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Comparison of Classical and Hybrid Methodologies for Model-based Design of Experiments

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Outline

- Introduction and motivation for investigating model-based design of experiments (MBDoE)
- Benchmark development
 - Ground truth CHO model
 - Iterative test strategy
- Results
 - Classical DoE (resolution III and IV fractional factorial, quadratic response surface)
 - DataHow hybrid model + (Latin hypercube, model-based design of experiments)
- Conclusions



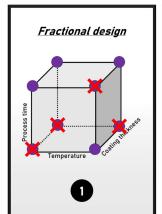
Introduction and Motivation

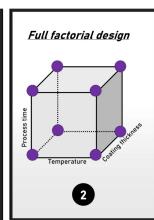
- Key mandate: add value to CMC development activities through data science
- How to get value from models (hybrid or otherwise)?
 - Reduction in number or cycles of experiments to get acceptable titer and product quality (PQ)
 - Improvement of titer, PQ, reduction of COGS through numerical optimization
- Goal: Speed up learning and optimization cycle through model-based design of experiments

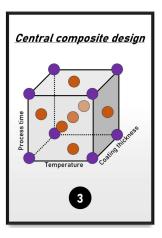


State of the art: Traditional DoE

- Traditional DoE is the standard of practice
 - Fractional factorial and response surface methodologies (RSM)
 - Process scientists pick relevant factors and ranges
 - Design experiments based on classical "canned" designs
 - Experiments allow for factors to be refined
 - Higher resolution experiments may be designed for further refinement
- Benefits: Causal identification of input-output relationships
- Limitations of these models
 - Limited representation of nonlinearity and dynamics
 - Input-output relationship ignores mass balance and cell-specific rates
 - Model uncertainty not directly handled (indirect factor screening)







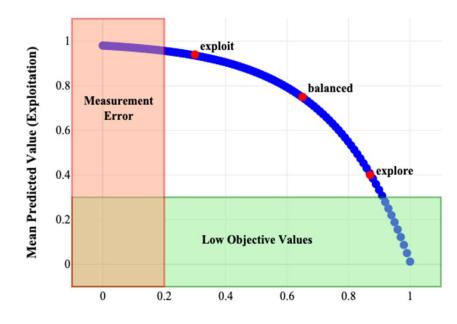
https://www.experimentaldesignhub.com/blog/advanced-doe-plans-part-1



Moving to a model-based DoE strategy



- Approach within DataHow
 - Latin hypercube sampling (LHS) for initial set
 - Multi-objective optimization to balance between uncertainty reduction (explore) and optimization (exploit)
- Benefits
 - Use of bioreactor mass balances converts problem into a smoother cell-specific derivative space
 - Naturally handles dynamics
 - Direct uncertainty description
- Unknowns
 - How do they compare to traditional DoE?



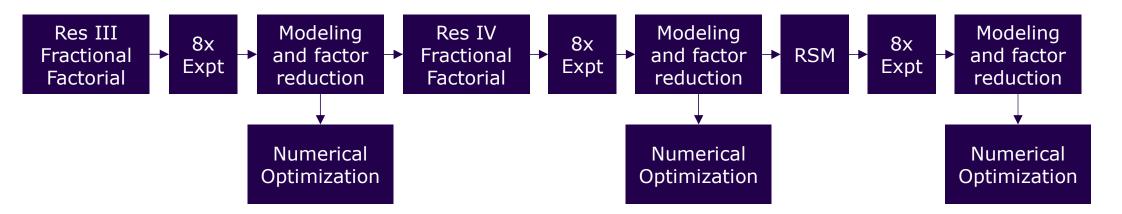
Standard Deviation (Exploration)



Ideal Empirical Strategy

Does the DataHow hybrid MBDoE outperform classical DoE in a media optimization setting?

