

Extending OpenKIM with an Uncertainty Quantification Toolkit for Molecular Modeling

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Karls, Mingjian Wen

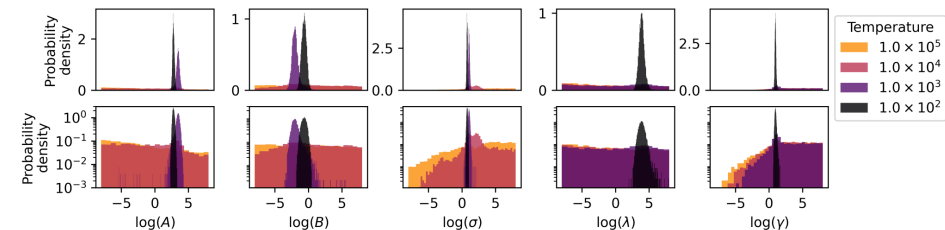
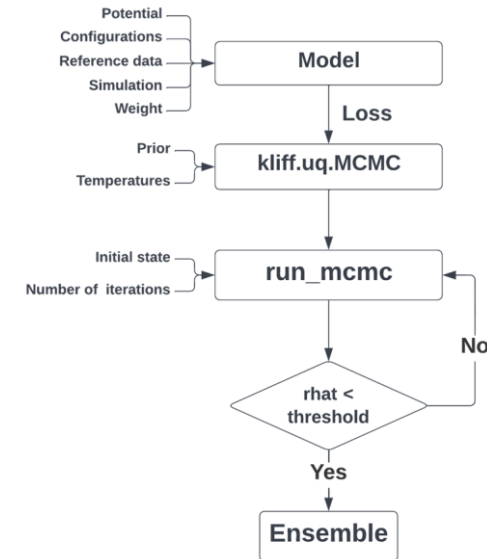
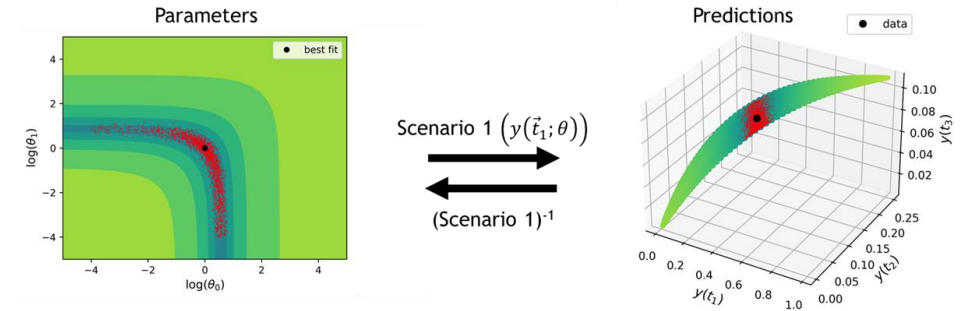
Contact: kurniawanyo@outlook.com

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Outline

1. The OpenKIM project
2. Introduction to uncertainty quantification
3. UQ extension to KLIFF
4. Demonstration: Study of SW potential
5. Conclusion and Future work



Welcome to the Open Knowledgebase of Interatomic Models!

OpenKIM is a curated repository of interatomic potentials and analytics for making classical molecular simulations of materials reliable, reproducible, and accessible. Content on OpenKIM is open source and freely available. [Read more](#)



OpenKIM is funded by the NSF.

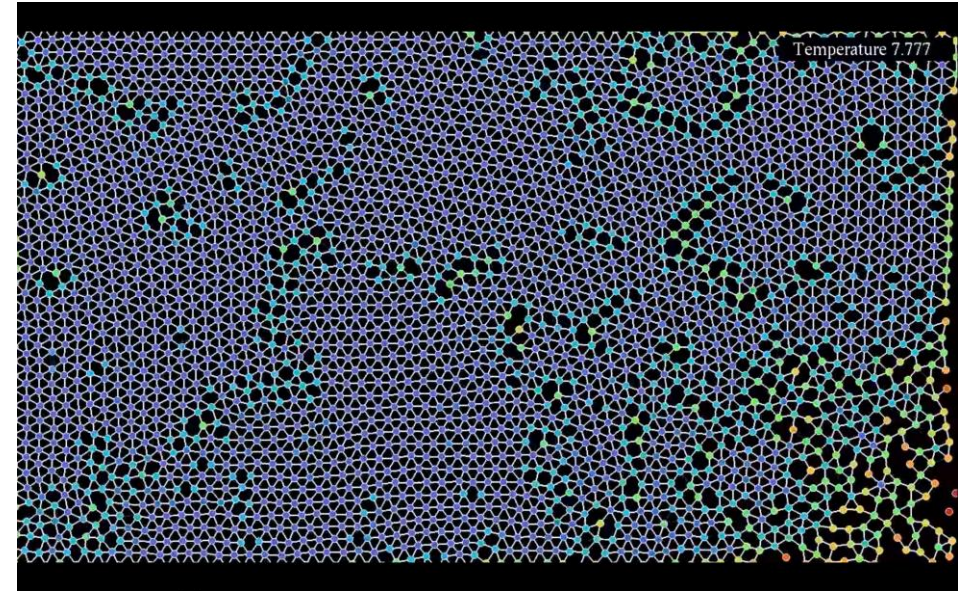
Image of a twisted graphene bilayer generated by Moon-ki Choi, University of Minnesota

The OpenKIM project

<https://openkim.org/>

Interatomic potential

- In atomistic scale simulation, the atoms are treated as classical particles.
- Interatomic potential (IP) approximates interaction energy between atoms.
- IPs are developed for specific applications, resulting in plethora of potentials.
- The functional forms of these potentials have limited scope, miss some physics, and thus introduce model errors.



