



**PYOMO  
DOE**

# **Science-based Design of Experiments**

**Alexander (Alex) Dowling**

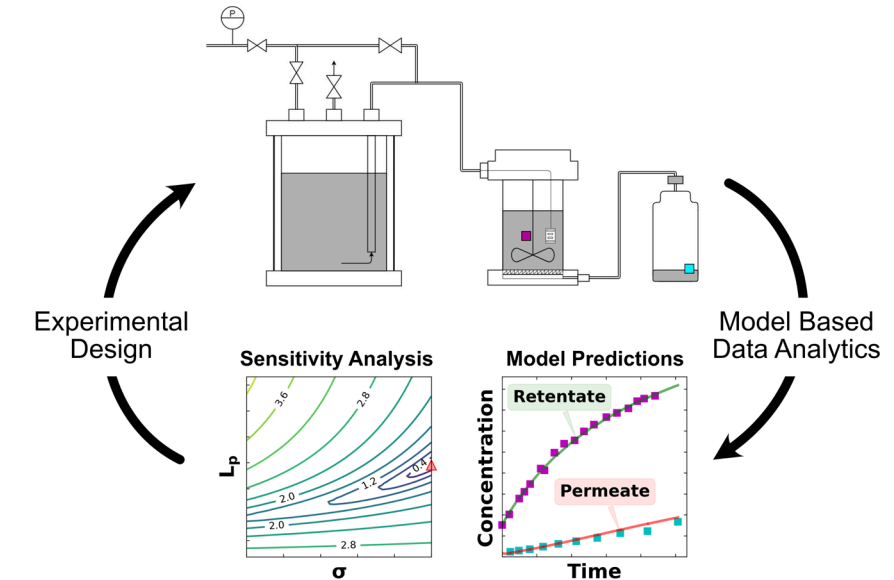
adowling@nd.edu

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Advanced PSE Workshop, Washington D.C.

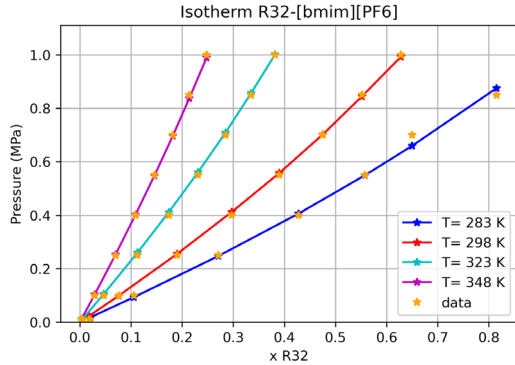
# Presentation Goals

1. When to use science-based (model-based) design of experiment?
2. Illustrate key ideas of SBDoE using membrane example
3. Summarize `Pyomo.DoE` and development plan



# What data are most valuable to optimize systems?

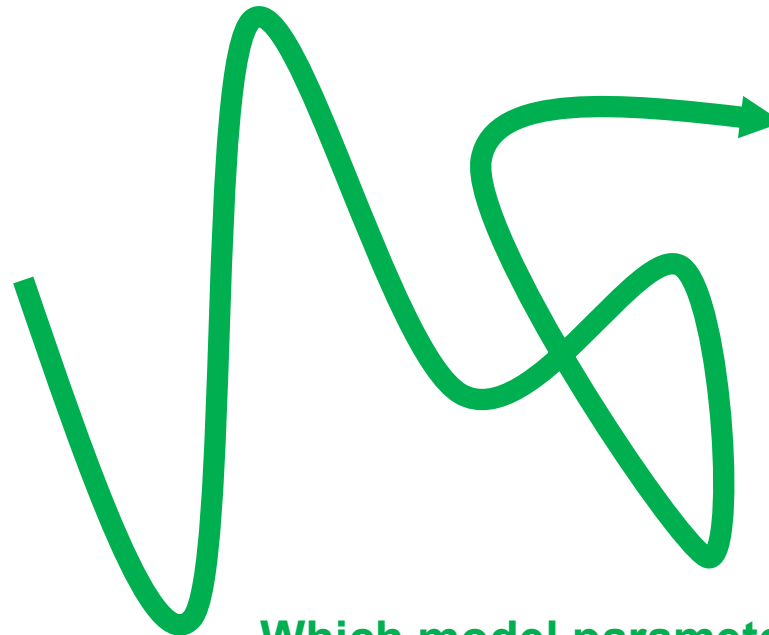
Experimental thermophysical property measurements



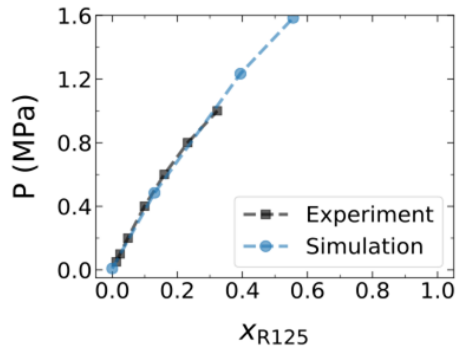
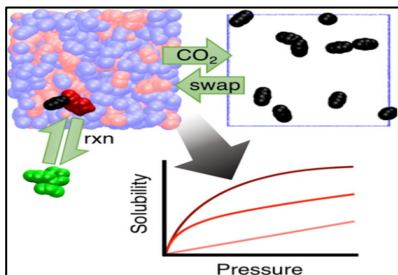
What uncertainties are acceptable?

System optimization, e.g., design, operation, control

$$\begin{aligned} \min \quad & f(x, y) \\ \text{s.t.} \quad & g(x, y) \leq 0 \\ & h(x, y) = 0 \\ & x \in \mathbb{R}^n, y \in \mathbb{Z}^m \end{aligned}$$

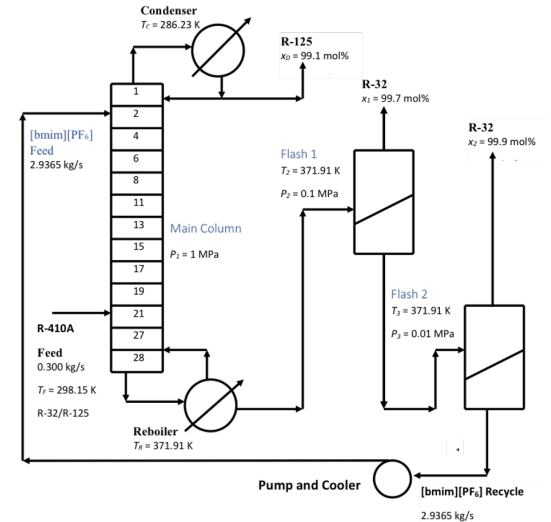


Molecular simulation predictions



Which model parameters to calibration from data?

How can optimization help straighten this tortuous path?



Beftor et al (2021), *J. Chem. Inf. Model*, 61(9), 440-4414.

Shiflett, M. B. and Yokozeki, A. *Chimica Oggi - Chemistry Today*, 24(2). (2006).  
Devaki, S. J. and Sasi, R. *IntechOpen*. (2007).

# Design of Experiments $\neq$ One-Factor-at-a-Time

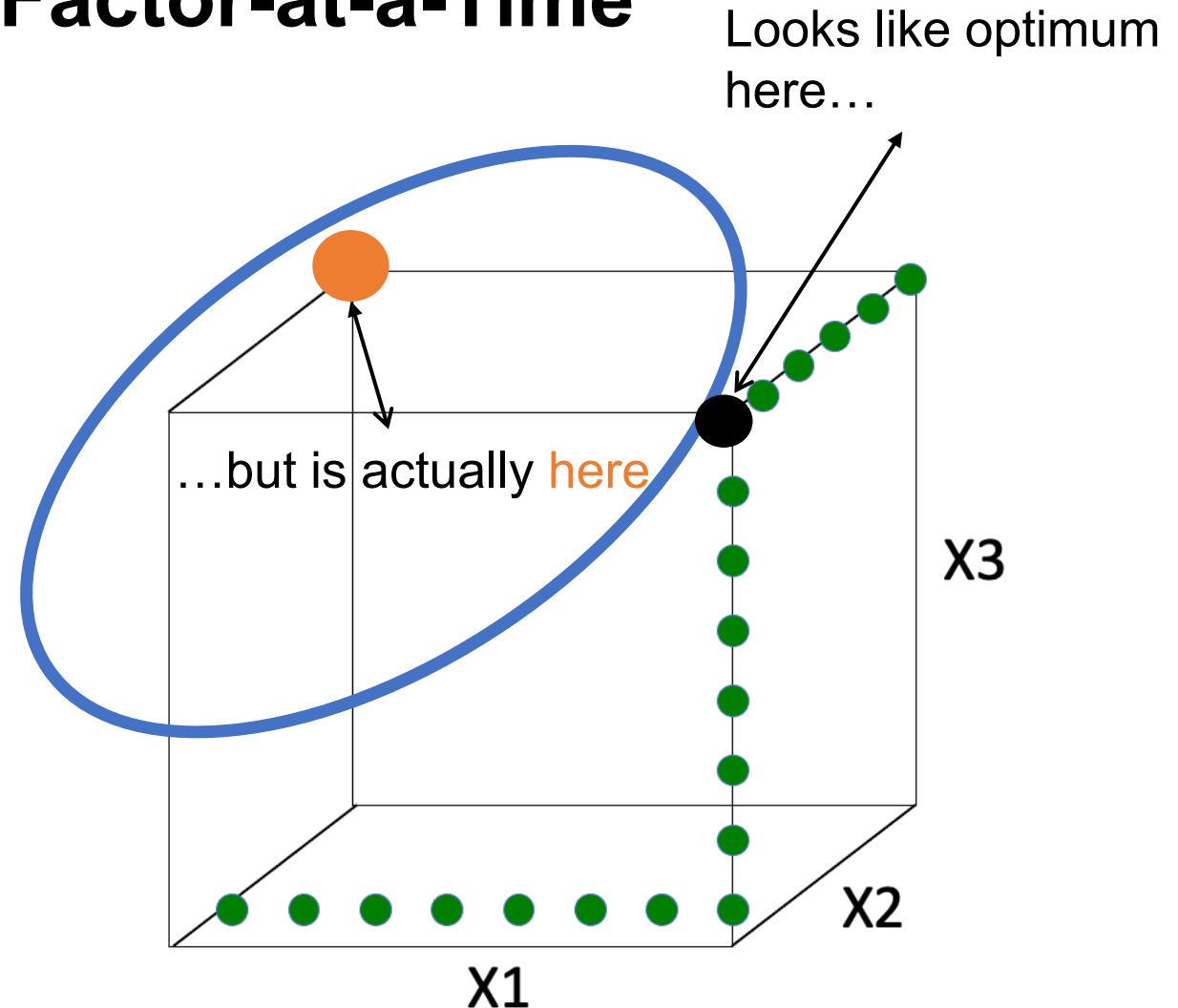
## OFAAT strategy:

- Change only one input (factor) at a time
- Hold all others constant

## Inefficient use of budget

## Cannot identify interactions

- Effect of one factor changes when another factor changes
- Finding optimal operating conditions is unlikely



# Design of Experiments $\neq$ One-Factor-at-a-Time

## OFAAT strategy:

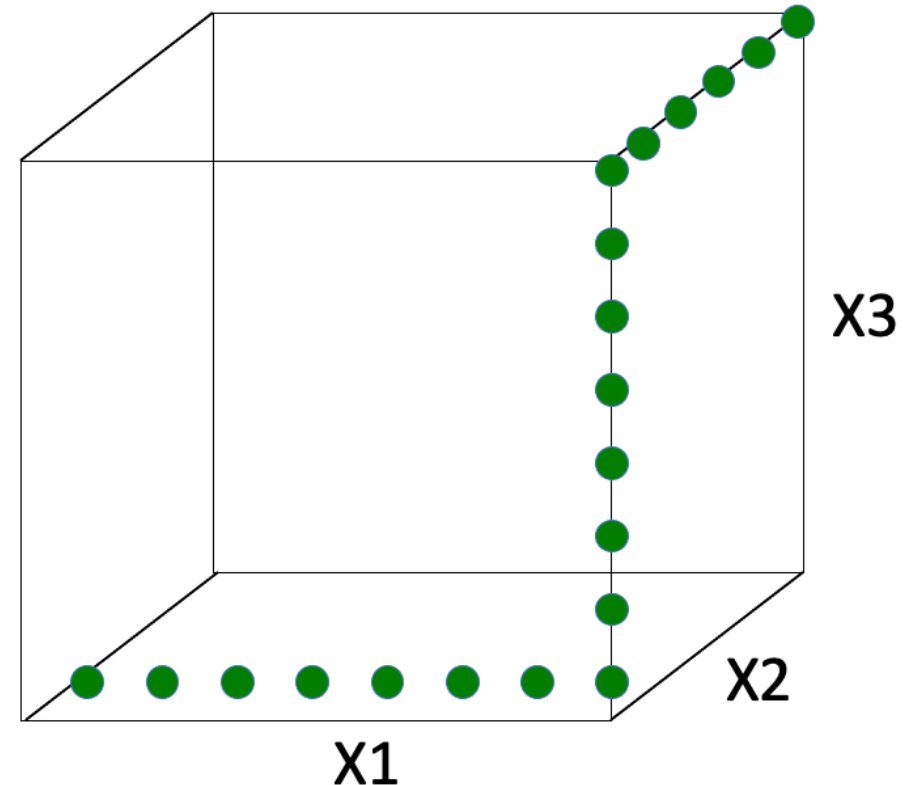
- Change only one input (factor) at a time
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Inefficient use of budget

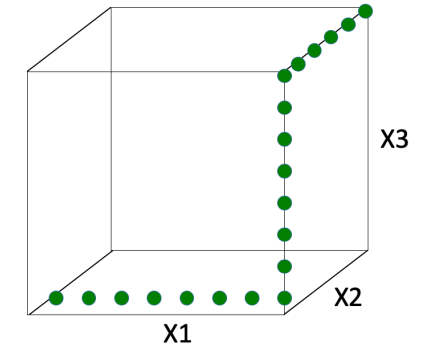
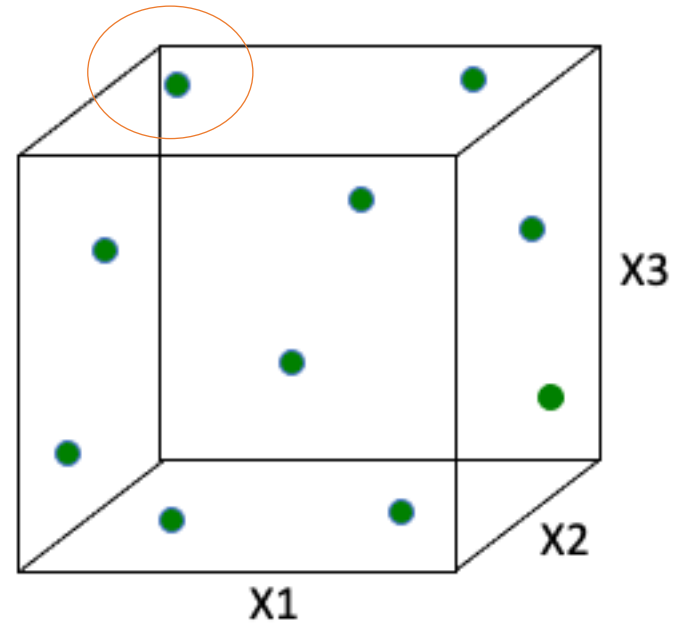
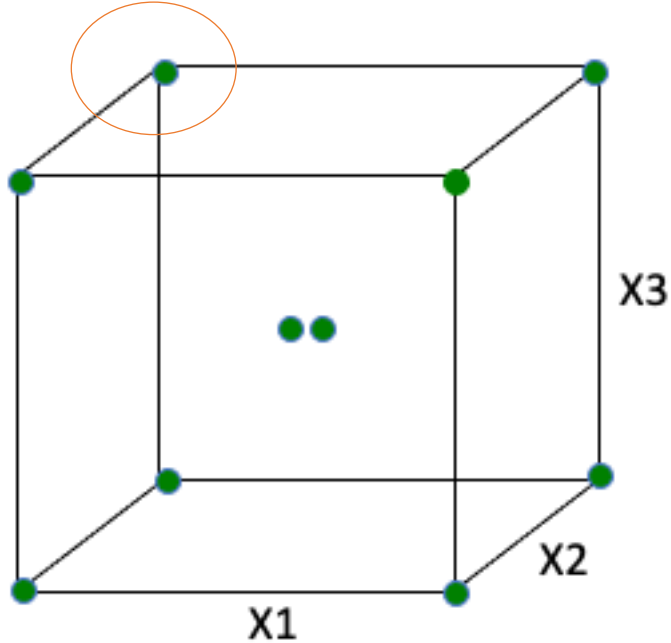
Cannot identify interactions