Control:



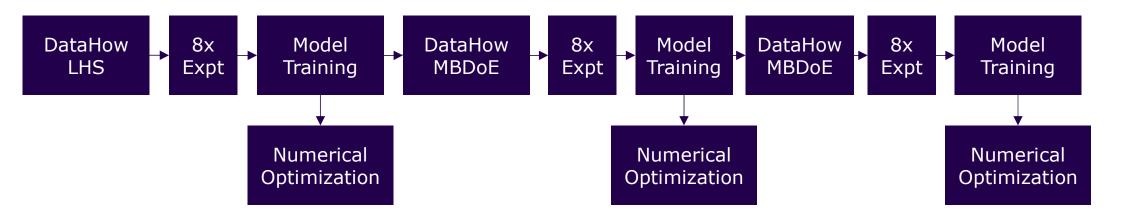
Record final optimized titer at each stage



Ideal Empirical Strategy

Does the DataHow hybrid MBDoE outperform classical DoE in a media optimization setting?

Treatment:



Record final optimized titer at each stage



Challenges with Empirical Comparison

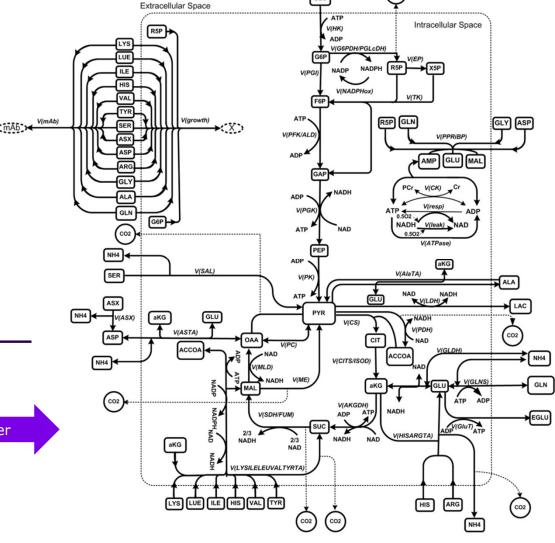
- High experimental burden (3 cycles * 8 experiments * 2 arms) for a single comparison
- Difficult to study non-ideal behavior
 - Analytical and biological variability
 - Block effects
 - Contamination events
- No access to ground truth, difficult to explain why the models behave the way they do
- Solution: Develop a ground truth model that replicates experimental work
 - Model does not need to accurately replicate Sanofi cell culture results
 - Only needs to capture representative dynamics, nonlinearity, and smoothness
 - Model can be used for evaluation of future model-based DoE tools



Ground Truth Model

Considered a combination of two closely related models from literature (Robitaille/Ghorbaniaghdam) modeling mAb production from CHO cells.

Modified to be fed-batch system, include cell death, and extended to 14 days in order to be closer to real world cell cultures.



Media/Feed Concentrations

Ground Truth Model (GTM)

mAb Titer



-) 2015, Robitaille, et al.; PLOS One
- 2) 2014, Ghorbaniaghdam, et al.; PLOS One

Ground Truth Model (Model Rates)

Reaction: GLC+ATP→G6P+ADP

$$v(HK) = v_{\text{max }HK} * \frac{GLC}{K_{mGLC} + GLC} * \frac{\frac{ATP}{ADP}}{K_{m\frac{ATP}{ADP}} + \frac{ATP}{ADP}} * \frac{K_{iG6P}}{K_{iG6P} + G6P}$$
Inhibition Terms

Reaction: PEP+ADP→PYR+ATP

$$v(PK) = v_{\text{max }PK} * \frac{PEP}{K_{mPEP} * \left(1 + \frac{Ka_{F6P}}{F6P}\right) + PEP} * \frac{\frac{ADP}{ATP}}{K_{m\frac{ADP}{ATP}} + \frac{ADP}{ATP}}$$