(Berglund, *Freezing and melting at the molecular scale: a representation with atomic bonds* 2021 <https://www.youtube.com/watch?v=LdTDlpRx0XQ>)

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- 18th IEEE International eScience Conference

<https://openkim.org/>

KIM-based Learning-Integrated Fitting Framework

- KLIFF is a general-purpose fitting framework for IPs.
- KLIFF employs the force-matching algorithm [1].
- The IPs are trained to match atomic forces of several configurations from first-principle simulation.
- The trained IPs conform to the KIM API.



<https://kliff.readthedocs.io/>

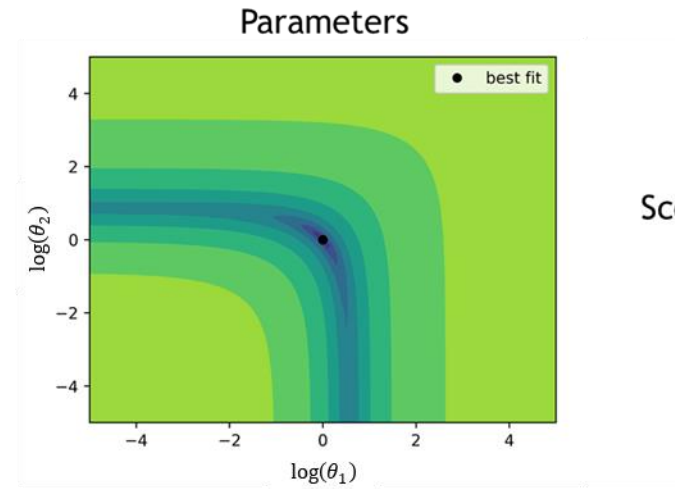
[1] F. Ercolessi and J. B. Adams, "Interatomic Potentials from First-Principles Calculations: The Force-Matching Method," *EPL*, vol. 26, no. 8, p. 583, Jun. 1994, doi: [10.1209/0295-5075/26/8/005](https://doi.org/10.1209/0295-5075/26/8/005).

Contribution

- We integrate an uncertainty quantification framework into KLIFF.

Goal

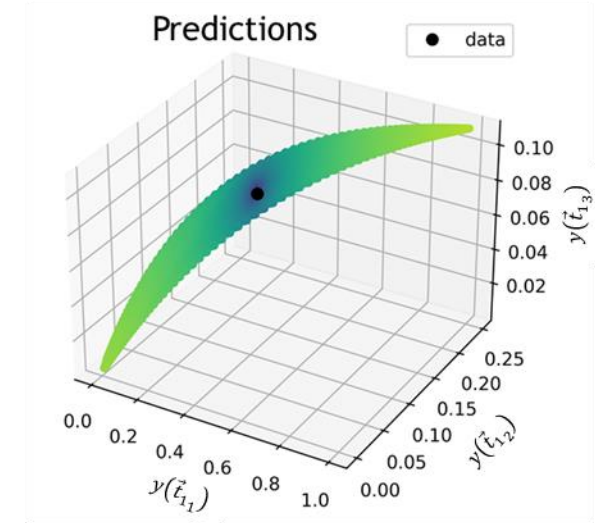
- We want to facilitate UQ studies for IPs.
- We hope that this integration can lead to more transparent and reproducible UQ analysis for IPs.



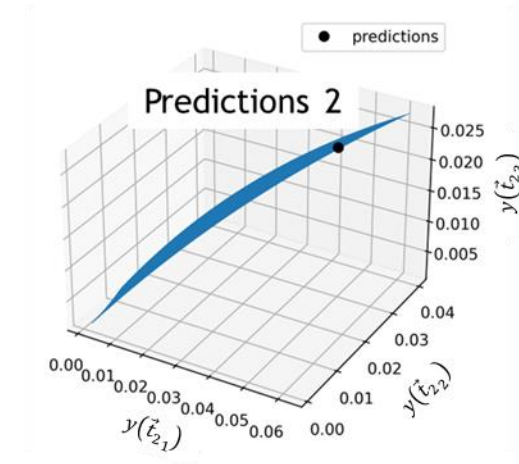
Scenario 1 ($y(\vec{t}_1; \theta)$)