## Not randomized

- Changing conditions can negatively affect the results



# Sequential DoE Avoids These Drawbacks and Is Always More Efficient



Uses **20 runs**

Two Different Sequential DoE Approaches  
Each uses **10 runs**

# Power of Adaptive Sequential Optimal Experiments

## Self-Driving Laboratories

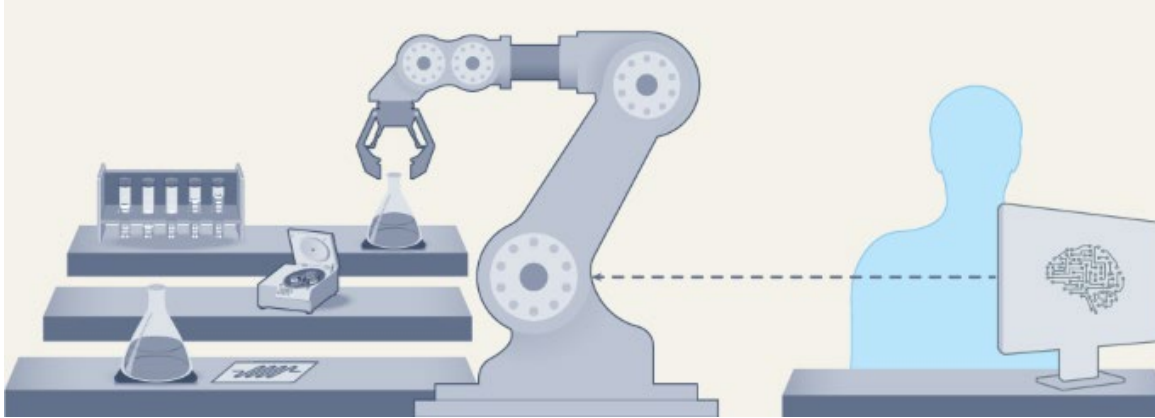


Figure: Abolhasani & Kumacheva (2023), *Nature Syn.*

Epps et al. (2022), *Advanced Materials*

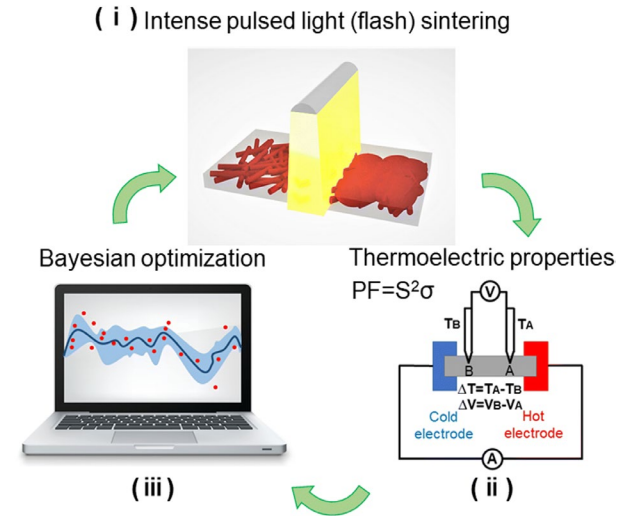
MacLeod et al. (2020), *Science Advances*

MacLeod et al. (2022), *Nature Communications*

Hase, Roch, Aspuru-Guzik (2019), *Trends in Chemistry*

Seifrid et al. (2022), *Acc. Chem. Res.*

## Bayesian Optimization



Saeidi-Javash et al.  
(2022), *Eng. Env. Sci*

## CO<sub>2</sub> Capture Technology Scale-Up



Morgan et al (2020),  
*Applied Energy*

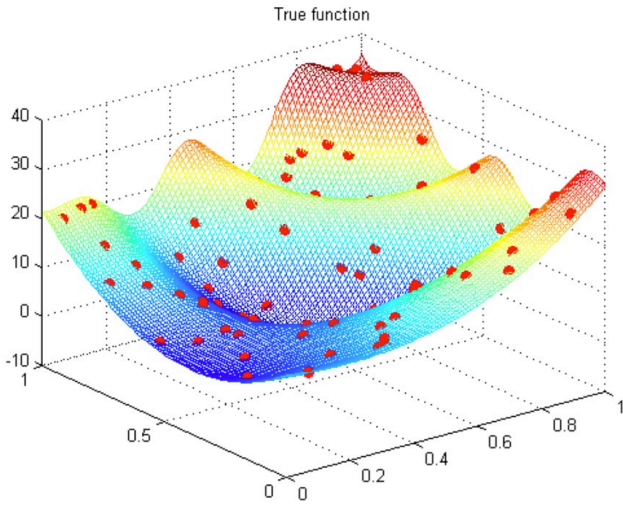
Photo: National Carbon  
Capture Center (AL)



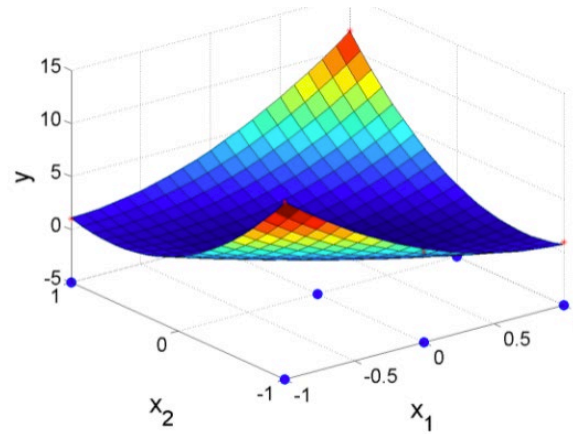
# Optimizing Computational & Physical Experiments

## Classical DoE

### Space Filling



### Response Surface

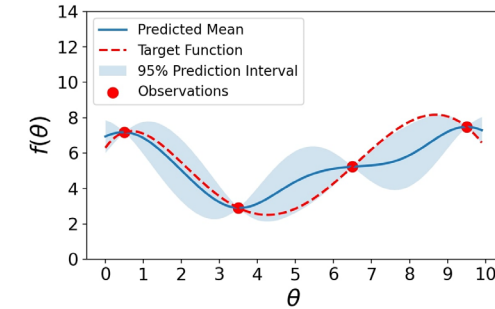


+ Decades of success,  
staple of applied statistics

- Static/fixed design that is not updated  
as data are collected

## Modern (Adaptative) DoE

### Bayesian Optimization



with a data-driven model

### Model-based

$$\rho C_p \frac{\partial T(t, X)}{\partial t} + \rho C_p V \cdot \nabla T(t, X) + \nabla \cdot (-k \nabla T(t, X)) = Q$$

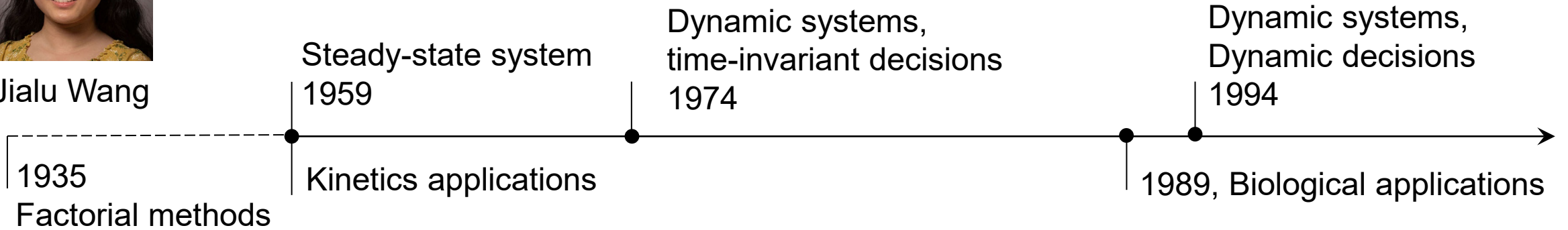
with a scientific model



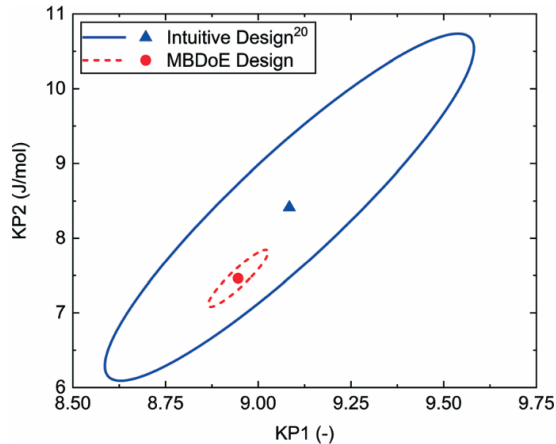
# Science-based (Model-Based) DoE History



Jialu Wang



MBDoe parameter confidence regions are...  
84% smaller than intuitive design  
40% smaller than factorial design



Waldron, C., et al., 2020, *Reaction Chemistry & Engineering*, 5(1), pp.112-123.

## Robust Formulations:

Bruwer, M.J. and MacGregor, J.F., 2006. Robust multi-variable identification: Optimal experimental design with constraints. *Journal of Process Control*, 16(6), pp.581-600.

Dette, H., Melas, V.B., Pepelyshev, A. and Strigul, N., 2005. Robust and efficient design of experiments for the Monod model. *Journal of theoretical biology*, 234(4), pp.537-550.

## Online (Automated) MBDoE:

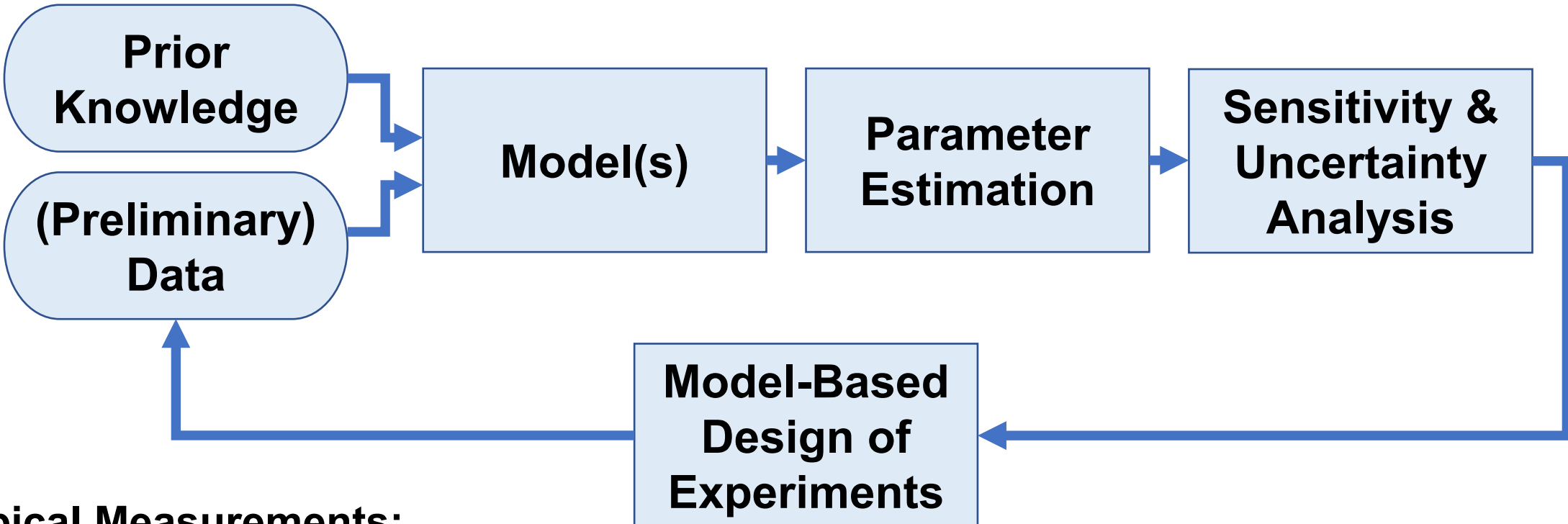
Quaglio, M., Waldron, C., Pankajakshan, A., Cao, E., Gavriilidis, A., Fraga, E.S. and Galvanin, F., 2019. An online reparametrisation approach for robust parameter estimation in automated model identification platforms. *Computers & Chemical Engineering*, 124, pp.270-284.

Waldron, C., Pankajakshan, A., Quaglio, M., Cao, E., Galvanin, F. and Gavriilidis, A., 2019. Closed-loop model-based design of experiments for kinetic model discrimination and parameter estimation: Benzoic acid esterification on a heterogeneous catalyst. *Industrial & Engineering Chemistry Research*, 58(49), pp.22165-22177.

## Recommended Review:

Franceschini, G. and Macchietto, S., 2008. Model-based design of experiments for parameter precision: State of the art. *Chemical Engineering Science*, 63(19), pp.4846-4872.

# Science-based Modeling Workflow



### Typical Measurements:

- Pressures
- Temperature
- Composition
- Flowrate

**Typical Models: Partial Differential Algebraic Equations**  
Transport                      Thermodynamics                      Reactions

# DATA: Diafiltration Apparatus for high-Throughput Analysis



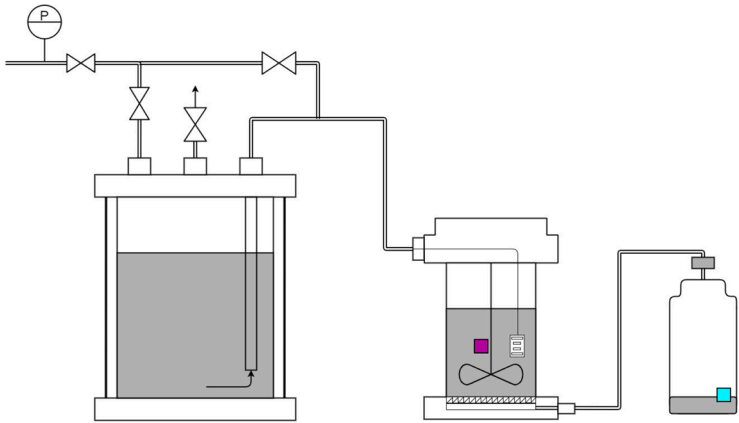
The research community should use **data science**, modeling, and simulation with **experimental measurements** to develop a fundamental understanding of separation materials in **complex environments** and at multiple scales.



**William Phillip**



**Alexander Dowling**



Experimental Design

Model Based Data Analytics

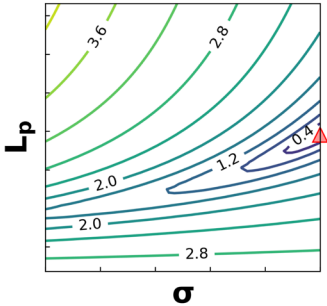


**Jonathan Ouimet**

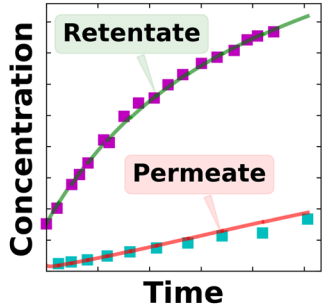


**Xinhong Liu**

Sensitivity Analysis



Model Predictions

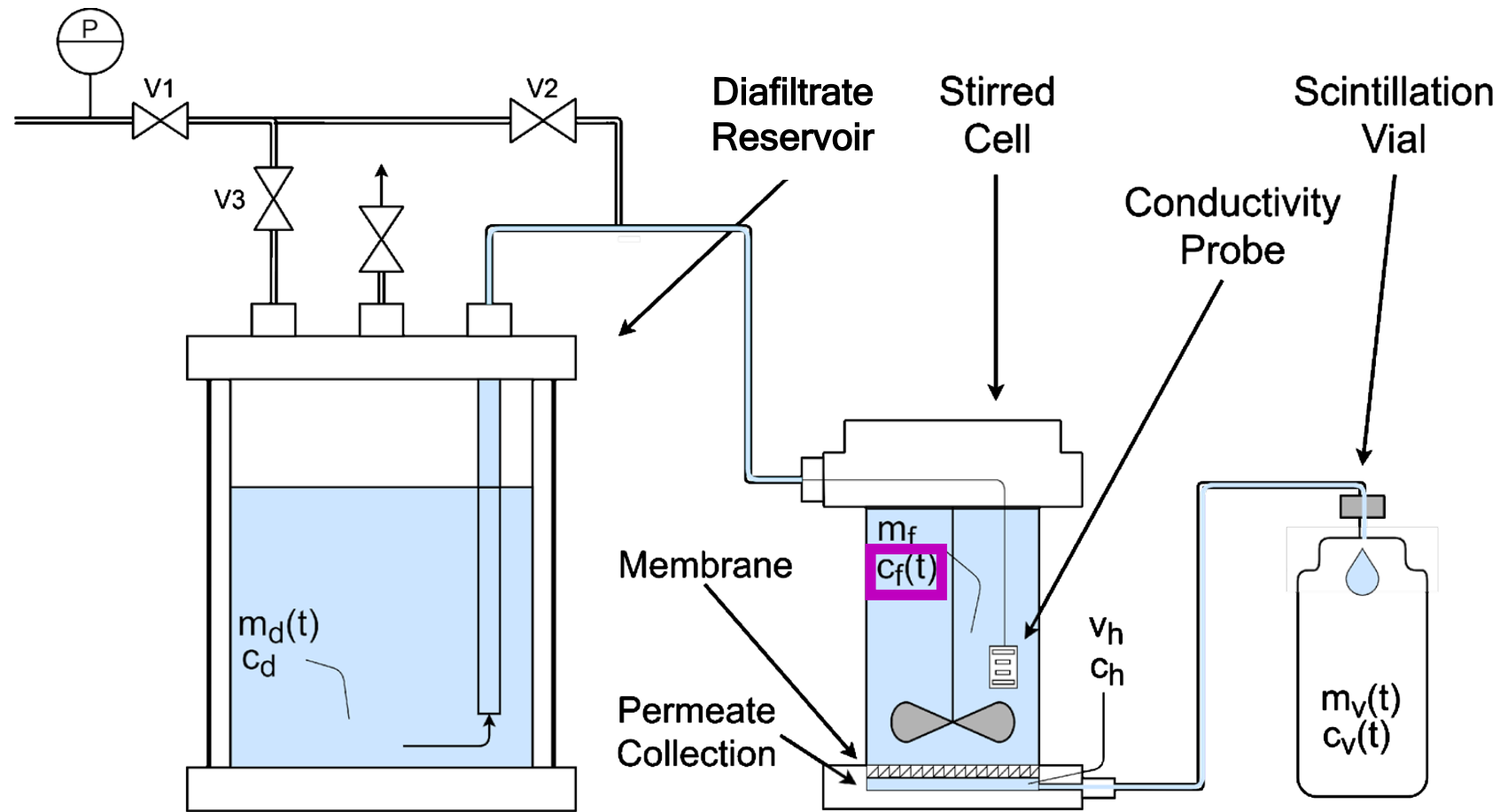


How to develop, accelerate, and improve the membrane characterization technique?  
 How to rapidly and precisely elucidate governing phenomena and mechanisms?