► Activation Terms

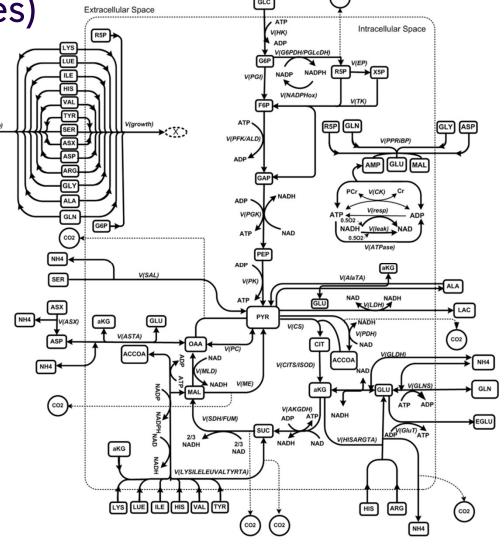
Reaction: 0.01GLU +0.01GLN+...+0.0145GLY+4ATP→mAb

$$v(mAb) = v_{maxmAb} * \frac{GLN}{K_{mGLNmab} + GLN} * \cdots * \frac{GLY}{K_{mGLY} + GLY} \frac{\frac{ATP}{ADP}}{K_{m\frac{ATP}{ADP}} + \frac{ATP}{ADP}}$$

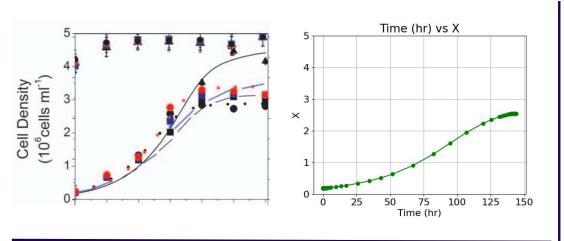
→ 16 Michaelis-Menten Terms

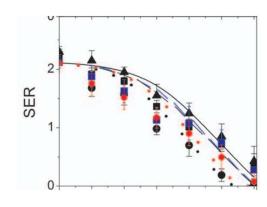


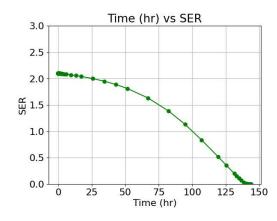
- 1) 2015, Robitaille, et al.; PLOS One
- 2) 2014, Ghorbaniaghdam, et al.; PLOS One

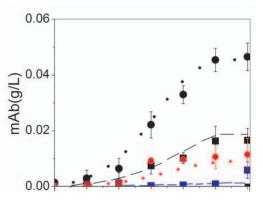


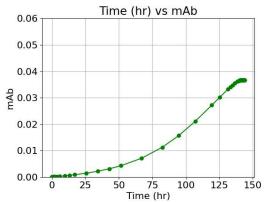
Ground Truth Model matches literature data



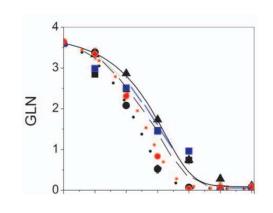














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Ground Truth Model (Modifications)

Modifications required to better reflect commercial cell culture

- Changing from a batch model to a bolus fed-batch model
- Cell death based on ammonia and lactate concentrations.

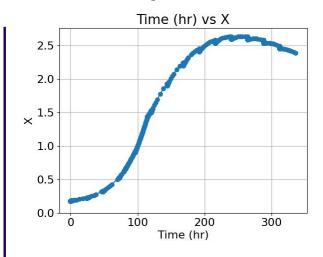
$$\mu_d = k_d \cdot \frac{[LAC]}{KD_{lac} + [LAC]} \cdot \frac{[AMM]}{KD_{amm} + [AMM]}$$

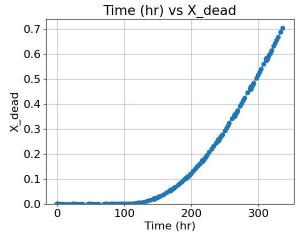
where

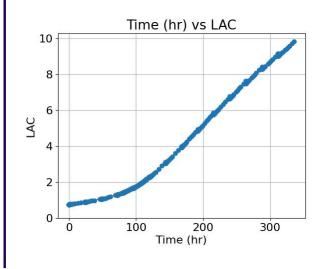
$$\begin{cases} \textit{if} \ [AMM] \leq 4.6 \ mM \ then \ KI_{amm} \gg [AMM], \\ KD_{amm} \gg [AMM] \\ \textit{if} \ [LAC] \leq 52 \ mM \ then \ KI_{lac} \gg [LAC], \ KI_{lac} \gg [LAC] \end{cases}$$