(Scenario 1)⁻¹

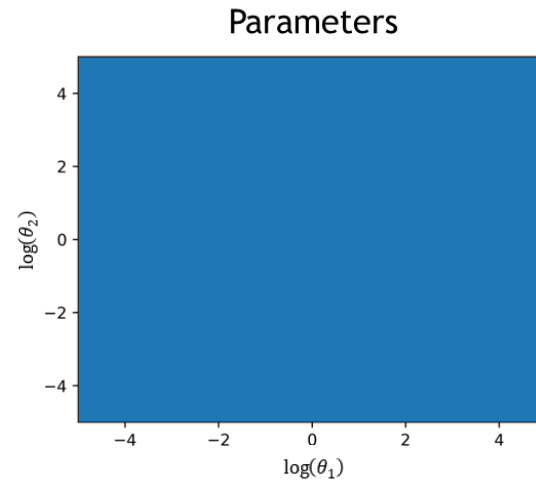
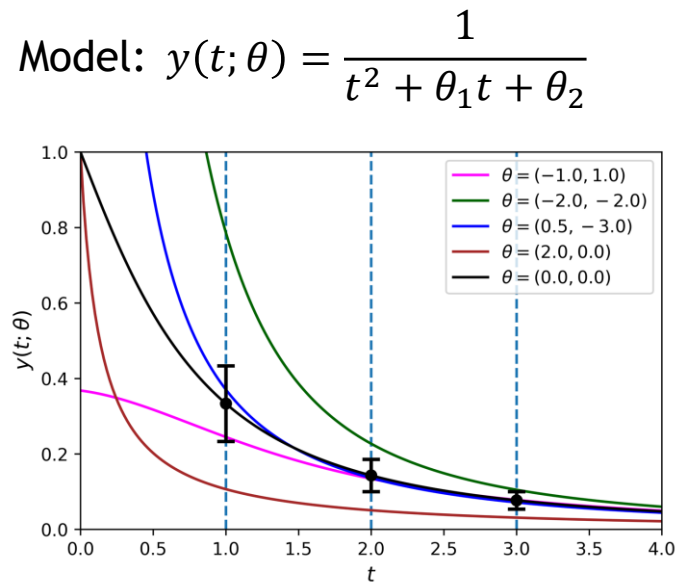


Scenario 2 ($y(\vec{t}_2; \theta)$)

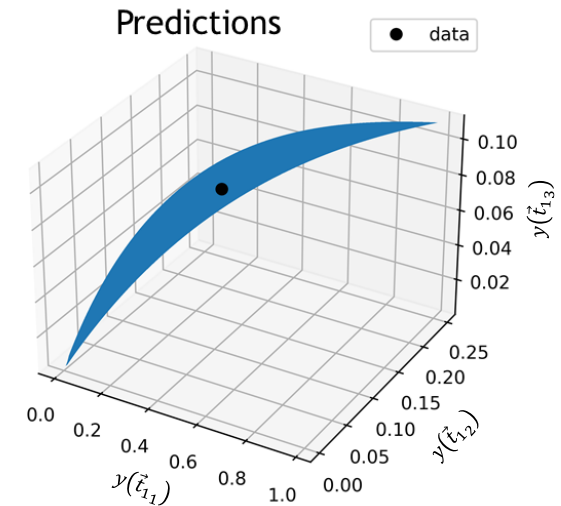


Introduction to uncertainty quantification

Geometry of a model

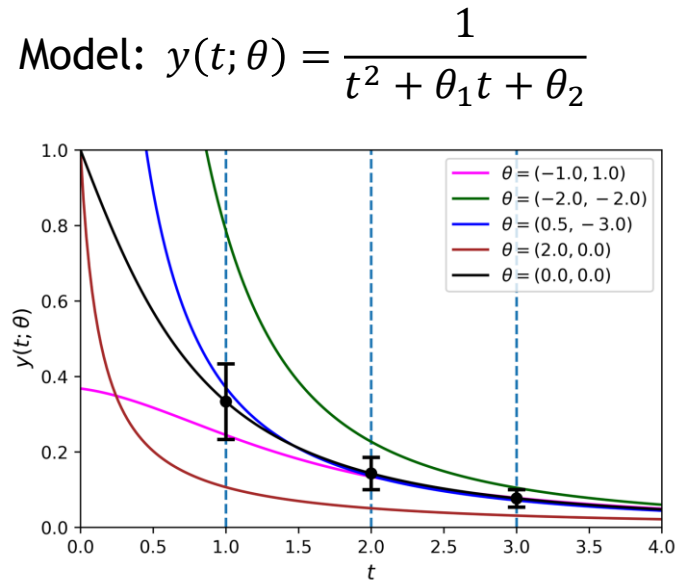


Scenario 1 ($y(\vec{t}_1; \theta)$)



- Model is a mapping from a parameter space to a prediction space.
- The model manifold is the range of the model map.

Loss function



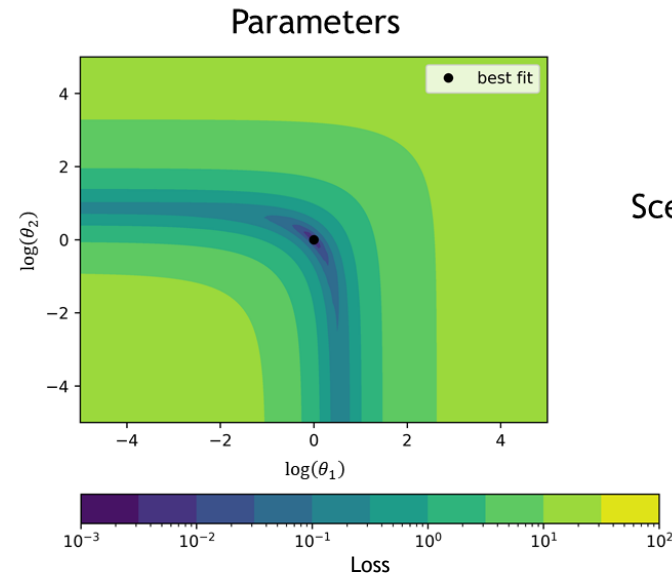
Assumptions:

$$d_m = y(t_m; \theta) + \xi_m$$

$$\xi_m \sim \mathcal{N}(0, \sigma_m)$$

Loss function:

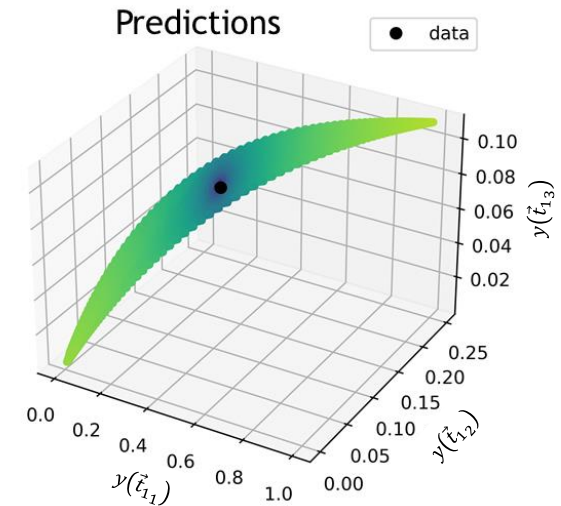
$$L(\theta) = \frac{1}{2} \sum_m \left(\frac{d_m - y(t_m; \theta)}{\sigma_m} \right)^2$$



Scenario 1 ($y(\vec{t}_1; \theta)$)

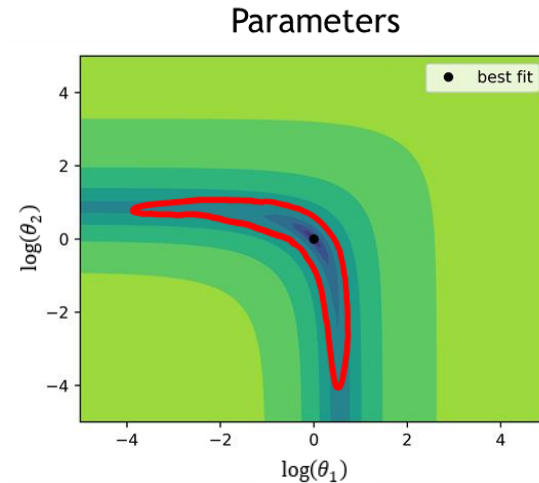
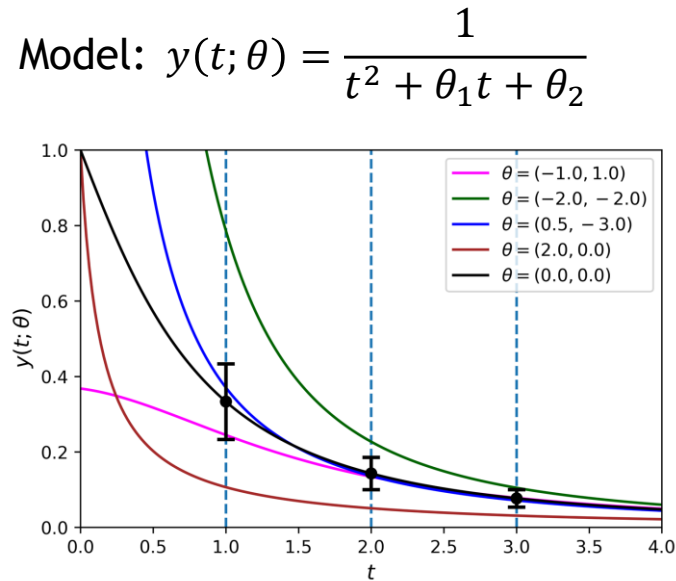
↔

(Scenario 1)⁻¹



- Loss function measures the quality of model predictions compared to the observed data.
- The best fit parameters minimize the loss function.