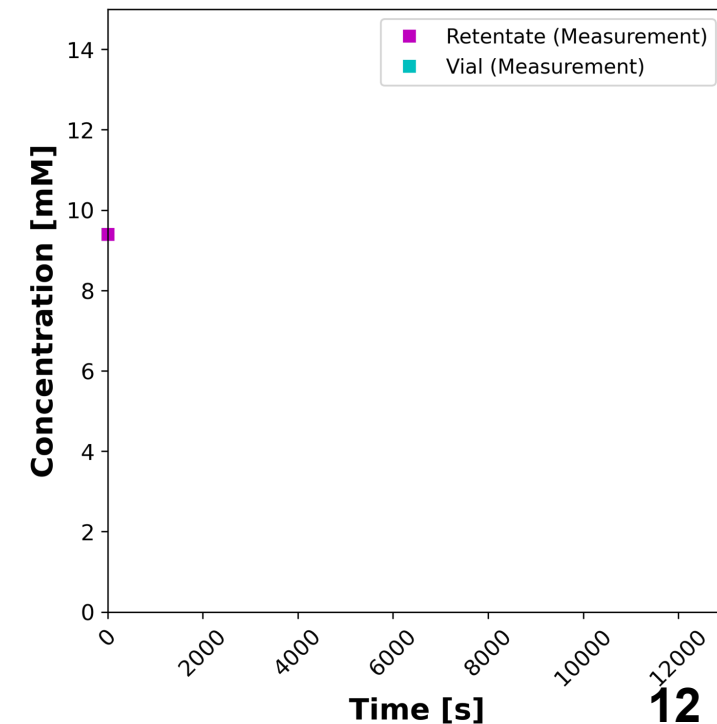
[Left Fig.] National Academies of Sciences, Engineering, and Medicine. (2019). The National Academies Press.  
 [Right Fig.] Ouimet, J. A. et. al., (2022). Journal of Membrane Science, 641, 119743.

# Dynamic Diafiltration Experiment Sweeps Larger Concentration Space

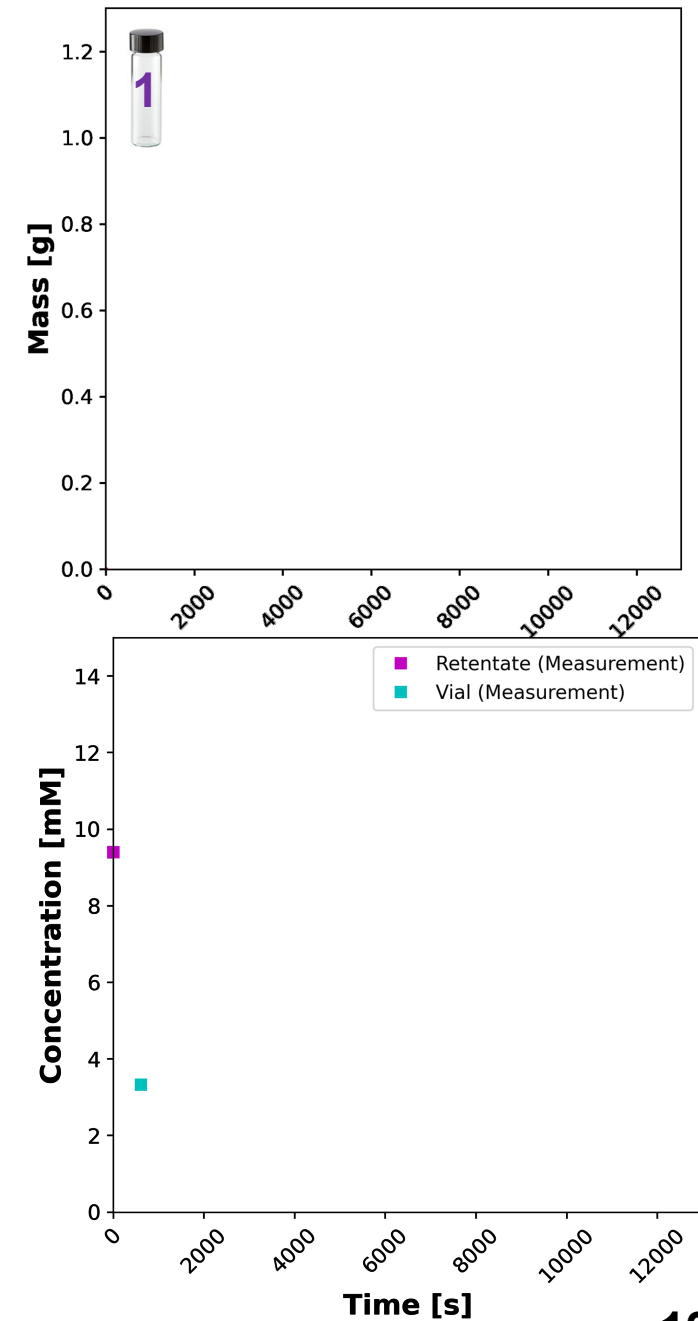
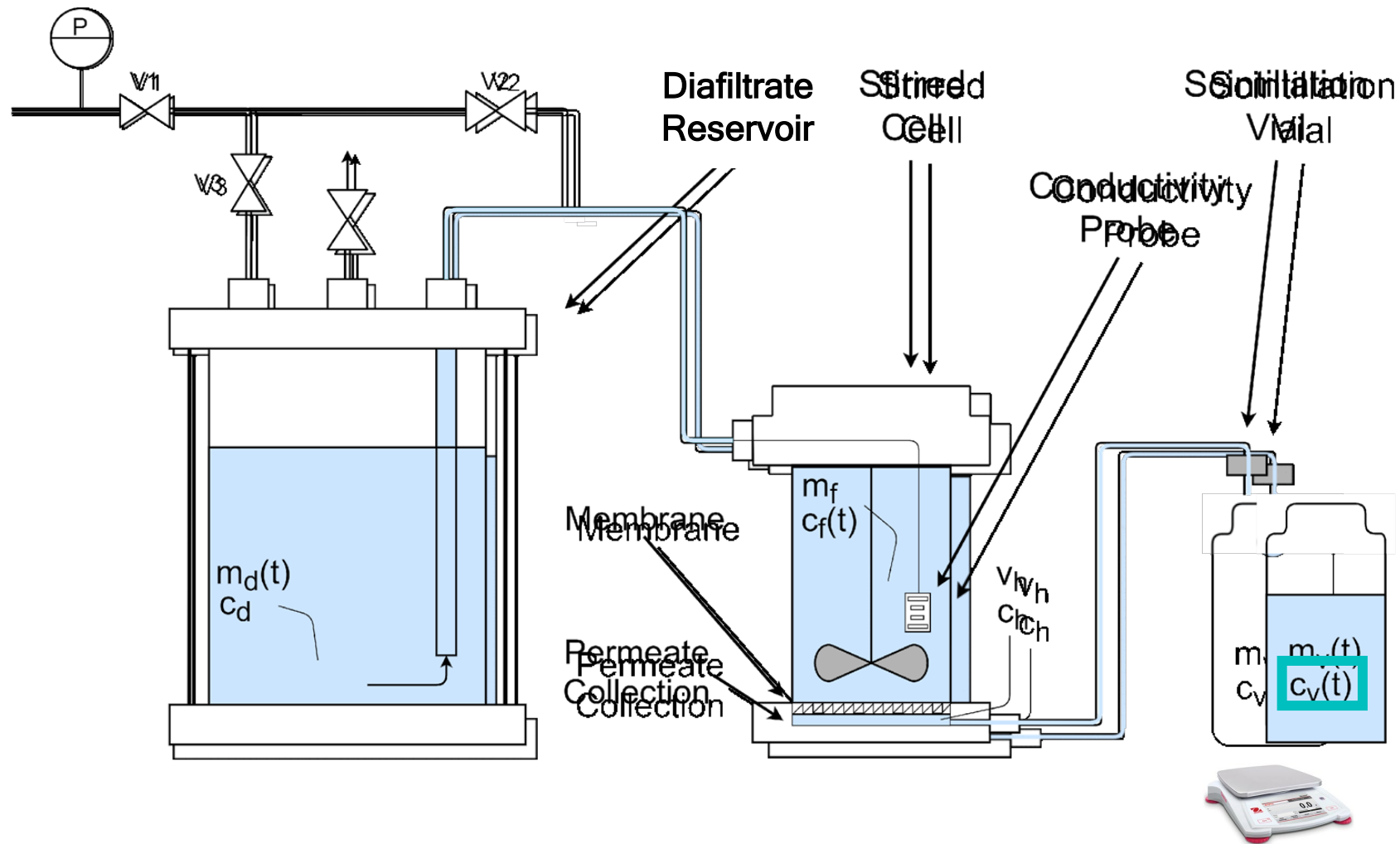
Experimental degrees of freedom: temperature, pressure, feed concentration, diafiltration concentration



Animation: experiment start-up

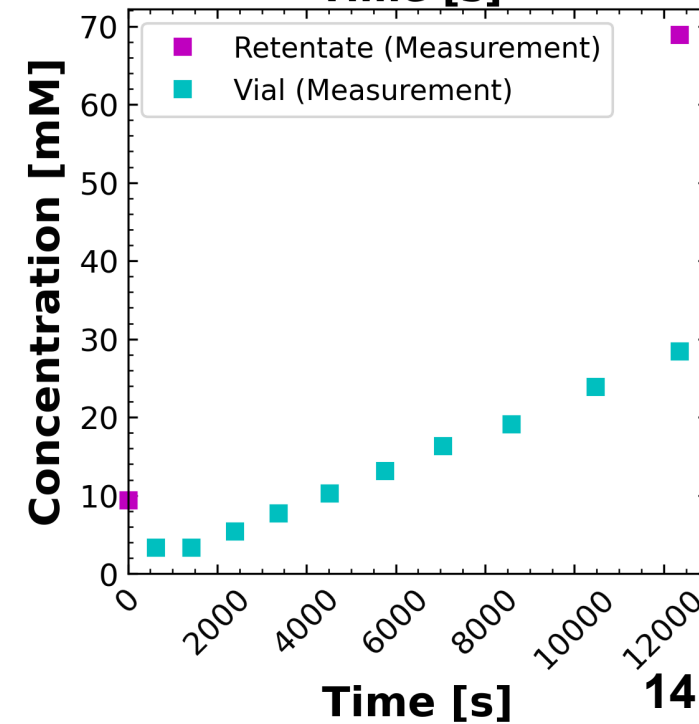
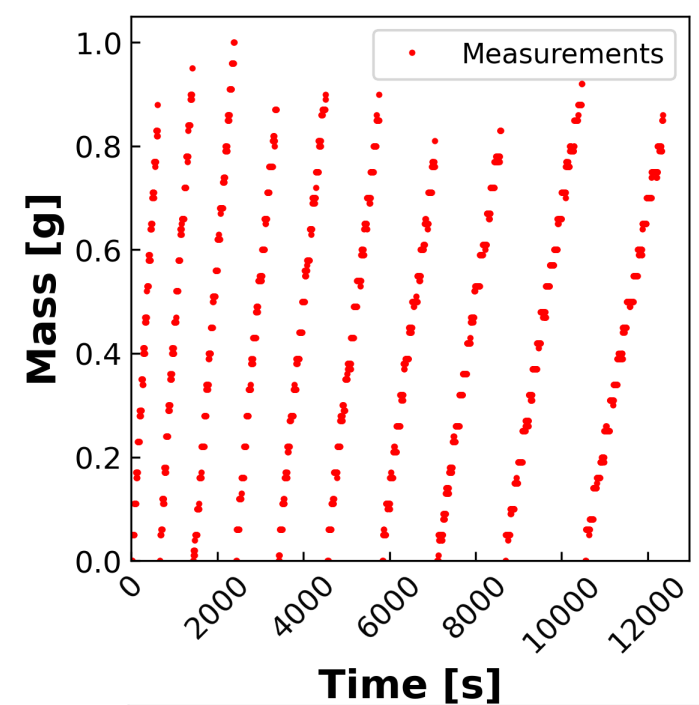
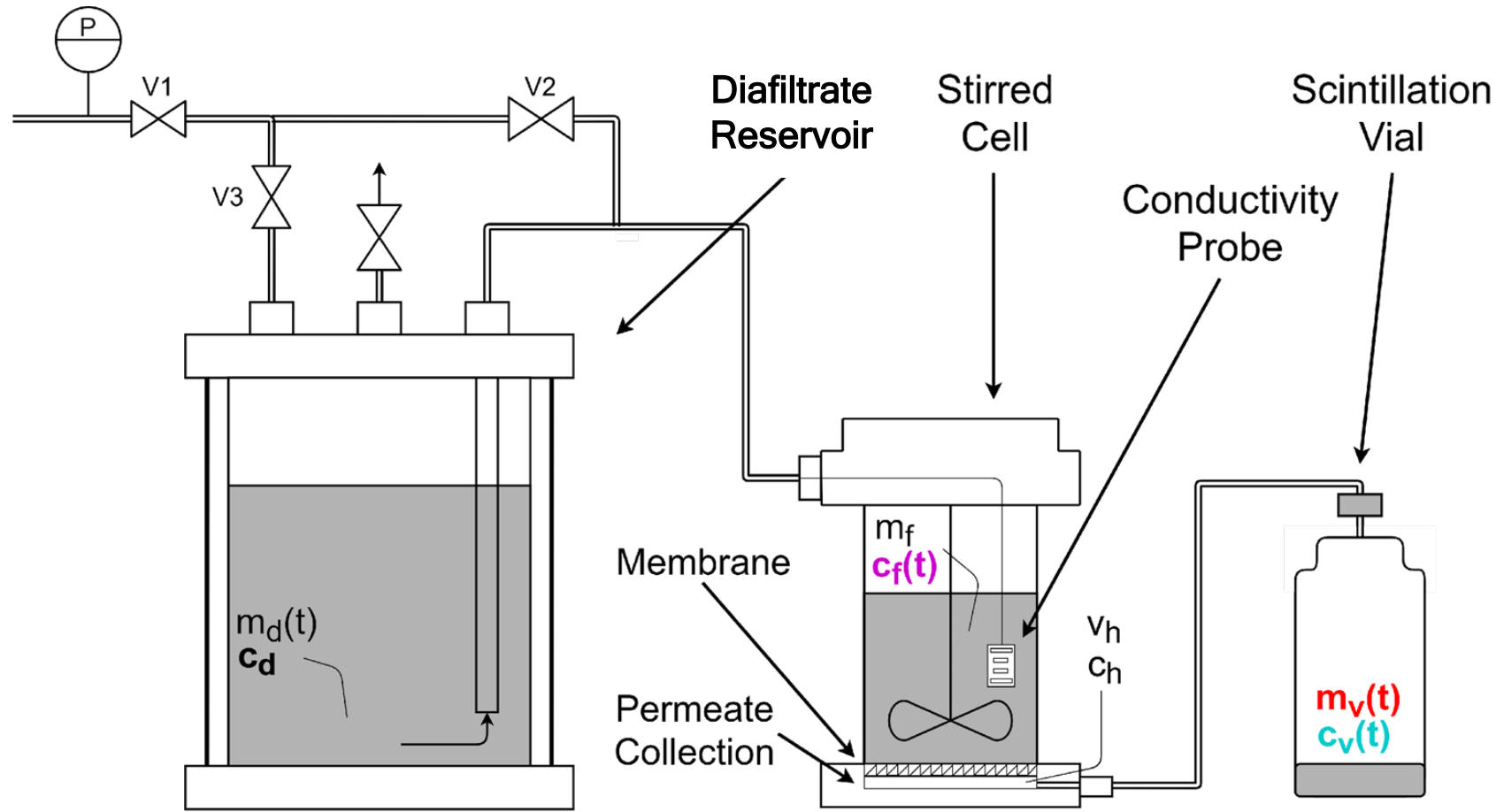