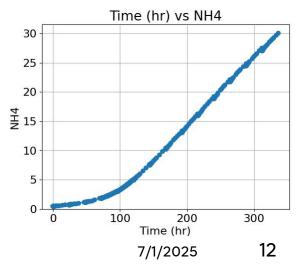
3) 2009, Xing, et al.; Biotechnology Progress

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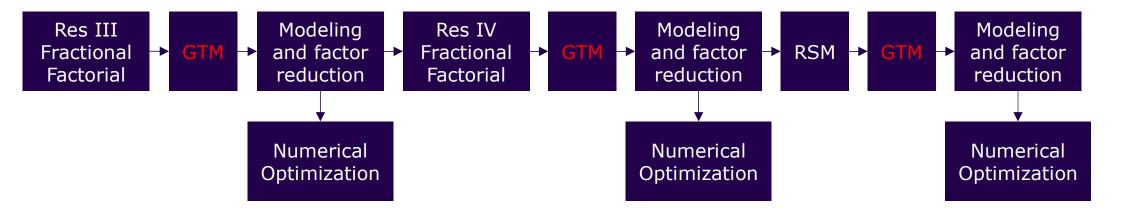




Actual Empirical Strategy

Does the DataHow hybrid MBDoE outperform classical DoE in a media optimization setting?

Control:

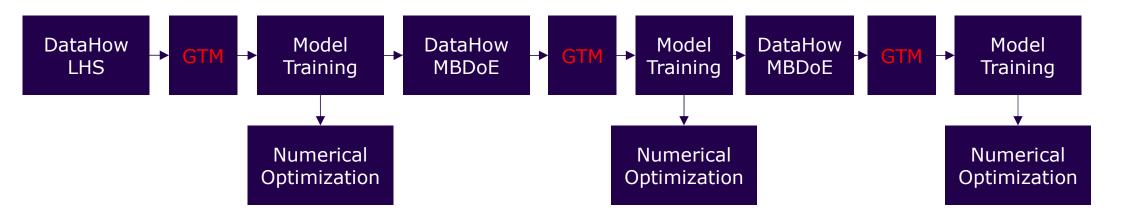




Actual Empirical Strategy

Does the DataHow hybrid MBDoE outperform classical DoE in a media optimization setting?

Treatment:





DoE Evaluations

DoE Parameter	Low	High
Feeding Start Time (hrs)	24	312
Basal Glutamine (mM)	0.01	7.2
Feed Glutamine (mM)	0.05	36
Glucose Setpoint (mM)	0.01	56
Basal Alanine (mM)	0.01	1.5
Feed Alanine (mM)	0.05	7.5

Choice of Parameter Ranges

- Glucose, glutamine, and alanine had interesting, nonlinear behavior
- Studied ranges of 0 2x of Gorbaniaghdam conditions
- Feed was 5x of the media conditions

Evaluation Criteria

- (Highest) sampled titer:
 - GTM titer across the batch of 8 experiments
- Model-optimized titer
 - Classical/hybrid model trained and optimized to suggest single optimal run
 - GTM titer for that single optimized run

