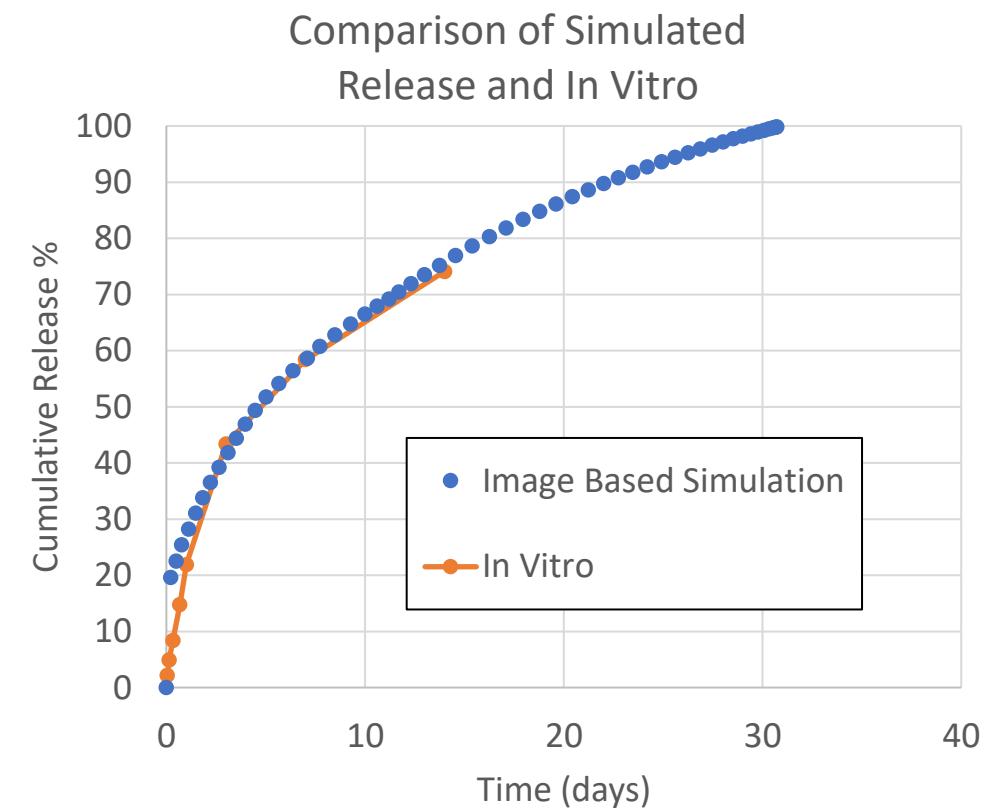
What are the other inputs, and how does predicted release compare to *in vitro*?

External inputs for simulation:

- Bulk diffusion coefficient
- Drug load
- Drug solubility
- Diameter



With Purdue and Eli Lilly
<https://doi.org/10.1016/j.jconrel.2023.05.049>



Blind simulation provides highly precise prediction without any data fitting or adjustments

What are the implications?

Use of this workflow can help solve a number of the biggest challenges in controlled release development...

Accelerate Formulation Selection

Optimize Manufacturing and Scale Up

Shorten Lengthy *In Vitro* Testing

Easier Troubleshooting

Seek *In vitro-In Vivo* Correlation

Demonstrate Generic Equivalence with RLD