# Dynamic Diafiltration Experiment Sweeps Larger Concentration Space

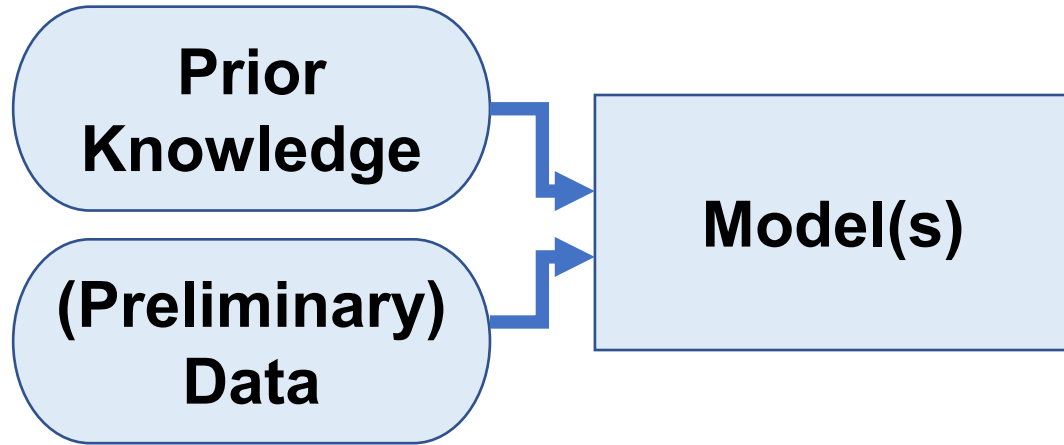


# Dynamic Diafiltration Experiment Sweeps

## Larger Concentration Space 5x Faster Than Filtration Experiments

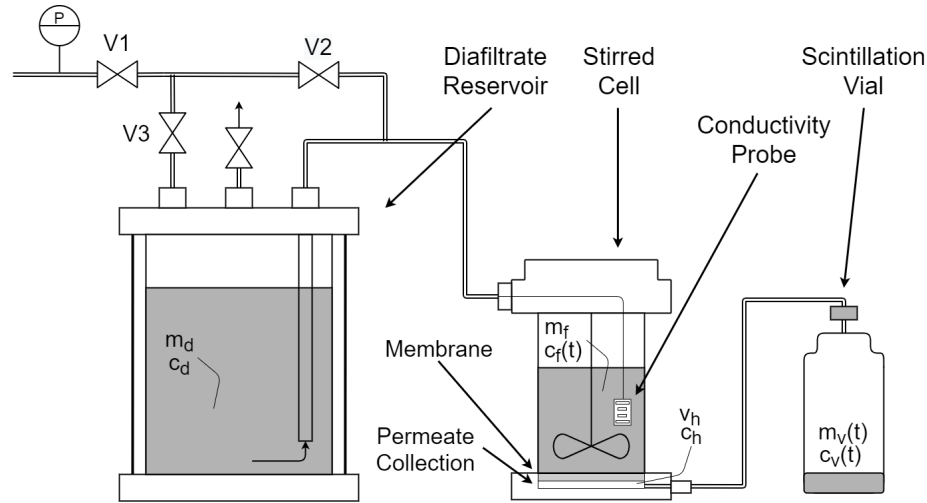


# Science-based Modeling Workflow





# Lumped Parameter Model Infers Membrane Properties



## Fitted Parameters

$L_p$ : hydraulic permeability

$B$ : solute permeability coefficient

$\sigma$ : reflection coefficient

$\Delta P$ : applied pressure drop

$n$ : number of dissolved species

$R$ : gas constant

$T$ : temperature

$A_m$ : area of the membrane

$\rho$ : density of solution

$k$ : Mass transfer coefficient of interest

$b$ : Diameter of the stirred cell

$D$ : Diffusion coefficient of the solute in the solvent

$\nu$ : Kinematic viscosity of the solvent

$\nu^0$ : Average velocity within the system

Water flux across the membrane

$$J_w = L_p (\Delta P - \sigma \Delta \pi)$$

$$\Delta \pi = nRT(c_{in} - c_h)$$

Solute flux across the membrane

$$J_s = B(c_{in} - c_h)$$

Concentration polarization

$$\frac{c_{in} - c_h}{c_f - c_h} = \exp\left(\frac{J_w}{k}\right)$$

$$\frac{kb}{D} = 0.23 \left(\frac{b\nu^0}{\nu}\right)^{0.57} \left(\frac{\nu}{D}\right)^{0.33}$$

Mass balance in dialysate reservoir

$$\frac{dm_d}{dt} = -A_m \rho J_w$$

Constant concentration in dialysate reservoir

$$\frac{dc_d}{dt} = 0$$

Constant mass in feed cell

$$\frac{dm_f}{dt} = 0$$

Solute balance in feed cell

$$\frac{d(c_f m_f)}{dt} = A_m \rho (J_w c_d - J_s)$$

Constant mass in holdup

$$\frac{dm_h}{dt} = 0$$

Solute balance in holdup

$$\frac{d(c_h m_h)}{dt} = A_m \rho J_s + \frac{dm_d}{dt} c_h$$

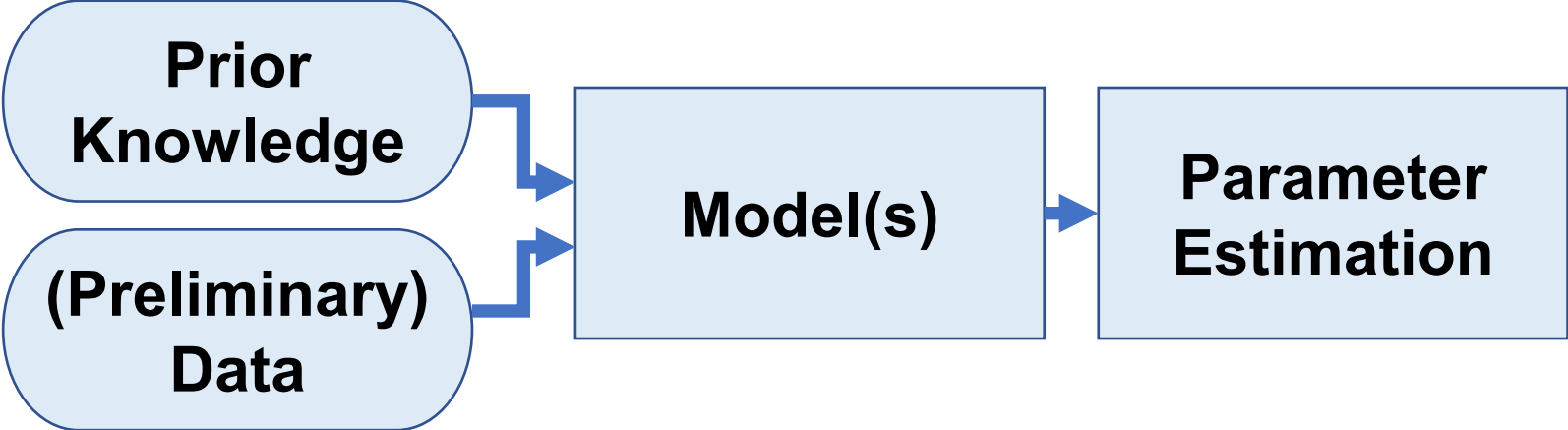
Mass balance in collecting vial

$$\frac{dm_v}{dt} = -\frac{dm_d}{dt} = A_m \rho J_w$$

Solute balance in collecting vial

$$\frac{d(c_v m_v)}{dt} = -\frac{dm_d}{dt} c_h = A_m \rho J_w c_h$$

# Science-based Modeling Workflow



# Nonlinear Regression as a 2-Stage Stochastic Program

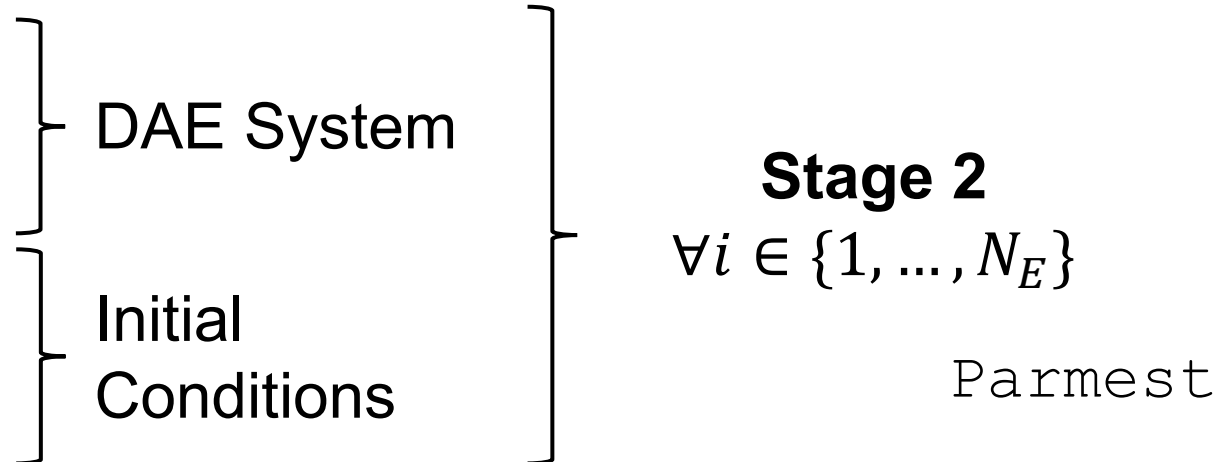
Pyomo: Bynum, Hackebeil, Hart, Laird, Nicholson, Sirola, Watson, Woodruff. Springer, 2021.

Parmest: Klise, Nicholson, Staid, Woodruff. *Computer Aided Chemical Engineering*, 47 (2019): 41-46.

Pyomo.DAE: Nicholson, Sirola, Watson, Zavala, and Biegler. *Mathematical Programming Computation* 10(2) (2018): 187-223.

$$\min \sum_{i=1}^{N_E} \sum_{t \in t_i} (\tilde{y}_i(t) - y_i(t))^T \Sigma_y (\tilde{y}_i(t) - y_i(t))$$

$$\text{s.t.} \quad \begin{aligned} \dot{x}_i(t) &= f(x_i(t), z_i(t), u_i(t), \bar{w}_i, \theta) \\ g(x_i(t), z_i(t), u_i(t), \bar{w}_i, \theta) &= \mathbf{0} \\ y(t) &= h(x_i(t), z_i(t), \theta) \\ f^0(\dot{x}_i(t_0), x_i(t_0), z_i(t_0), u_i(t_0), \bar{w}_i, \theta) &= \mathbf{0} \\ g^0(x_i(t_0), z_i(t_0), u_i(t_0), \bar{w}_i, \theta) &= \mathbf{0} \\ y^0(t_0) &= h(x_i(t_0), z_i(t_0), \theta) \end{aligned}$$



Stage 2 (for experiment  $i$ )

$N_E$  Number of experiments

$t_i$  Measurement times for experiment  $i$

$\Sigma_y$  Covariance matrix for measurement errors

$\theta$  Estimated parameters (stage 1)

$\tilde{y}_i$  &  $y_i$  Measurements & model responses (parameter, variable)

$x_i$  Time-dependent differential state variables

$z_i$  Time-dependent algebraic state variables

$u_i$  Time-varying control (parameter)

$\bar{w}_i$  Time-invariant control (parameter)

# Weighted Least-Squared Parameter Estimation

$$\hat{\theta} = \arg \min_{\theta} \sum_i w_{m_v,i} (m_{v,i} - \hat{m}_{v,i})^2$$

Mass collected

$$+ \sum_j w_{c_v,j} (c_{v,j} - \hat{c}_{v,j})^2$$

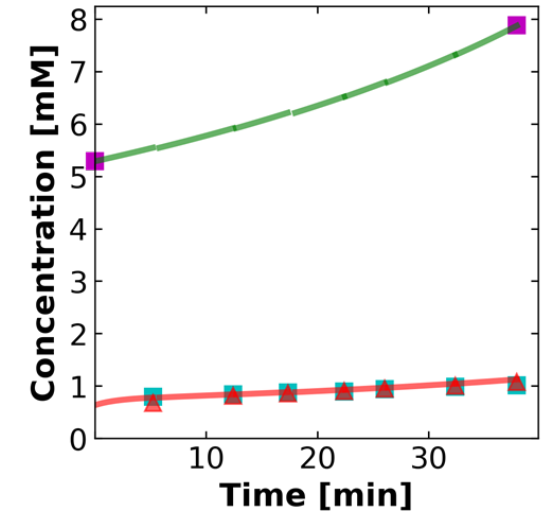
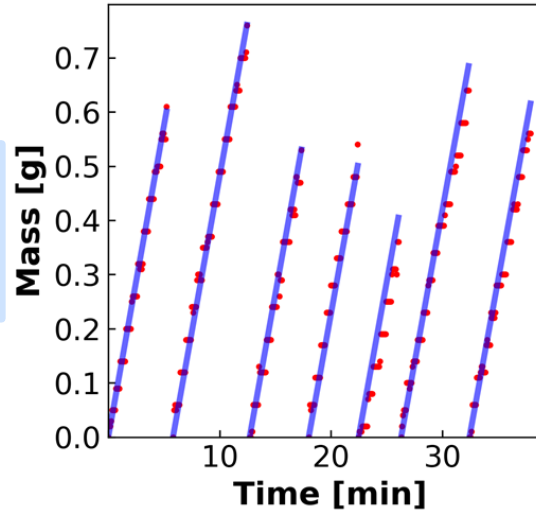
Vial concentration

$$+ \sum_k w_{c_f,k} (c_{f,k} - \hat{c}_{f,k})^2$$

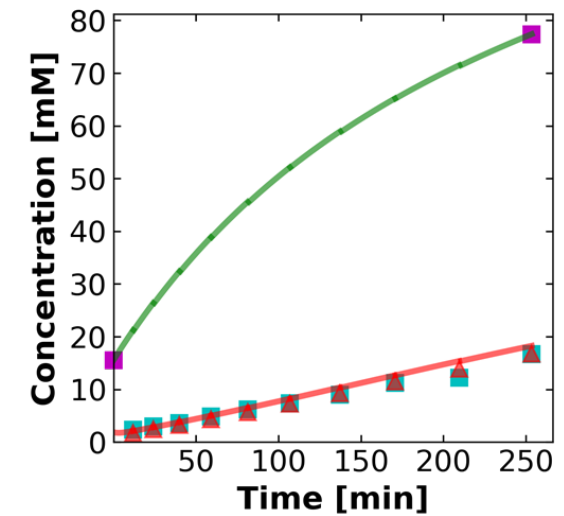
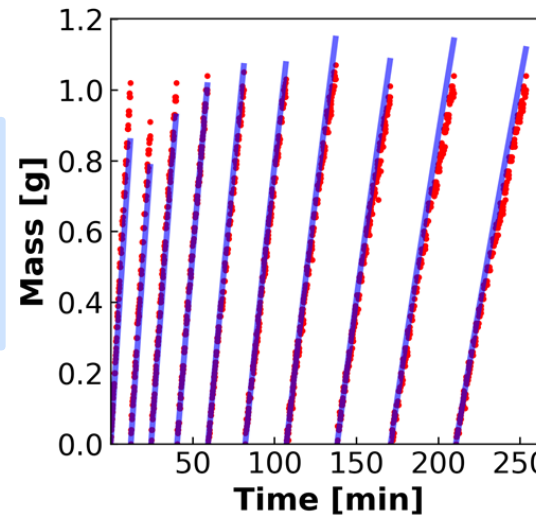
Retentate concentration

	$L_p$	B	$\sigma$
Filtration	4.30 L/(m <sup>2</sup> -hr-bar)	0.67 μm/s	0.83 dimensionless
Diafiltration	3.25 L/(m <sup>2</sup> -hr-bar)	0.31 μm/s	1 dimensionless

**Filtration mode**  
Feed: 5mM KCl



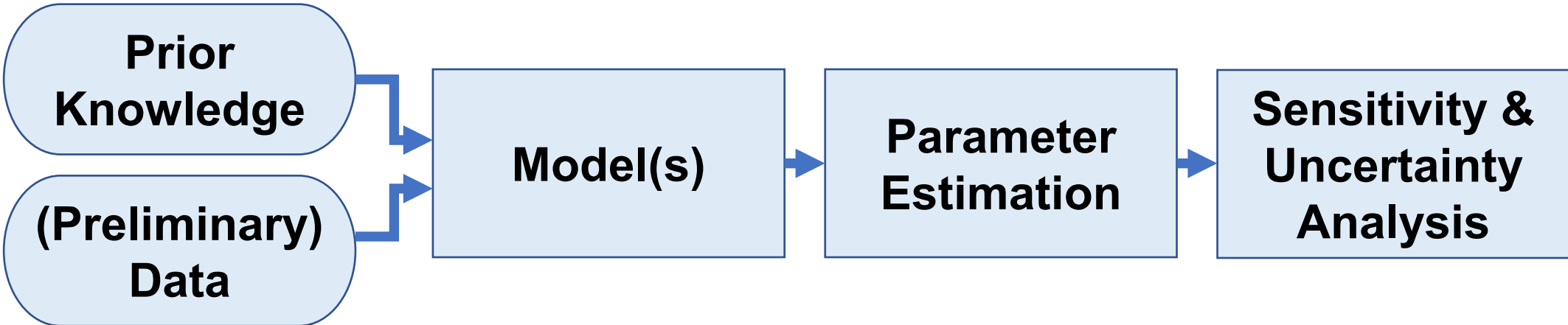
**Diafiltration mode**  
Feed: 5mM KCl  
Dialysate: 80mM KCl



• Measurements  
— Predictions

■ Retentate (Measurement)  
— Retentate (Prediction)  
■ Vial (Measurement)  
▲ Vial (Prediction)  
— Permeate (Prediction)

# Science-based Modeling Workflow



# Parameter Estimation and Uncertainty

Assume a model and error structure:

$$y_i = f(x_i, \theta) + \epsilon_i \quad \epsilon \sim N(0, \sigma_\epsilon^2)$$