Uncertainty quantification

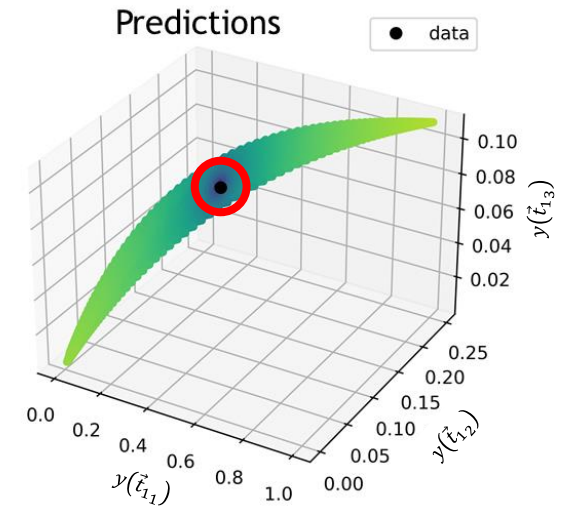


Distribution
of data

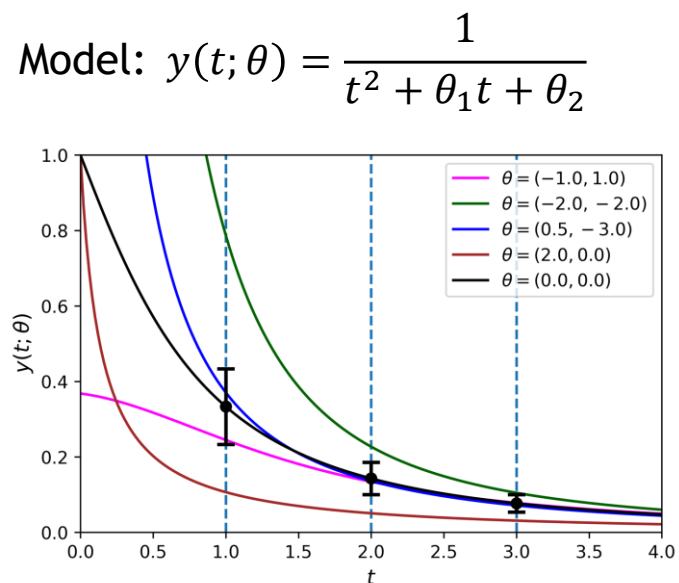


Distribution
of parameters

Scenario 1 ($y(\vec{t}_1; \theta)$)
 \longleftrightarrow
 (Scenario 1) $^{-1}$



Uncertainty quantification



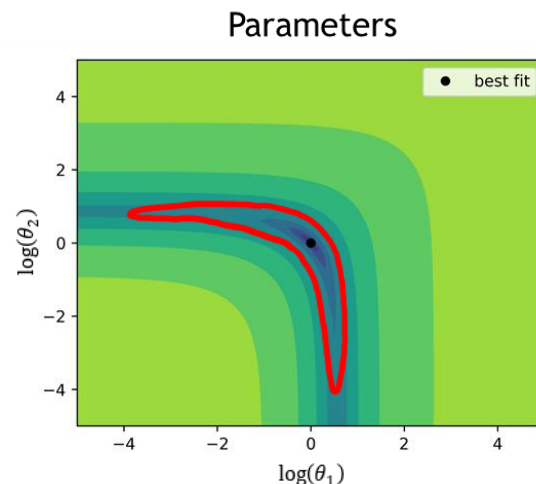
Assumptions:

$$d_m = y(t_m; \theta) + \xi_m$$

$$\xi_m \sim \mathcal{N}(0, \sigma_m)$$

Loss function:

$$L(\theta) = \frac{1}{2} \sum_m \left(\frac{d_m - y(t_m; \theta)}{\sigma_m} \right)^2$$



Distribution
of data

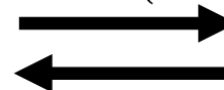


Distribution
of parameters

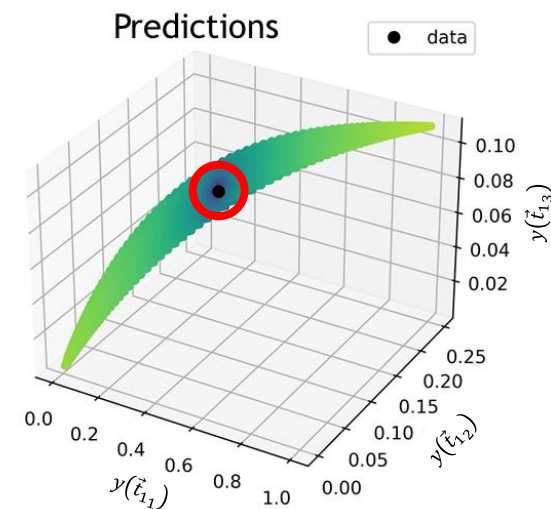


Distribution
of predictions

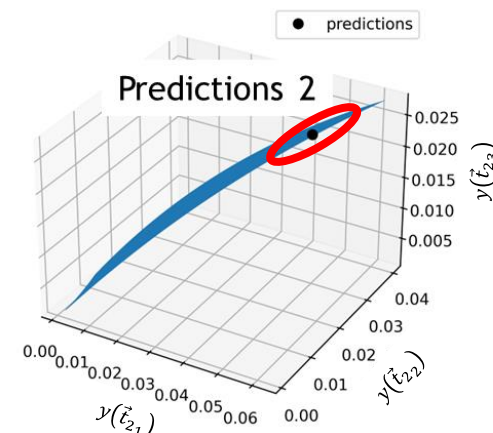
Scenario 1 ($y(\vec{t}_1; \theta)$)



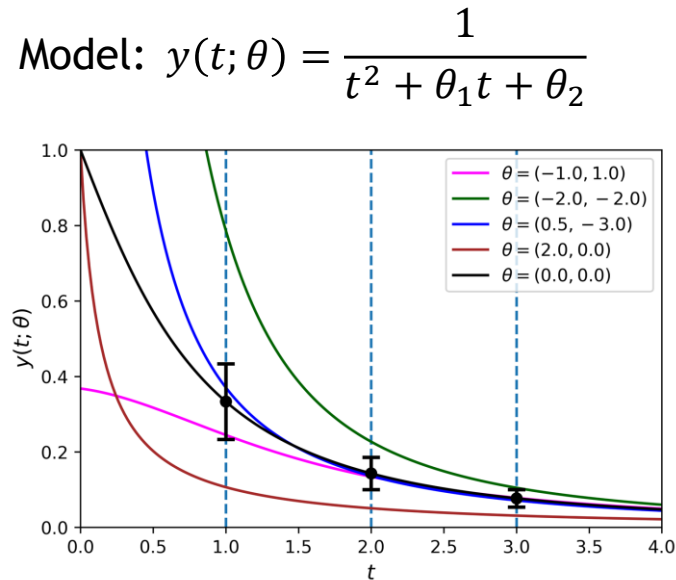
(Scenario 1)⁻¹



Scenario 2 ($y(\vec{t}_2; \theta)$)