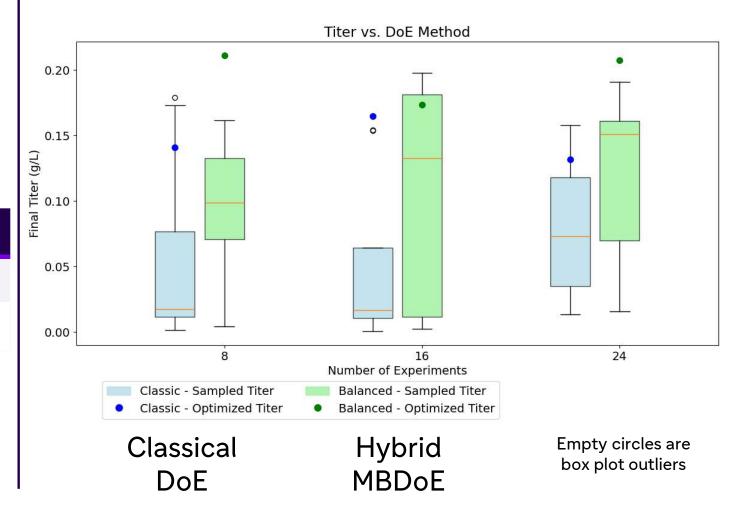


Results

Overall Results

 Hybrid-balanced approach has better final titers at each stage of design

Titer (g/L)	Highest Sampled	Model Optimum
Classic	0.179	0.165
Hybrid (Balanced)	0.198 (+11%)	0.211 (+28%)



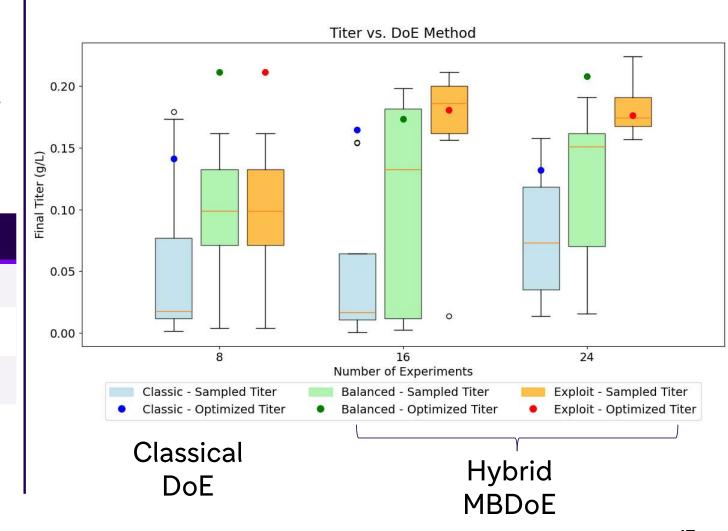


Results

Overall Results

 Hybrid-exploit approach improves sampled titer at each stage of design

Titer (g/L)	Highest Sampled	Model Optimum
Classic	0.179	0.165
Hybrid (Balanced)	0.198 (+11%)	0.211 (+28%)
Hybrid (Exploit)	0.224 (+25%)	0.211 (+28%)

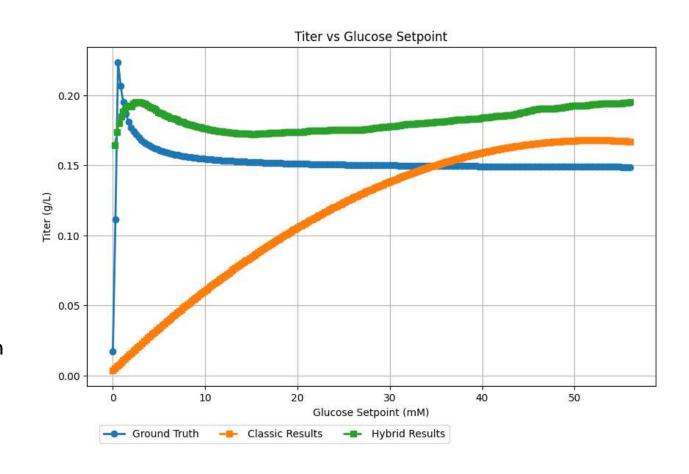




Results

Model System Mismatch Analysis

- Glucose setpoint has a very sharp peak near the optimum
- Classical models cannot capture that behavior using quadratic RSM models
- Hybrid methodology has the necessary complexity to pick up on the peak like behavior, resulting in better experiments.





Conclusions

- Role of modeling in CMC development for cell culture
 - Get to better processes faster
- Design of experiments using hybrid models is a rational approach
 - Cell-specific rates, mass balance, uncertainty description
- Developed a ground truth mechanistic model to evaluate DoE methodologies
 - Does not need to replicate in-house processes, just needs to capture smoothness and nonlinearity
- Hybrid model-based methodology outperforms classical DoE under the benchmark test
- Next steps
 - Experimental implementation on assets
 - Impact of non-ideal behavior (variability, contamination)



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