What values of model parameters  $\theta$  best fit the data  $x$  and  $y$ ?

$$\hat{\theta} = \operatorname{argmin}_{\theta} \sum_i [y_i - f(x_i, \theta)]^2$$

best fit estimates

How sensitive is the least-squares objective  $\Psi$  to perturbations in  $\theta$ ?

$$H = \begin{bmatrix} \frac{\partial^2 \Psi}{\partial \theta_1^2} & \cdots & \frac{\partial^2 \Psi}{\partial \theta_n \partial \theta_1} \\ \vdots & \ddots & \vdots \\ \frac{\partial^2 \Psi}{\partial \theta_1 \partial \theta_m} & \cdots & \frac{\partial^2 \Psi}{\partial \theta_m^2} \end{bmatrix}$$

Hessian matrix

$$H \approx Q^T Q$$

$$Q(\theta) = \begin{bmatrix} \frac{\partial f(x_1, \theta)}{\partial \theta_1} & \cdots & \frac{\partial f(x_1, \theta)}{\partial \theta_m} \\ \vdots & \ddots & \vdots \\ \frac{\partial f(x_n, \theta)}{\partial \theta_1} & \cdots & \frac{\partial f(x_n, \theta)}{\partial \theta_m} \end{bmatrix}$$

sensitivity matrix

How does measurement uncertainty  $\epsilon$  propagate into uncertainty about the regressed parameters  $\hat{\theta}$ ?

covariance matrix for  $\hat{\theta}$

$$V_{\hat{\theta}} \approx \sigma_\epsilon^2 H^{-1} \approx \sigma_\epsilon^2 (Q^T Q)^{-1}$$

**Key Insight.** If model predictions  $f(x_i, \theta)$  are **insensitive to  $\theta_j$** , then:

- sensitivity  $Q$ , Hessian  $H$ , and covariance  $V_{\hat{\theta}}$  matrices are (numerically) **rank deficient**
- data  $x$  and  $y$  **cannot identify  $\theta_j$**  in model  $f$
- **large uncertainty** in  $\theta_j$  (and corresponding elements of  $V_{\hat{\theta}}$ )

# Finding 1: Diafiltration Experiments Identify the Reflection Coefficient $\sigma$

Water flux across the membrane

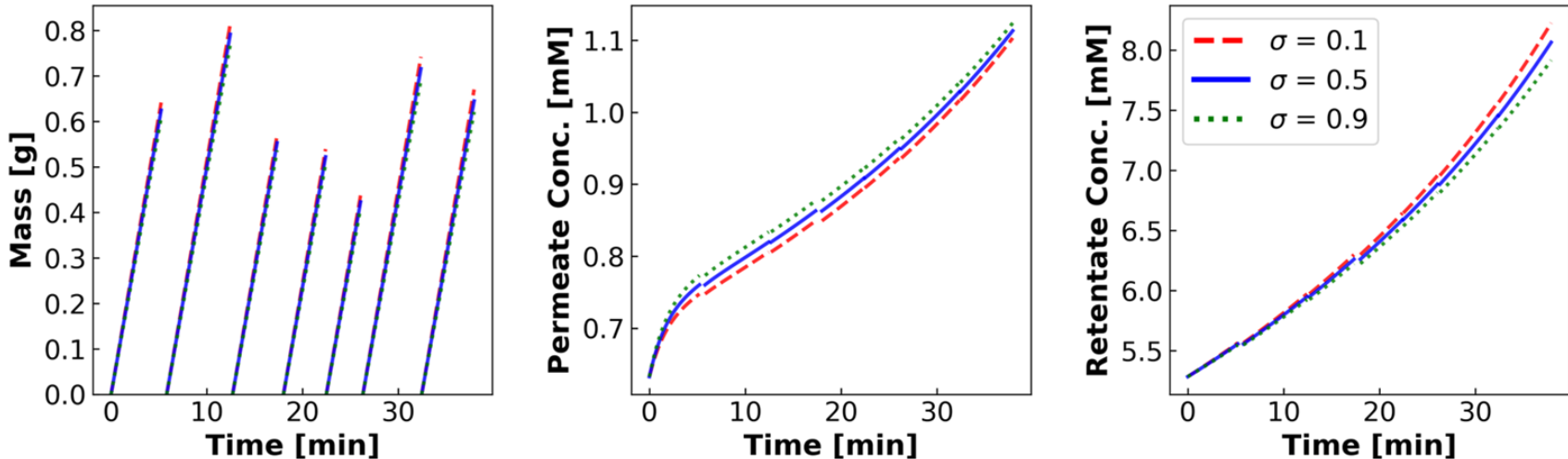
$$J_w = L_p(\Delta P - \sigma \Delta\pi)$$

$$\Delta\pi = nRT(c_{in} - c_h)$$

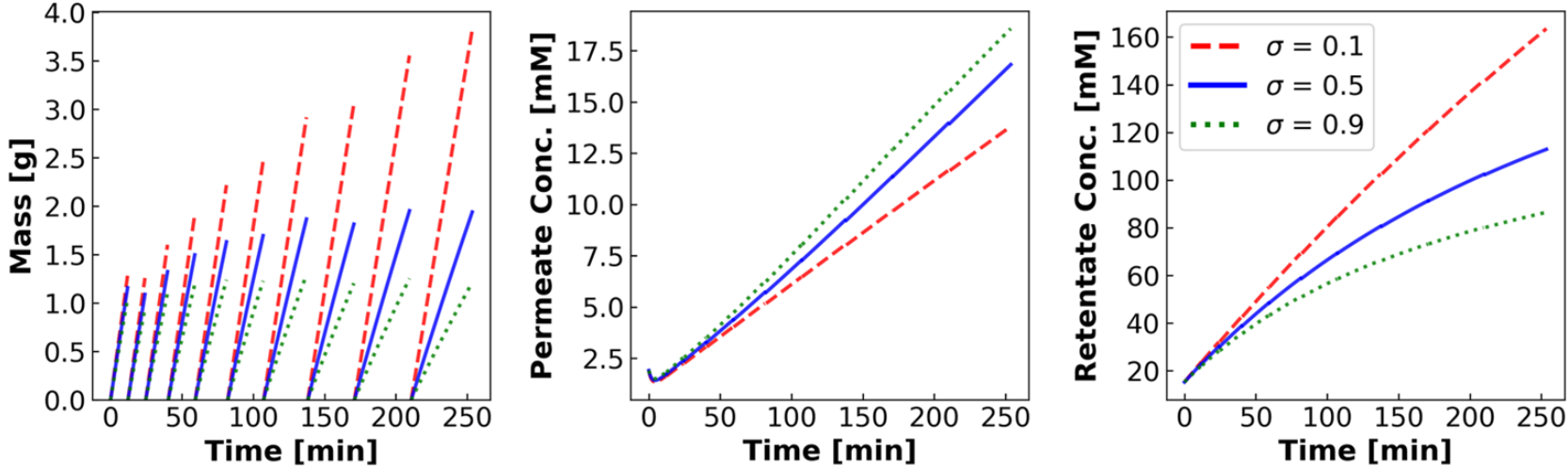
Solute flux across the membrane

$$J_s = B(c_{in} - c_h)$$

Filtration experiment (low concentrations)



Diafiltration experiment (high concentrations)





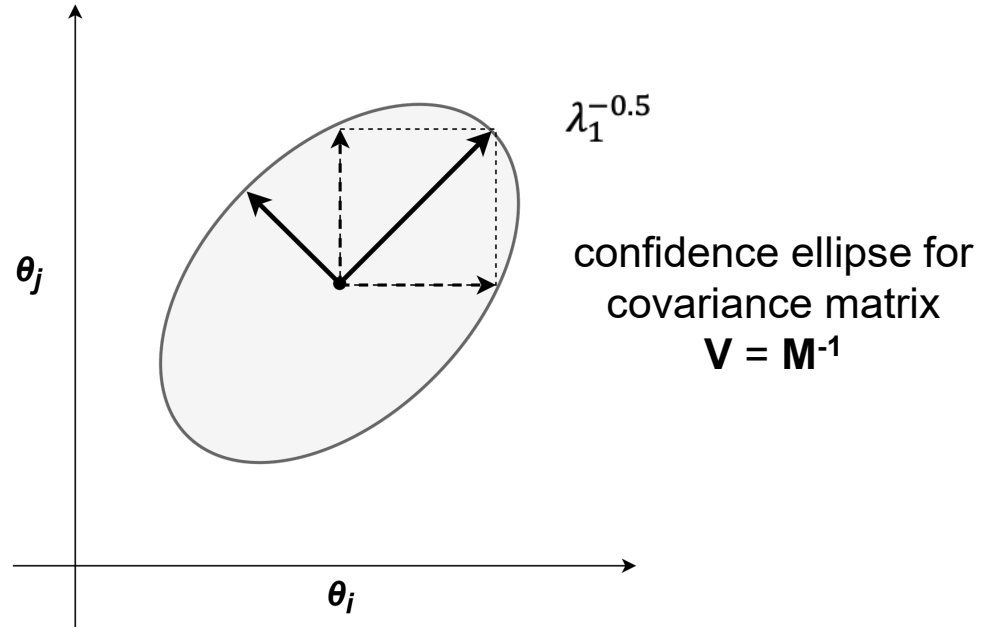
# Finding 1 continued:

## Fisher Information Matrix (FIM)

### Quantifies the Information Content

$$M \approx V_{\hat{\theta}}^{-1} \approx \sigma_{\epsilon}^{-2} H \approx \sigma_{\epsilon}^{-2} Q^T Q$$

$$Q(\theta) = \begin{bmatrix} \frac{\partial f(x_1, \theta)}{\partial \theta_1} & \dots & \frac{\partial f(x_1, \theta)}{\partial \theta_m} \\ \vdots & \ddots & \vdots \\ \frac{\partial f(x_n, \theta)}{\partial \theta_1} & \dots & \frac{\partial f(x_n, \theta)}{\partial \theta_m} \end{bmatrix}$$



- M** Fisher information matrix
- $\hat{\theta}$  Estimated parameters
- $V_{\hat{\theta}}$  Covariance matrix for  $\hat{\theta}$
- H** Hessian matrix for regression opt.
- Q** Model sensitivity matrix
- $\sigma_{\epsilon}^2$  Measurement error variance

FIM Eigen values Eigenvectors

		$L_p$	B	$\sigma$
--	--	-------	---	----------

Filtration	$\times 10^9$	$4.85 \times 10^5$	0.086	0.0033	0.9963		
	4.3352	0.0074	-0.3742	1.63 x 10 <sup>7</sup>	0.002	-1	0.0032
	0.0074	0.0163	-0.0007	4.37 x 10 <sup>9</sup>	-0.9963	-0.0017	0.086
Diafiltration	$\times 10^{10}$	$1.96 \times 10^8$	-0.1405	-0.9468	-0.2894		
	1.7779	0.3004	-1.8364	1.27 x 10 <sup>10</sup>	0.9503	-0.211	0.2289
	0.3004	0.5567	-1.9033	8.18 x 10 <sup>10</sup>	-0.2778	-0.2429	0.9294
	-1.8364	-1.9033	7.1384				

# Multi-objective Parameter Estimation Trade-offs

Mass Residual  
Objective [log g<sup>2</sup>]

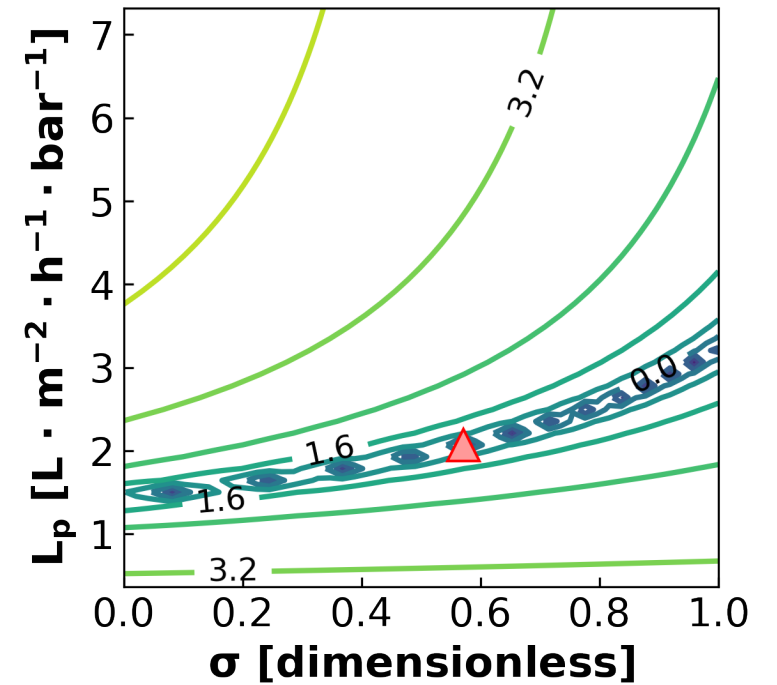
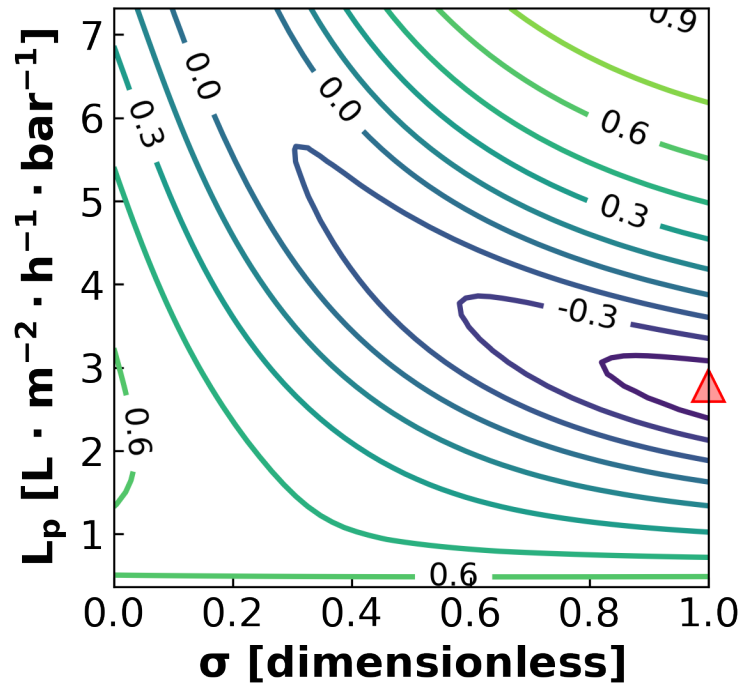
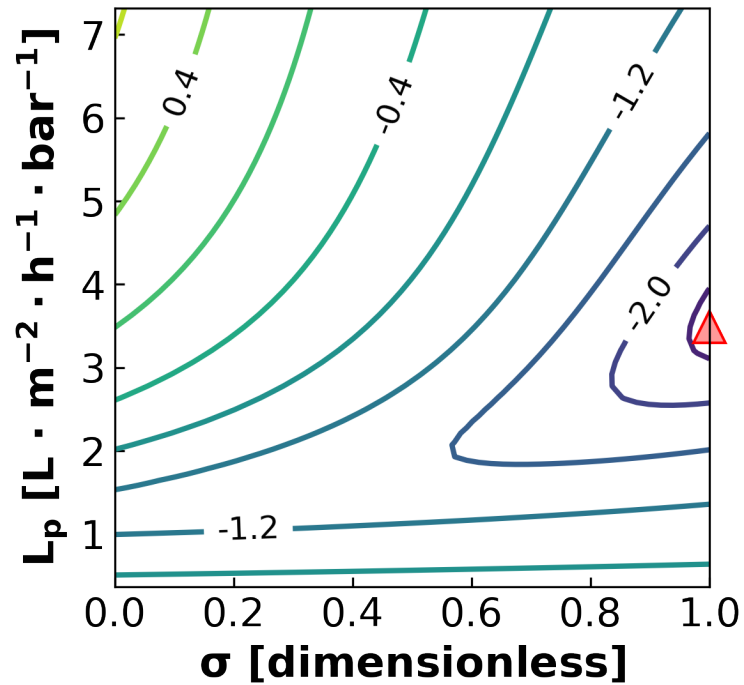
$$\log_{10} \sum_i (m_{v,i} - \hat{m}_{v,i})^2$$

Vial Conc. Residual  
Objective [log mM<sup>2</sup>]

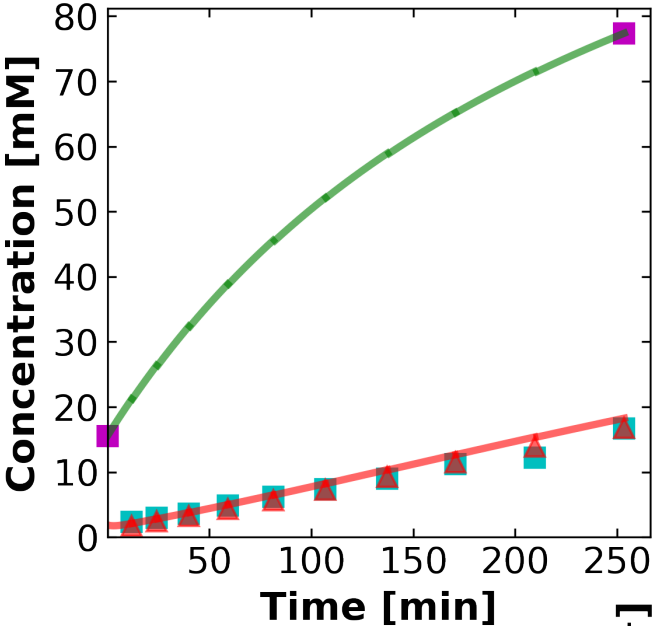
$$\log_{10} \sum_j (c_{v,j} - \hat{c}_{v,j})^2$$