Markov Chain Monte Carlo



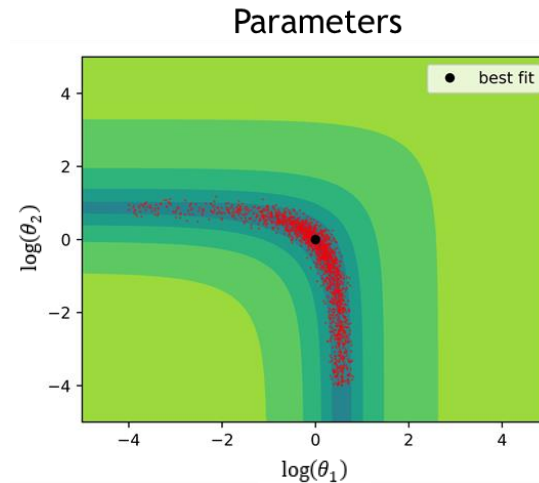
Assumptions:

$$d_m = y(t_m; \theta) + \xi_m$$

$$\xi_m \sim \mathcal{N}(0, \sigma_m)$$

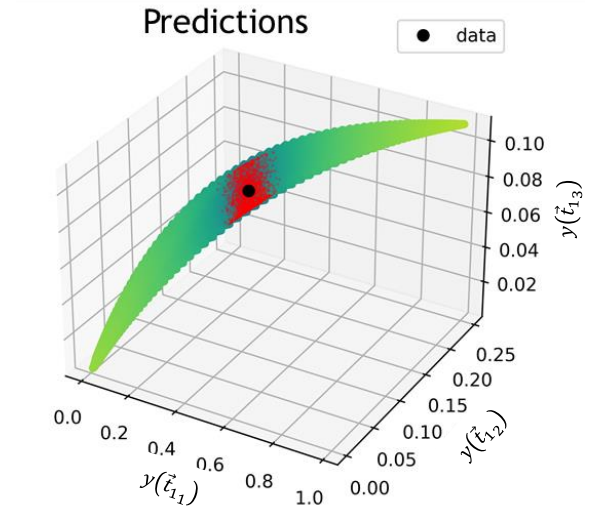
Loss function:

$$L(\theta) = \frac{1}{2} \sum_m \left(\frac{d_m - y(t_m; \theta)}{\sigma_m} \right)^2$$



Scenario 1 ($y(\vec{t}_1; \theta)$)

Scenario 1⁻¹



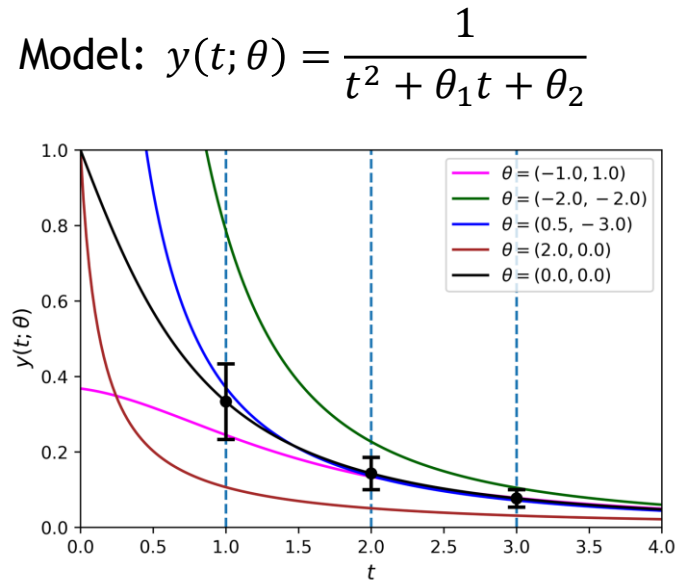
- Bayes' rule:

$$P(\theta | \vec{d}) \propto \mathcal{L}(\theta | \vec{d}) \times \pi(\theta),$$

$$\mathcal{L}(\theta | \vec{d}) \propto \exp(-L(\theta))$$

- Use MCMC algorithm to sample the posterior $P(\theta | \vec{d})$.

Markov Chain Monte Carlo



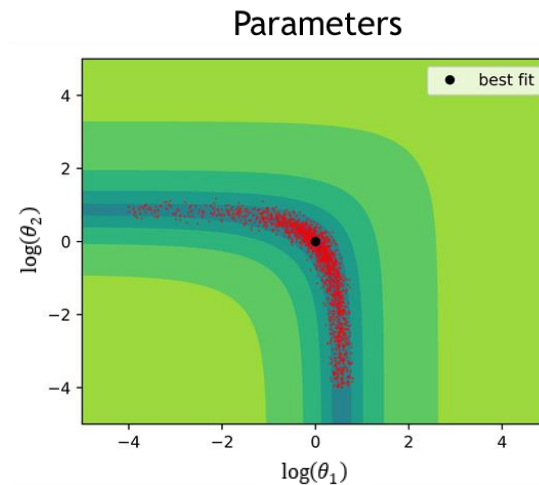
Assumptions:

$$d_m = y(t_m; \theta) + \xi_m$$

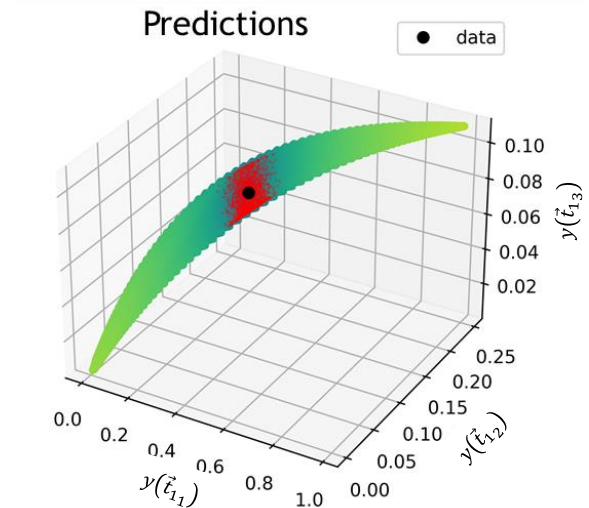
$$\xi_m \sim \mathcal{N}(0, \sigma_m)$$

Loss function:

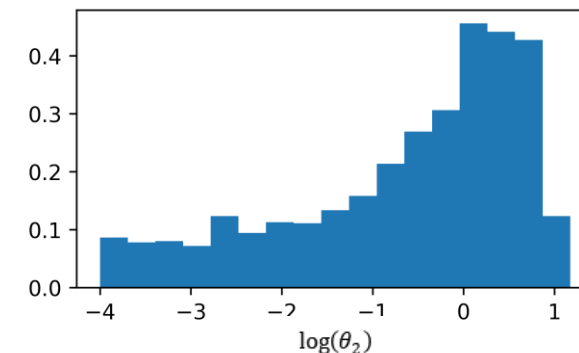
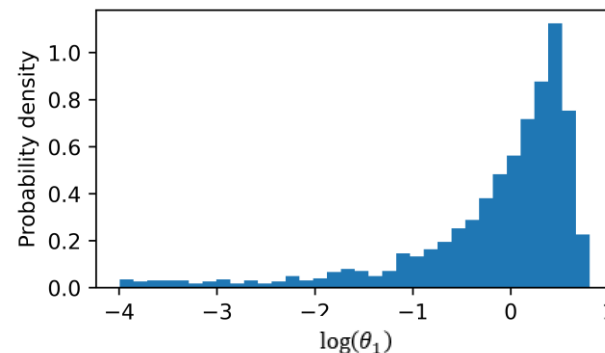
$$L(\theta) = \frac{1}{2} \sum_m \left(\frac{d_m - y(t_m; \theta)}{\sigma_m} \right)^2$$



Scenario 1 ($y(\vec{t}_1; \theta)$)
 \longleftrightarrow
 (Scenario 1) $^{-1}$

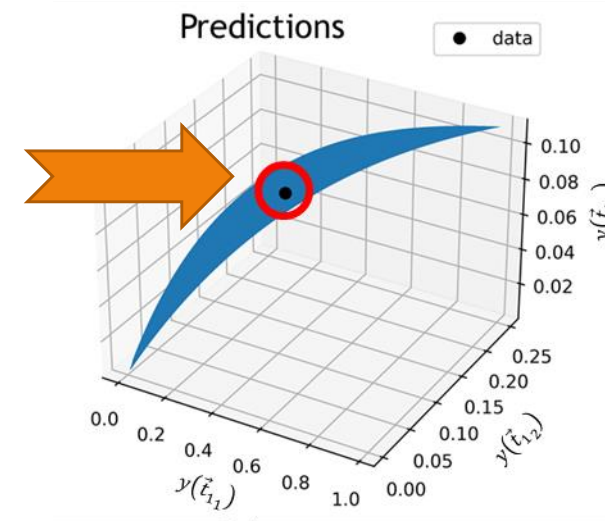


- Distribution of the parameters is inferred from the resulting samples.



Model inadequacy

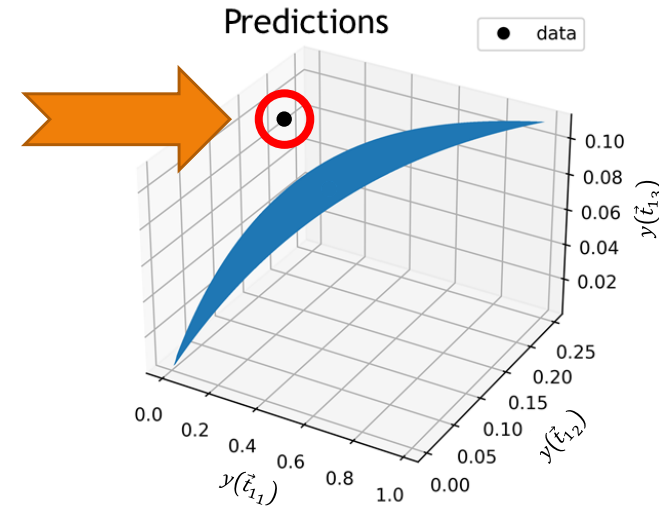
- $\mathcal{L}(\theta|\vec{d}) \propto \exp(-L(\theta))$ assumes the model can reproduce the data within the error bar.
- The high-density circle/sphere intersects the manifold.



Adequate
model

Model inadequacy

- In some cases, this assumption is invalid.
- The data is far from the manifold; the high-density circle/sphere doesn't intersect the manifold.
- We need to fix the UQ formulation to include model inadequacy.