Retentate Conc. Residual  
Objective [log mM<sup>2</sup>]

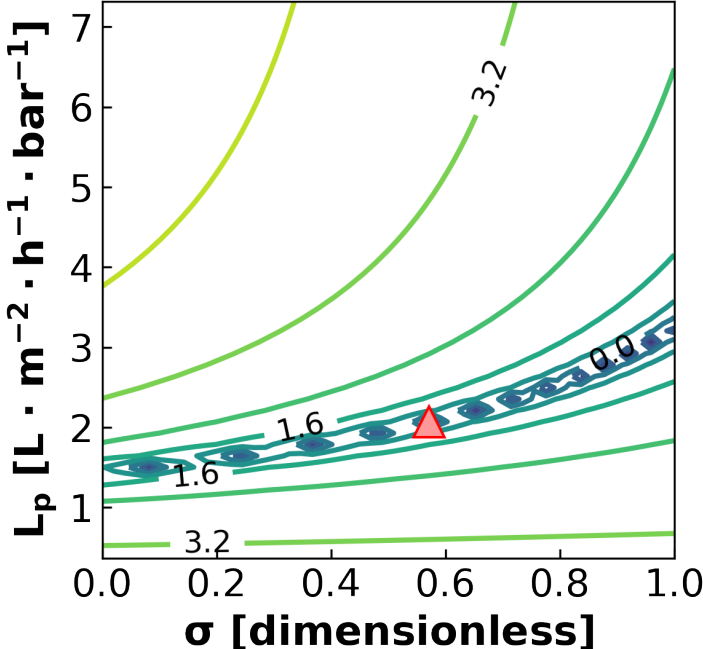
$$\log_{10} \sum_k (c_{f,k} - \hat{c}_{f,k})^2$$



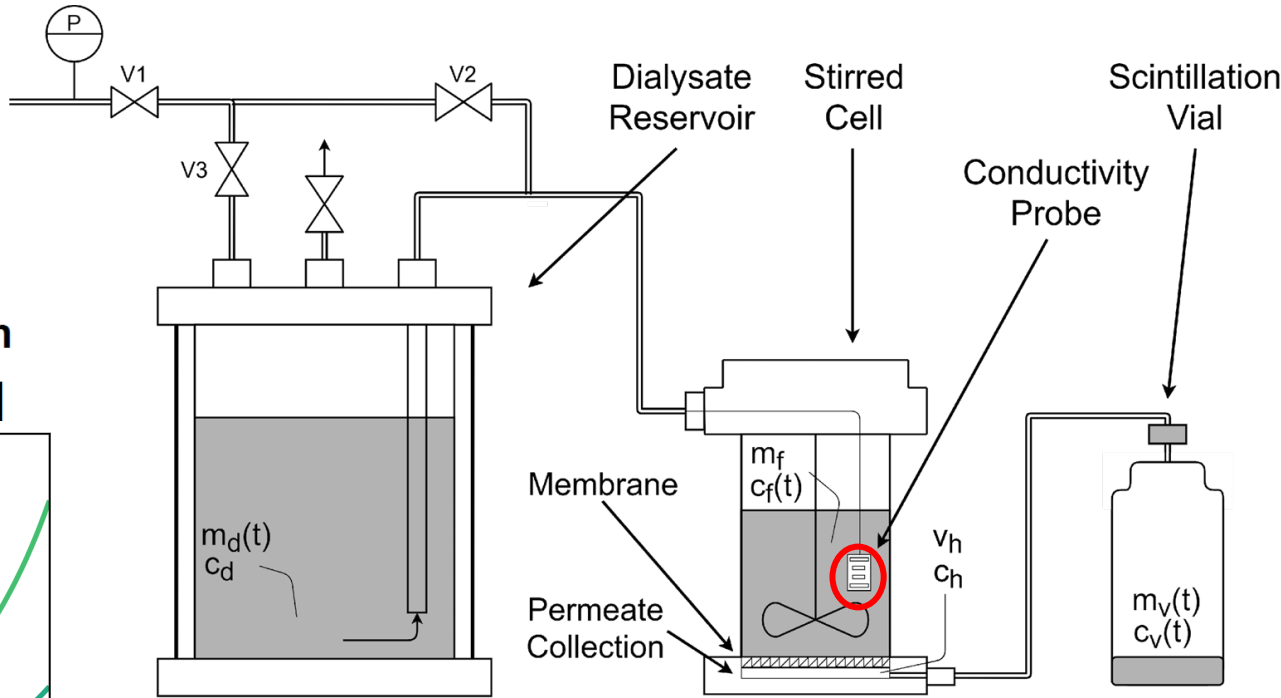
# Finding 2: Add Retentate Concentration Measurements



Log<sub>10</sub> transform  
Retentate Concentration  
Residual Squared [mM<sup>2</sup>]

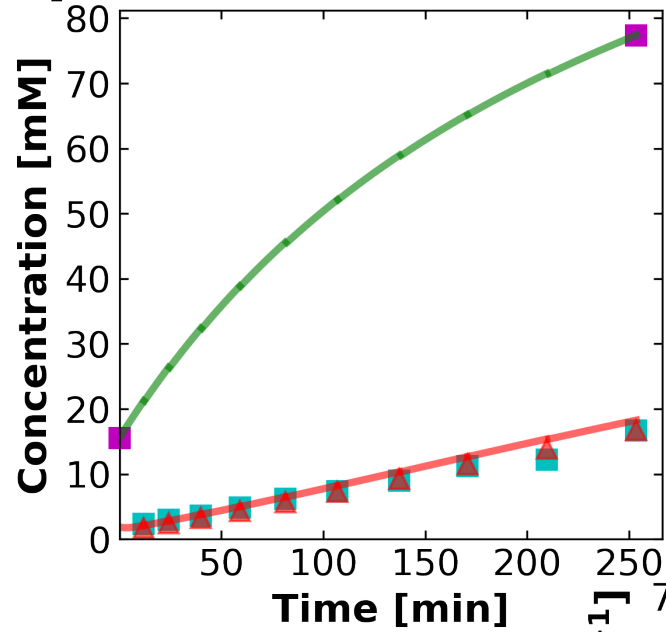


Several sets of parameters could give good predictions



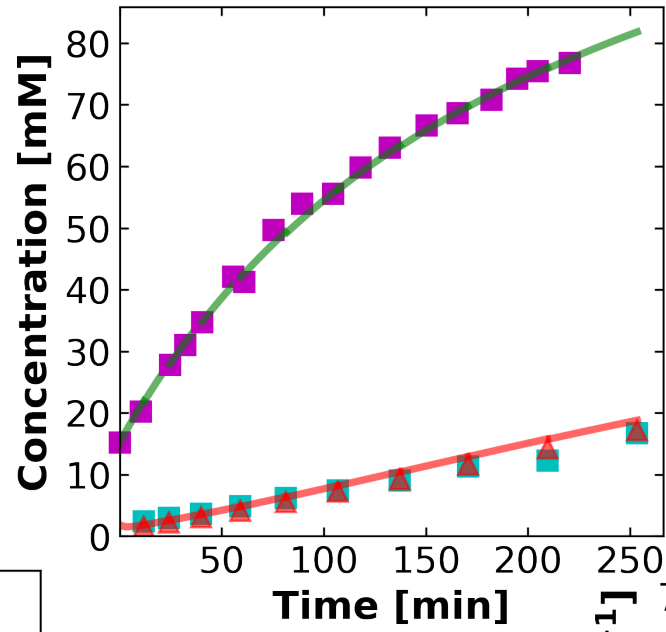
The conductivity probe enables more data collected

# Finding 2: Add Retentate Concentration Measurements improves precision of $\sigma$ estimation

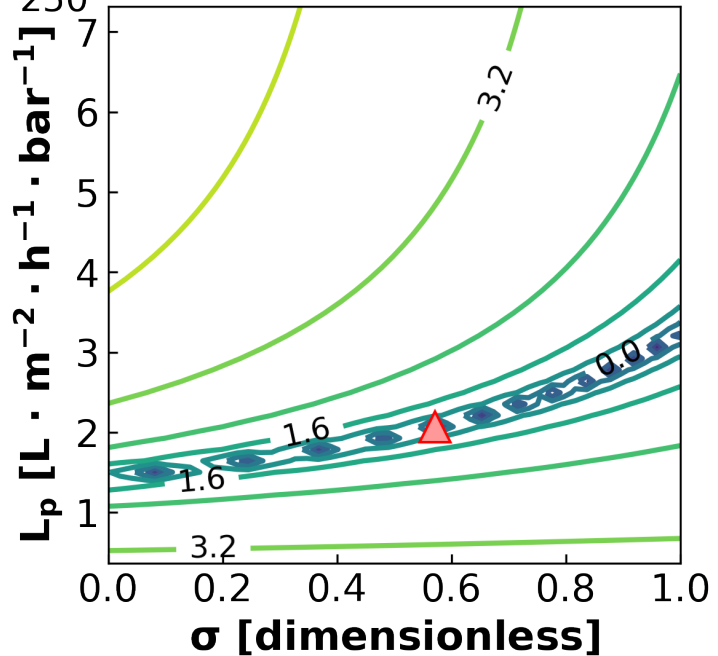


Retentate concentration data added

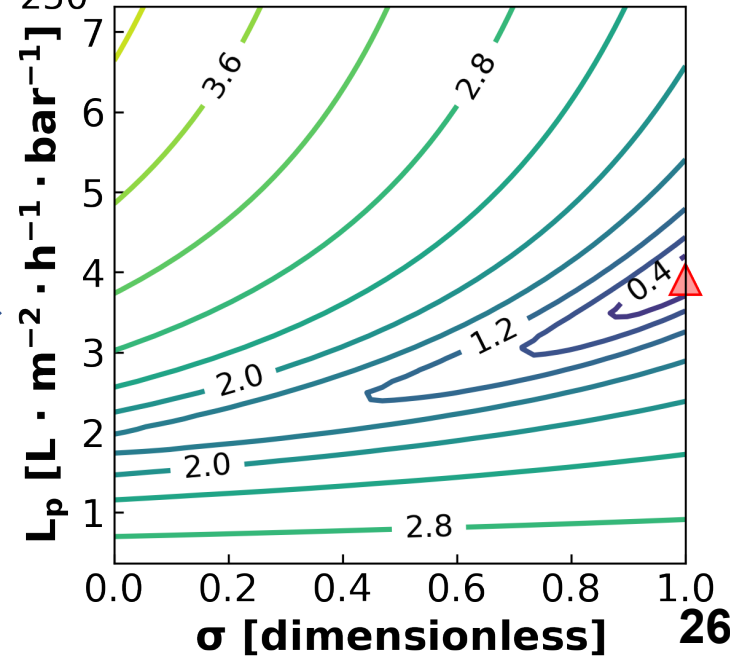
Retentate Conc. Residual Objective [log mM<sup>2</sup>]

$$\log_{10} \sum_k (c_{f,k} - \hat{c}_{f,k})^2$$


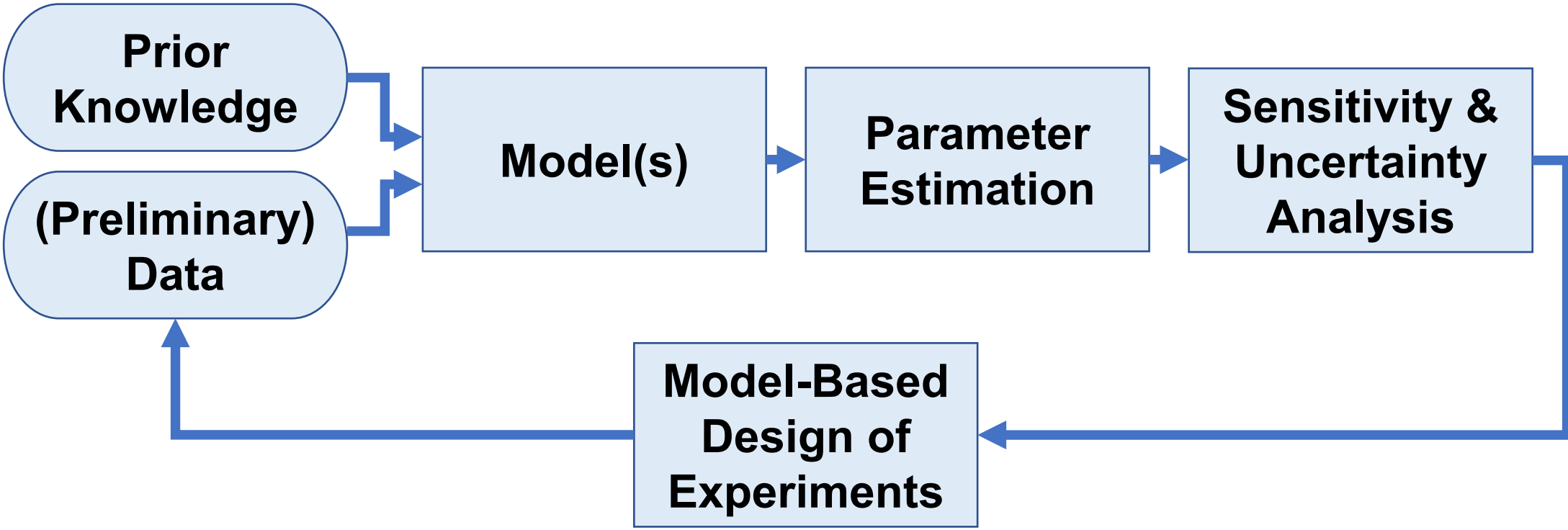
Retentate Conc. Residual Objective [log mM<sup>2</sup>]

$$\log_{10} \sum_k (c_{f,k} - \hat{c}_{f,k})^2$$


Local minima eliminated



# Science-based Modeling Workflow



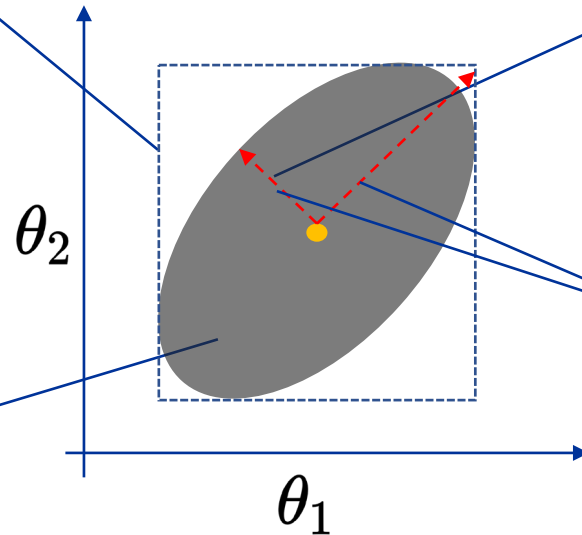
# Alphabetic Design Criteria Measure Information Content

Figure adapted from: Franceschini, G., & Macchietto, S. (2008). *Chem. Eng. Sci.*, 63(19), 4846-4872.

**A-optimality**  
max trace(M)  
enclosing box volume  
poor choice for highly correlated  $\theta$

**D-optimality**  
max det(M)  
ellipsoid volume  
robust to linear transformations

confidence ellipsoid for  
covariance matrix  $\mathbf{V} = \mathbf{M}^{-1}$



**E-optimality**  
max min(eig(M))  
major axis  
recommended if M is ill-conditioned

**ME-optimality**  
min  $\kappa(M) = \max(\text{eig}(M)) / \min(\text{eig}(M))$   
ratio of major to minor axes  
recommended if M is ill-conditioned

## Model Discrimination

Hunter, W.G. and Reiner, A.M., 1965. Designs for discriminating between two rival models. *Technometrics*, 7(3), pp.307-323.

Buzzi-Ferraris, G. and Forzatti, P., 1983. A new sequential experimental design procedure for discriminating among rival models. *Chemical engineering science*, 38(2), pp.225-232.

Ferraris, G.B., Forzatti, P., Emig, G. and Hofmann, H., 1984. Sequential experimental design for model discrimination in the case of multiple responses. *Chemical engineering science*, 39(1), pp.81-85.

## Joint Parameter Precision and Model Discrimination

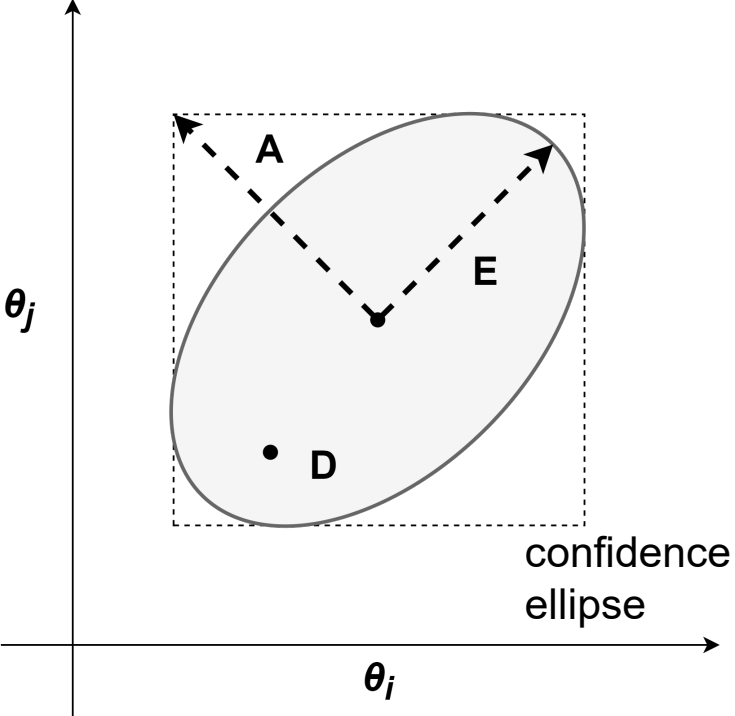
Alberton, A.L., Schwaab, M., Lobão, M.W.N. and Pinto, J.C., 2011. Experimental design for the joint model discrimination and precise parameter estimation through information measures. *Chemical Engineering Science*, 66(9), pp.1940-1952.

Galvanin, F., Cao, E., Al-Rifai, N., Gavriilidis, A. and Dua, V., 2016. A joint model-based experimental design approach for the identification of kinetic models in continuous flow laboratory reactors. *Computers & Chemical Engineering*, 95, pp.202-215.

Galvanin, F., Cao, E., Al-Rifai, N., Dua, V. and Gavriilidis, A., 2015. Optimal design of experiments for the identification of kinetic models of methanol oxidation over silver catalyst. *Chimica Oggi-Chemistry Today*, 33(3), pp.51-56.

Pankajakshan, A., Waldron, C., Quaglio, M., Gavriilidis, A. and Galvanin, F., 2019. A Multi-Objective Optimal Experimental Design Framework for Enhancing the Efficiency of Online Model Identification Platforms. *Engineering*, 5(6), pp.1049-1059.

# Finding 2 Continued: Fisher Information Matrix Quantifies Benefits of New Retentate Concentration Measurements (Conductivity Probe)



- A** dimensions of the enclosing box
- D** area or volume
- E** size of major axis
- ME** ratio of minor axis over major axis

A-optimal Maximize	D-optimal Maximize	E-optimal Maximize	ME-optimal Minimize
Trace	Determinant	Smallest eigenvalue	Condition number

w/o conductivity probe

w/ conductivity probe

9.47 x 10 <sup>10</sup>	2.03E+29	1.96E+08	417.745
1E+11 <b>6%↑</b>	2.69E+29 <b>32%↑</b>	2.17E+08 <b>11%↑</b>	393 <b>6%↓</b>

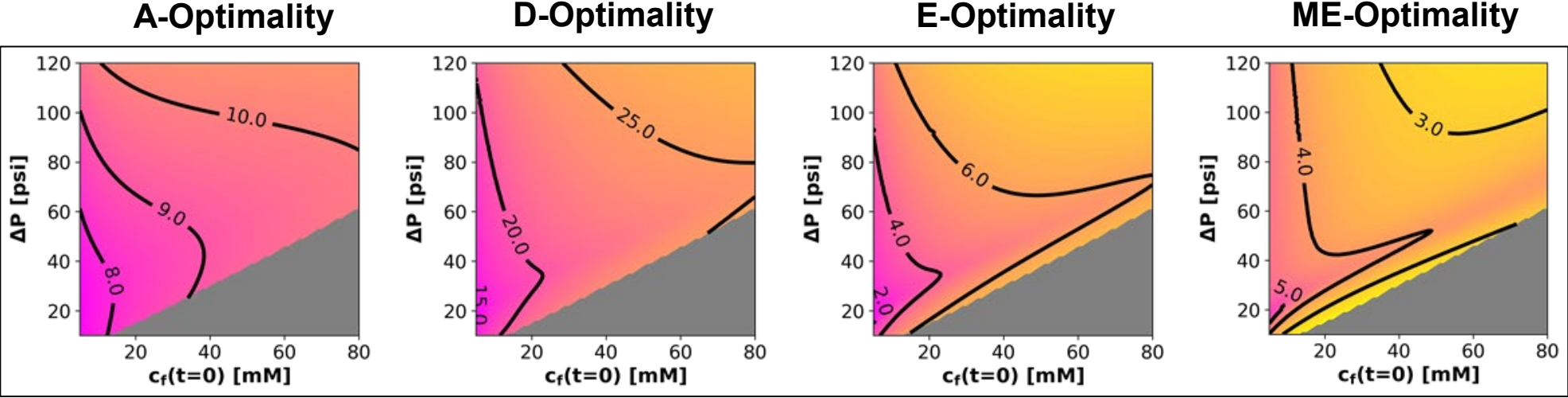
improved parameter precision

less correlation

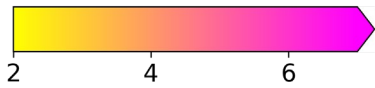
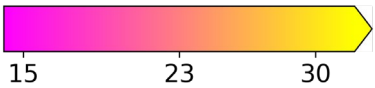
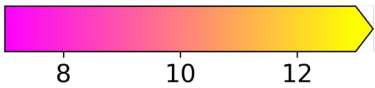
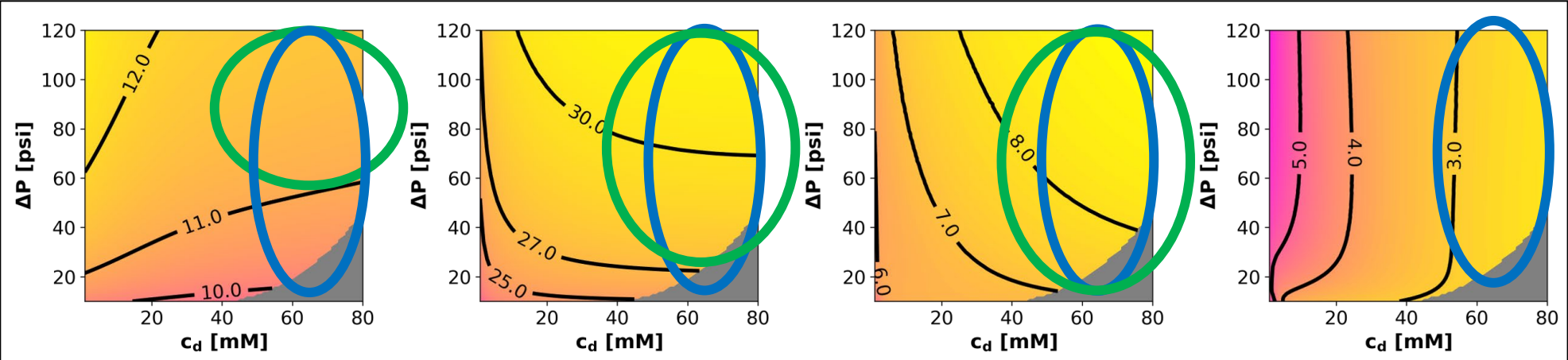


# Finding 3: Recommend Diafiltration Experiments with $c_d \geq 50$ mM and $\Delta P \geq 45$ psi

Filtration mode



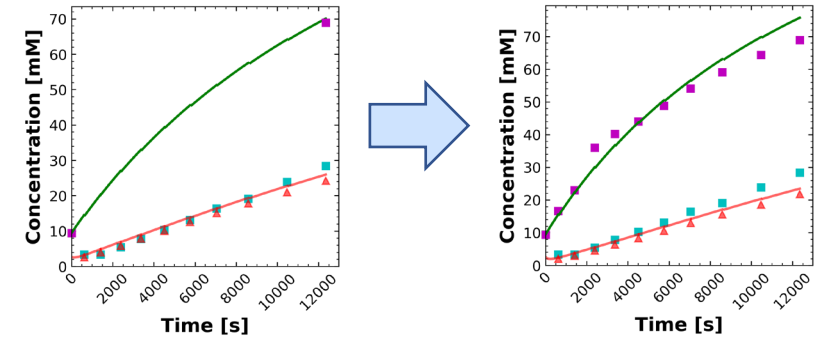
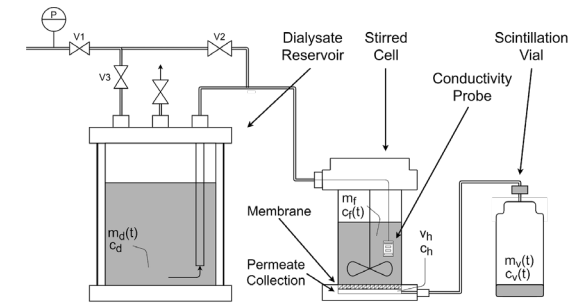
Diafiltration mode



# Membrane Tutorial Conclusions

FIM analysis quantifies benefits of new sensors *a priori*

FIM eigendecomposition gives insights into model identifiability



Ouimet, J. A., Liu, X., Brown, D. J., Eugene, E. A., Poppo, T., Muetzel, Z. W., Dowling, A. W., & Phillip, W. A. (2022). **DATA: Diafiltration Apparatus for high-Throughput Analysis.** *Journal of Membrane Science*, 641, 119743.

Liu, X., Wang, J., Ouimet, J. A., Phillip, W. A., & Dowling, A. W. (2022). **Accelerating Membrane Characterization with Model-Based Design of Experiments.** *Computer Aided Chemical Engineering, PSE2021+*.



The Patrick and Jana Eiler's Graduate Student Fellowship (E. Eugene)

The Vincent P. Slatt Fellowship for Undergraduate Research in Energy Systems and Processes (N. Wamle)



Award CBET-1941596

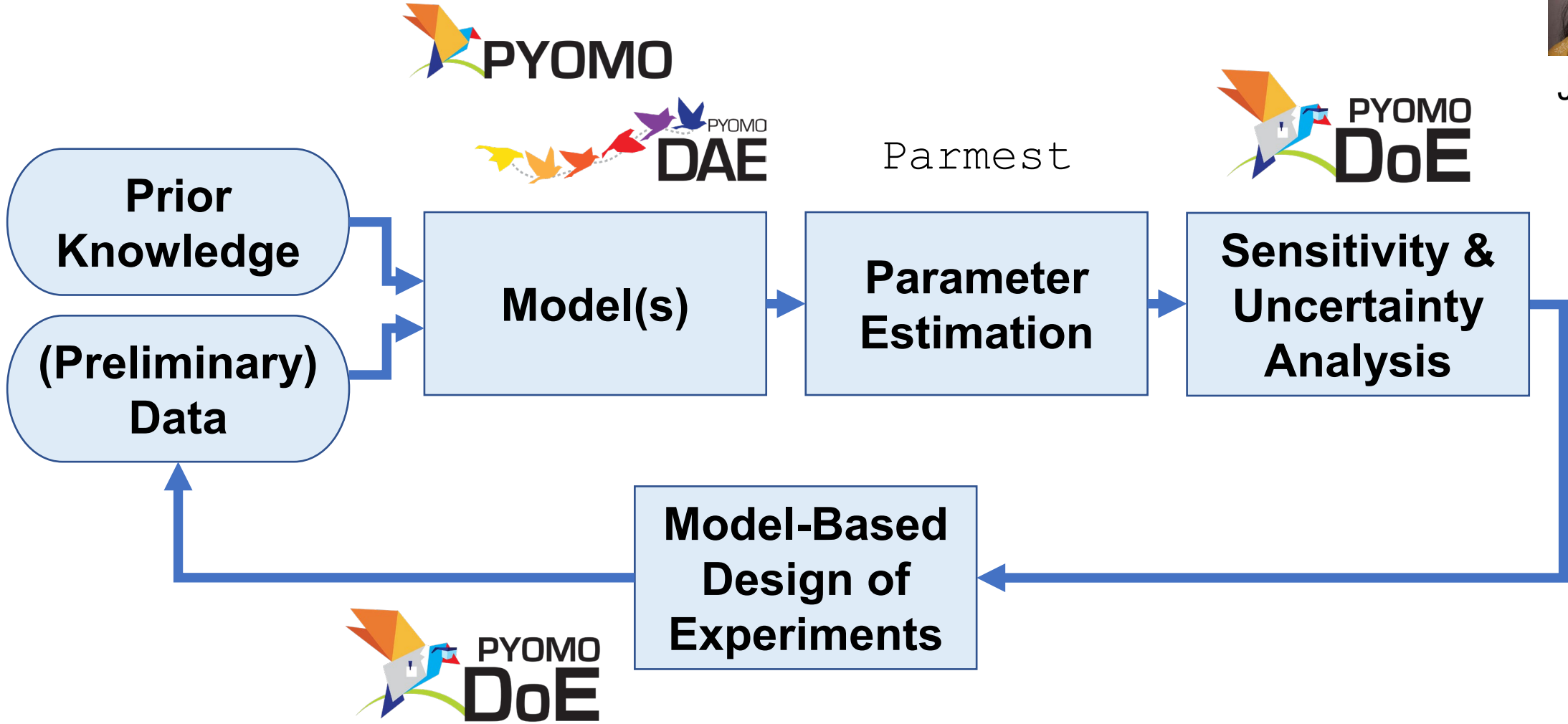
Award CMMI-1932206-S001



# Science-based Modeling Workflow



Jialu Wang



# SBDoE Optimization Formulation



Jialu Wang

$$\begin{aligned}
 & \max_{\varphi} \quad \Psi[ M(\hat{\theta}, \varphi) ] \\
 \text{s. t.} \quad & \left. \begin{aligned}
 & \dot{x}(t) = f(x(t), z(t), u(t), \bar{w}, \hat{\theta}) \\
 & g(x(t), z(t), u(t), \bar{w}, \hat{\theta}) = \mathbf{0} \\
 & y(t) = h(x(t), z(t), \hat{\theta})
 \end{aligned} \right\} \text{DAE System} \\
 & \left. \begin{aligned}
 & f^0(\dot{x}(t_0), x(t_0), z(t_0), u(t_0), \bar{w}, \hat{\theta}) = \mathbf{0} \\
 & g^0(x(t_0), z(t_0), u(t_0), \bar{w}, \hat{\theta}) = \mathbf{0} \\
 & y^0(t_0) = h(x(t_0), z(t_0), \hat{\theta})
 \end{aligned} \right\} \text{Initial Conditions}
 \end{aligned}
 \quad \left. \vphantom{\begin{aligned} \max_{\varphi} \end{aligned}} \right\} m(x(t), y(t), z(t), u(t), \bar{w}, \hat{\theta}) = \mathbf{0}$$

- $y$  Measurements (model responses)
- $\hat{\theta}$  Estimated parameters
- $x$  Time-dependent differential state variables
- $z$  Time-dependent algebraic state variables
- $u$  Time-varying control variables
- $\bar{w}$  Time-invariant control variable

**Fisher information matrix (FIM):**

$$M \approx V_{\hat{\theta}}^{-1} \approx \sigma_{\epsilon}^{-2} H \approx \sigma_{\epsilon}^{-2} Q^T Q$$

**MBDoE Decisions:**

$$\varphi = (u(t), x(t_0), z(t_0), \bar{w}, t)$$

# Pyomo.DoE Formulation: MBDoE as 2-Stage Stochastic Program

max  $\log \det(\mathbf{M}(\hat{\boldsymbol{\theta}}, \boldsymbol{\varphi})) = 2 \sum_{i=1}^{N_p} \log L_{ii}$  D-optimality

s.t.  $\mathbf{M} = \sum_r \sum_{r'} \tilde{\sigma}_{r,r'} \mathbf{Q}_r^T \mathbf{Q}_{r'}$  Stage 1

$\mathbf{M} = \mathbf{L}\mathbf{L}^T, L_{ii} \geq \epsilon$  Cholesky factorization

$q_{r,p}(t) = \frac{y_{r,p}^+(t) - y_{r,p}^-(t)}{2\epsilon_p}$  Central finite difference

$\mathbf{m}(x_p^+(t), y_p^+(t), z_p^+(t), \mathbf{u}(t), \bar{\mathbf{w}}, \boldsymbol{\theta}_p^+) = \mathbf{0}$  Two model evaluations

$\mathbf{m}(x_p^-(t), y_p^-(t), z_p^-(t), \mathbf{u}(t), \bar{\mathbf{w}}, \boldsymbol{\theta}_p^-) = \mathbf{0}$

$\boldsymbol{\theta}_p^+ = \hat{\boldsymbol{\theta}} + \mathbf{e}_p \epsilon_p$  Up and down perturbations

$\boldsymbol{\theta}_p^- = \hat{\boldsymbol{\theta}} - \mathbf{e}_p \epsilon_p$

**Stage 2**

$\forall p \in \{1, \dots, N_p\}$

## Model Sensitivity

$$\mathbf{Q}_r = \begin{bmatrix} \frac{\partial y_r(t_1)}{\partial \theta_1} & \dots & \frac{\partial y_r(t_1)}{\partial \theta_{N_p}} \\ \vdots & \ddots & \vdots \\ \frac{\partial y_r(t_n)}{\partial \theta_1} & \dots & \frac{\partial y_r(t_n)}{\partial \theta_{N_p}} \end{bmatrix} = [\mathbf{q}_{r,1} \quad \dots \quad \mathbf{q}_{r,N_p}]$$

$$\mathbf{q}_{r,p} = \left[ \frac{\partial y_r(t_1)}{\partial \theta_p} \quad \dots \quad \frac{\partial y_r(t_n)}{\partial \theta_p} \right]^T$$

- $\mathbf{y}$  Measurements (model responses)
- $\mathbf{Q}_r$  Dynamic sensitivity for response  $r$
- $\mathbf{m}(\cdot)$  DAE model
- $\hat{\boldsymbol{\theta}}$  Estimate for parameters
- $\mathbf{M}$  Fisher information matrix
- $\mathbf{L}$  Lower triangular Cholesky factorization
- $\epsilon_p$  Small perturbation for parameter  $p$
- $\mathbf{e}_p$  Unit vector with "1" in position  $p$

# New: Pyomo.DoE Extends ParmEst Interface

```
create_model
```

Create Pyomo model for DAE  
Compatible with parmest

```
DesignVariables
```

Specify the MBDoe degrees  
of freedom and their bounds

```
MeasurementVariables
```

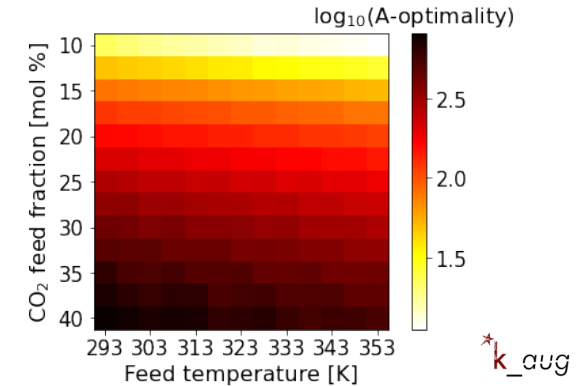
Specify the MBDoe  
measurement variables and  
observation error covariance  
matrix

```
DesignOfExperiment
```



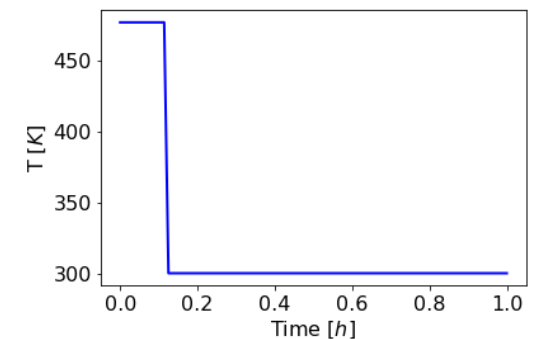
```
compute_FIM
```

Fast exploratory analysis



```
stochastic_program
```

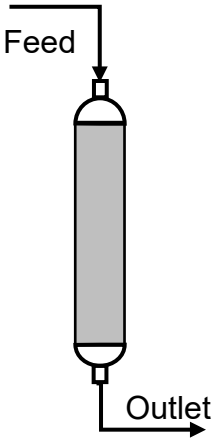
Dynamic optimization



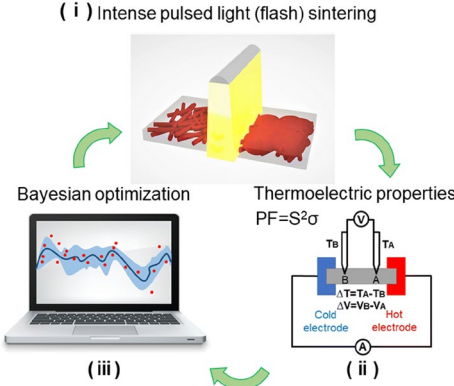


# SBDoE Facilitates Collaborations

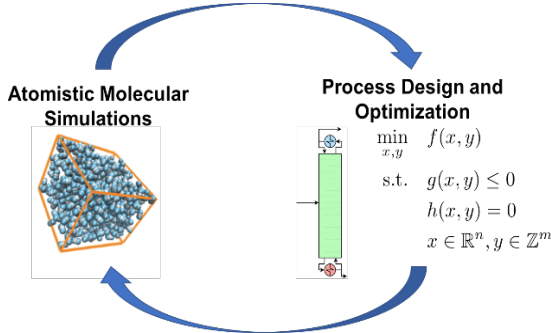
## CO<sub>2</sub> Capture



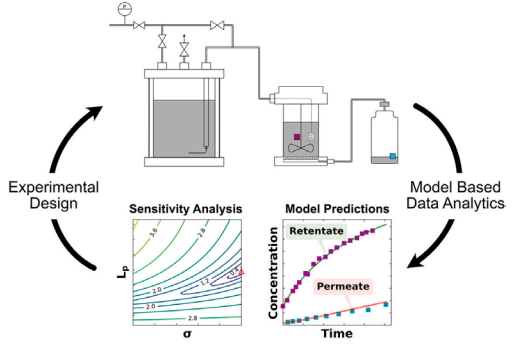
## Additive Manufacturing of Thermoelectric Devices



## Molecular Design (Ionic Liquids)



## Rapid/Automated Membrane Characterization



Jialu Wang



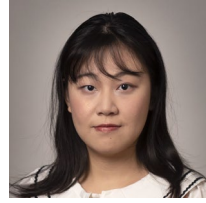
Ke Wang



Bridgette Befort



Xinhong Liu



Wang, J. and Dowling, A.W. (2022), *AIChE J.* e17813.

Wang K., Zhang M., Wang, J., Shang, W., Zhang, Y., Luo, T., Dowling, A.W. (2023), *Digital Chemical Engineering*

Befort, B.J., Garciadiego, A., Wang, J., Wang, K., Maginn, E.J., Dowling, A.W. (2023), *Fluid Phase Equilibria.*

Ouimet, J.A, Xinhong, L., Brown, D.J., Eugene, E.A., Pops, T., Muetzel, Z.W., Dowling, A.W., Phillip, W.A., (2022). *J. Membrane Science.*

# Getting Started with Pyomo.DoE

Documentation: [https://pyomo.readthedocs.io/en/stable/contributed\\_packages/dae/dae.html](https://pyomo.readthedocs.io/en/stable/contributed_packages/dae/dae.html)

Tutorial: [https://colab.research.google.com/github/Pyomo/pyomo/blob/main/pyomo/contrib/dae/examples/fim\\_doe\\_tutorial.ipynb](https://colab.research.google.com/github/Pyomo/pyomo/blob/main/pyomo/contrib/dae/examples/fim_doe_tutorial.ipynb)

The image shows a composite view of Pyomo.DoE resources. On the left is a navigation sidebar for the Pyomo.DoE documentation, listing various solvers and interfaces. The main content area displays the 'Pyomo.DoE' page, which includes a methodology overview and a list of contributors. On the right is a Google Colab notebook titled 'fim\_doe\_tutorial.ipynb', which serves as a tutorial for the 'Reaction Kinetics Example'. The notebook content includes a list of authors (Jialu Wang, Alex Dowling, and Hailey Lynch), their affiliations, and a detailed description of the notebook's purpose: to demonstrate Pyomo.DoE features using a reaction kinetics example. It outlines a three-step process: 1) Importing the necessary modules, 2) Implementing the mathematical model, and 3) Defining the model inputs.

Community Detection for Pyomo models

- Pyomo.DoE
  - Methodology Overview
  - Pyomo.DoE Required Inputs
  - Pyomo.DoE Solver Interface
  - Pyomo.DoE Usage Example
- GDPopt logic-based solver
- Infeasible Irreducible System (IIS) Tool
- Incidence Analysis
- MindtPy Solver
- MPC
- Multistart Solver
- Nonlinear Preprocessing Transformations
- Parameter Estimation with parmest
- PyNumero
- PyROS Solver
- Sensitivity Toolbox
- Trust Region Framework Method Solver
- MC++ Interface
- z3 SMT Sat Solver Interface

Read the Docs v: stable

## Pyomo.DoE

Pyomo.DoE (Pyomo Design of experiments using science-based methodology)

Pyomo.DoE was developed by Jialu Wang, Alex Dowling, and Hailey Lynch at the University of Notre Dame as part of the Carbon Capture Simulation for Industry Impact (CCSI) program through the U.S. Department of Energy.

If you use Pyomo.DoE, please cite the following paper:

[Wang and Dowling, 2022] Wang, Jialu, and Alex Dowling. "Pyomo.DoE: A Python Package for Model-based Design of Experiments." *AIChE J.* <https://doi.org/10.1002/aic.17811>

## Methodology Overview

Model-based Design of Experiments (MDO) is a methodology for designing experiments by directly using simulation. It is one key component in the model-based design of experiments (MDO) methodology.

Prior knowledge, preliminary data → Model

### fim\_doe\_tutorial.ipynb

File Edit View Insert Runtime Tools Help

+ Code + Text Copy to Drive

## Pyomo.DoE Tutorial: Reaction Kinetics Example

Jialu Wang ([jwang44@nd.edu](mailto:jwang44@nd.edu)), Alex Dowling ([adowling@nd.edu](mailto:adowling@nd.edu)), and Hailey Lynch ([hlynch@nd.edu](mailto:hlynch@nd.edu))  
University of Notre Dame

This notebook demonstrates the main features of Pyomo.DoE (model-based design of experiments) using a reaction kinetics example. See [Wang and Dowling \(2022\), AIChE J.](#), for more information.

The user will be able to learn concepts involved with model-based design of experiments (MDO) and practice using Pyomo.DoE from methodology in the notebook. Results will be interpreted throughout the notebook to connect the material with the Pyomo implementation.

The general process that will follow throughout this notebook:

- Import Modules
  - Step 0: Import Pyomo and Pyomo.DoE Module
- Problem Statement
  - Step 1: Import Reaction Kinetics Example Mathematical Model
- Implementation in Pyomo
  - Step 2: Implement Mathematical Model in Pyomo
  - Step 3: Define Inputs for the Model

Methodology



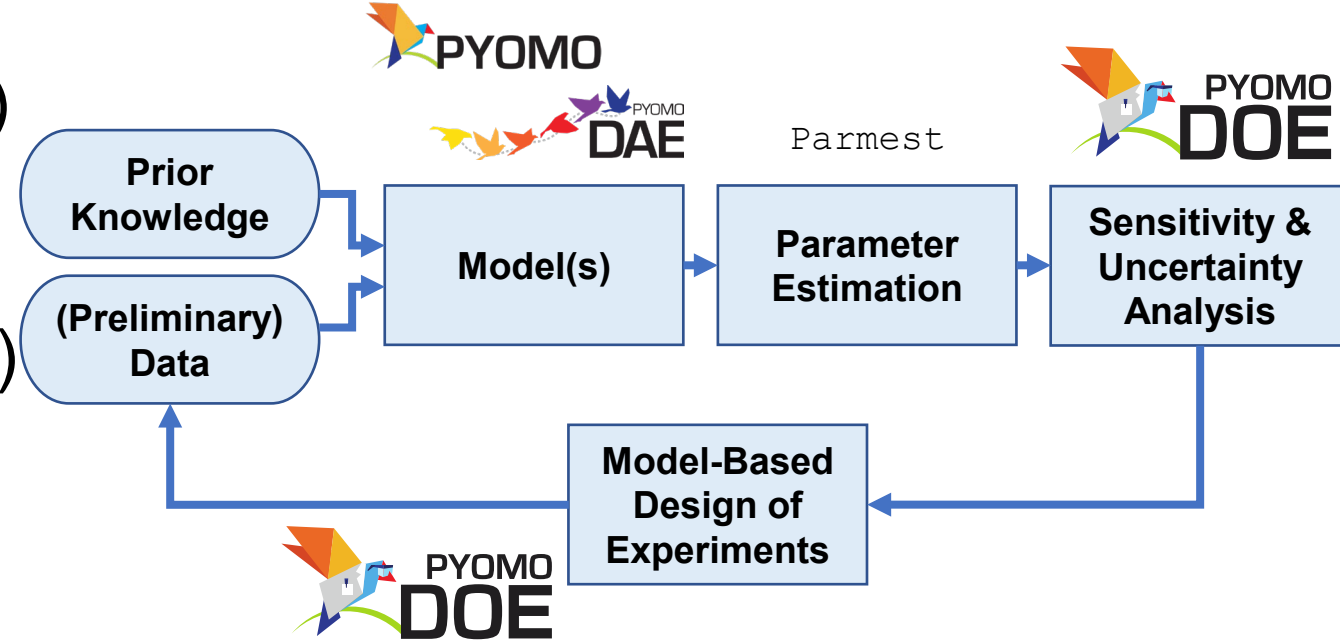
# Next Steps for Pyomo .DoE Development

Explore NLP decomposition algorithms (exploit 2-stage structure)

Optimization under uncertainty (e.g., MBDoe with uncertainty parameters)

External model function evaluations (linear algebra) to speed up Pyomo.DoE, solve measurement optimization MINLP

Refactor Pyomo .DoE and `parmes` interfaces to improve user experience



Wang, J. and Dowling, A.W. (2022), *AIChE J.* e17813.

Wang K., Zhang M., Wang, J., Shang, W., Zhang, Y., Luo, T., Dowling, A.W. (2023), *Digital Chemical Engineering*

Befort, B.J., Garciadiego, A., Wang, J., Wang, K., Maginn, E.J., Dowling, A.W. (2023), *Fluid Phase Equilibria*.

And more to come!

# Acknowledgements

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This project was funded by the Department of Energy, National Energy Technology Laboratory an agency of the United States Government, through a support contract. Neither the United States Government nor any agency thereof, nor any of their employees, nor the support contractor, nor any of their employees, makes any warranty, express or implied, or assumes any legal liability or responsibility for the accuracy, completeness, or usefulness of any information, apparatus, product, or process disclosed, or represents that its use would not infringe privately owned rights. Reference herein to any specific commercial product, process, or service by trade name, trademark, manufacturer, or otherwise does not necessarily constitute or imply its endorsement, recommendation, or favoring by the United States Government or any agency thereof. The views and opinions of authors expressed herein do not necessarily state or reflect those of the United States Government or any agency thereof.

## **Pyomo Team (SNL):**

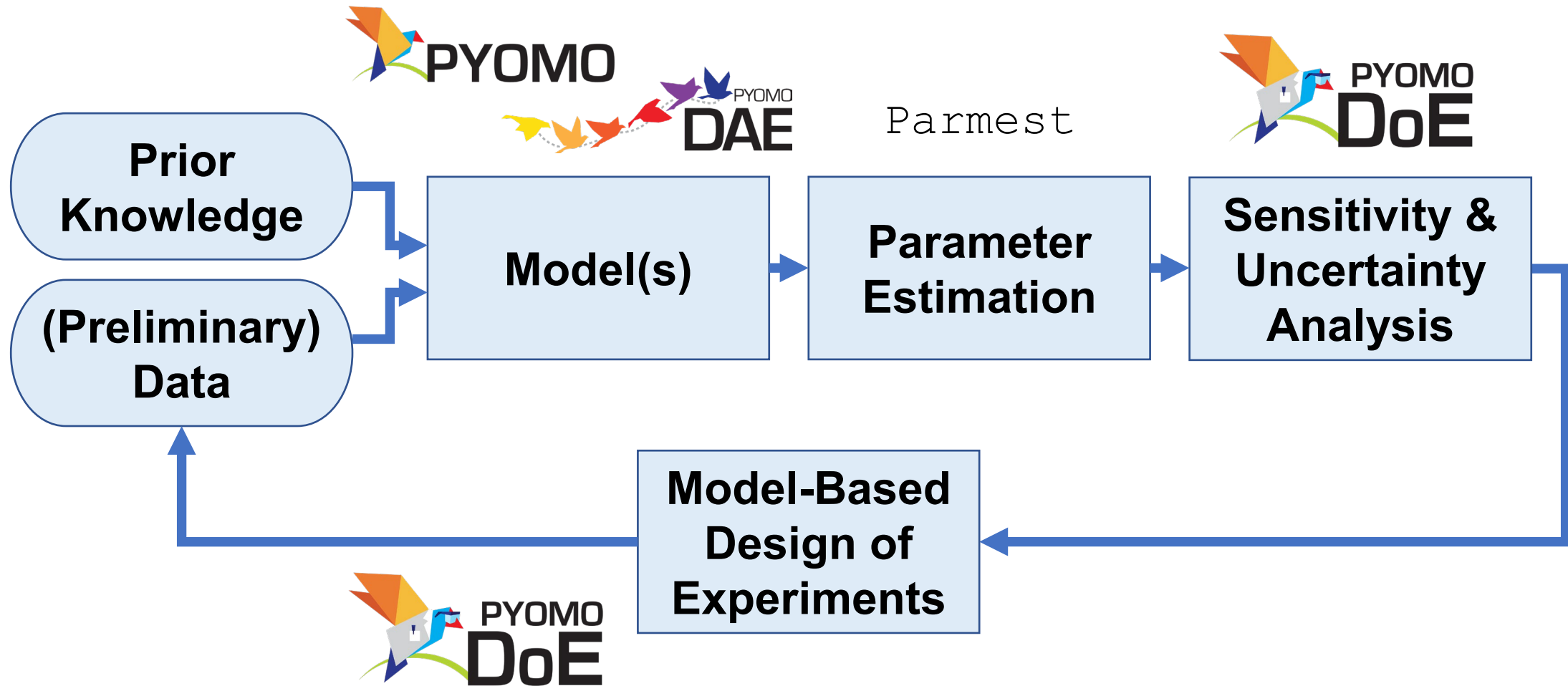
- Bethany Nicholson
- Miranda Mundt
- John Siirola

## **Membrane Case Study (ND):**

- William Phillip
- Jonathan Ouimet

# Take Away Message

*While there is some “art” to building science-based models...*



*...statistical tools can maximize the usefulness of data*