Inadequate
model

Model inadequacy

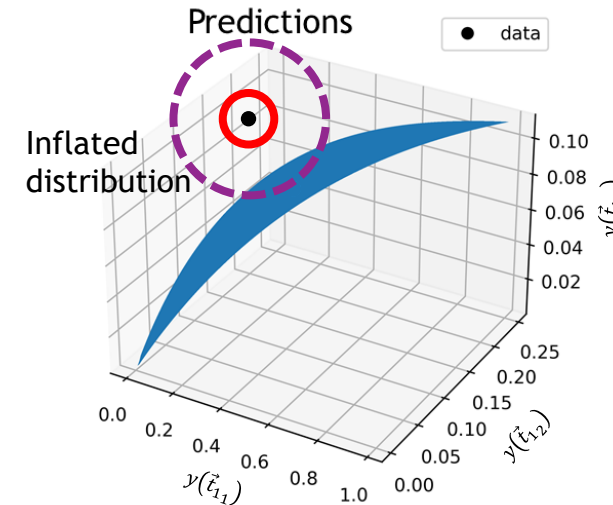
- Suggestion: Inflate the likelihood [2]:

$$\mathcal{L}(\theta|\vec{d}) \propto \exp\left(-\frac{L(\theta)}{T_0}\right), \quad T_0 = \frac{2L_0}{N}$$

$$L(\theta) = \frac{1}{2} \sum_m \left(\frac{d_m - y(t_m; \theta)}{\sigma_m} \right)^2$$

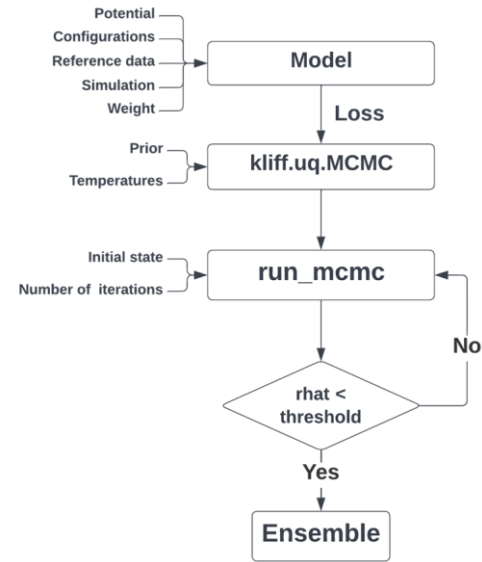
$L_0 \equiv$ minimum loss

$N \equiv$ number of parameters.



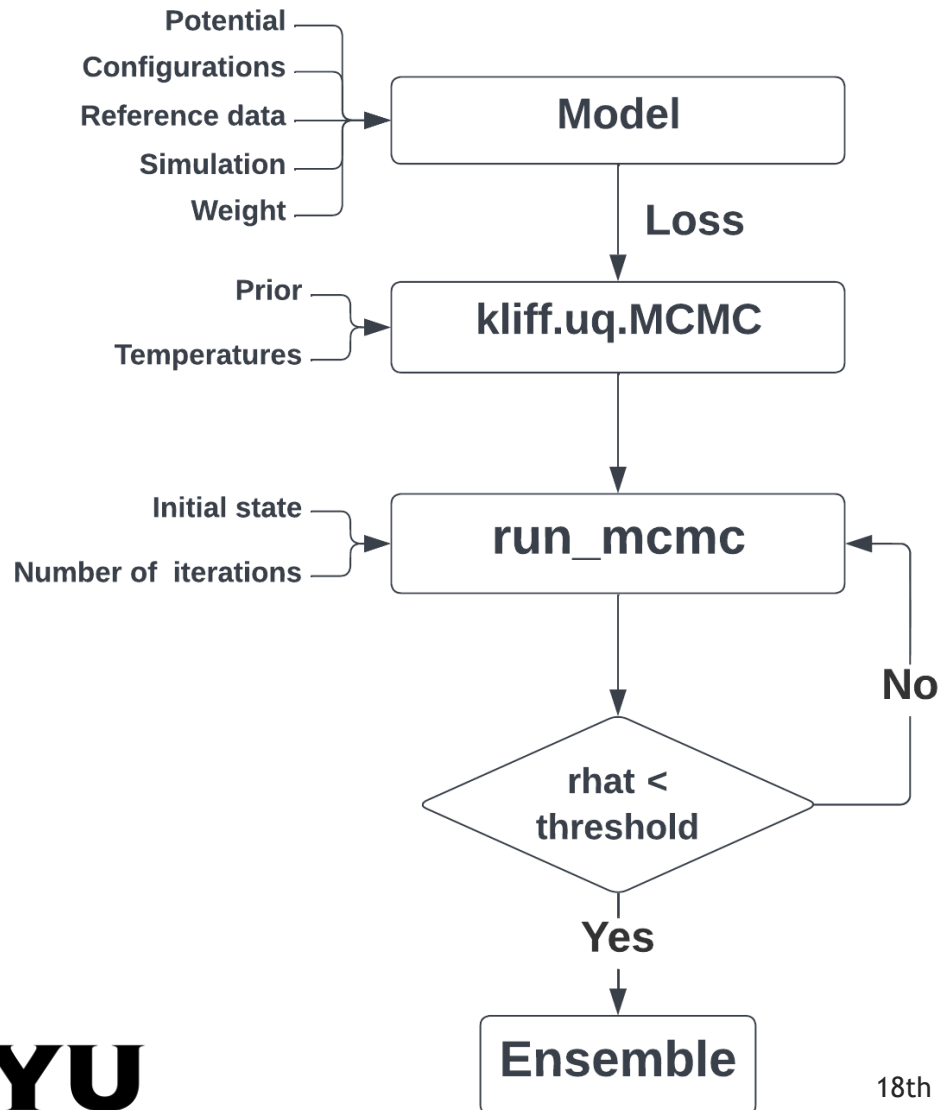
Inadequate model

[2] P. Pernot and F. Cailliez, "A critical review of statistical calibration/prediction models handling data inconsistency and model inadequacy," *AIChE Journal*, vol. 63, no. 10, pp. 4642-4665, 2017, doi: [10.1002/aic.15781](https://doi.org/10.1002/aic.15781).



UQ extension to KLIFF

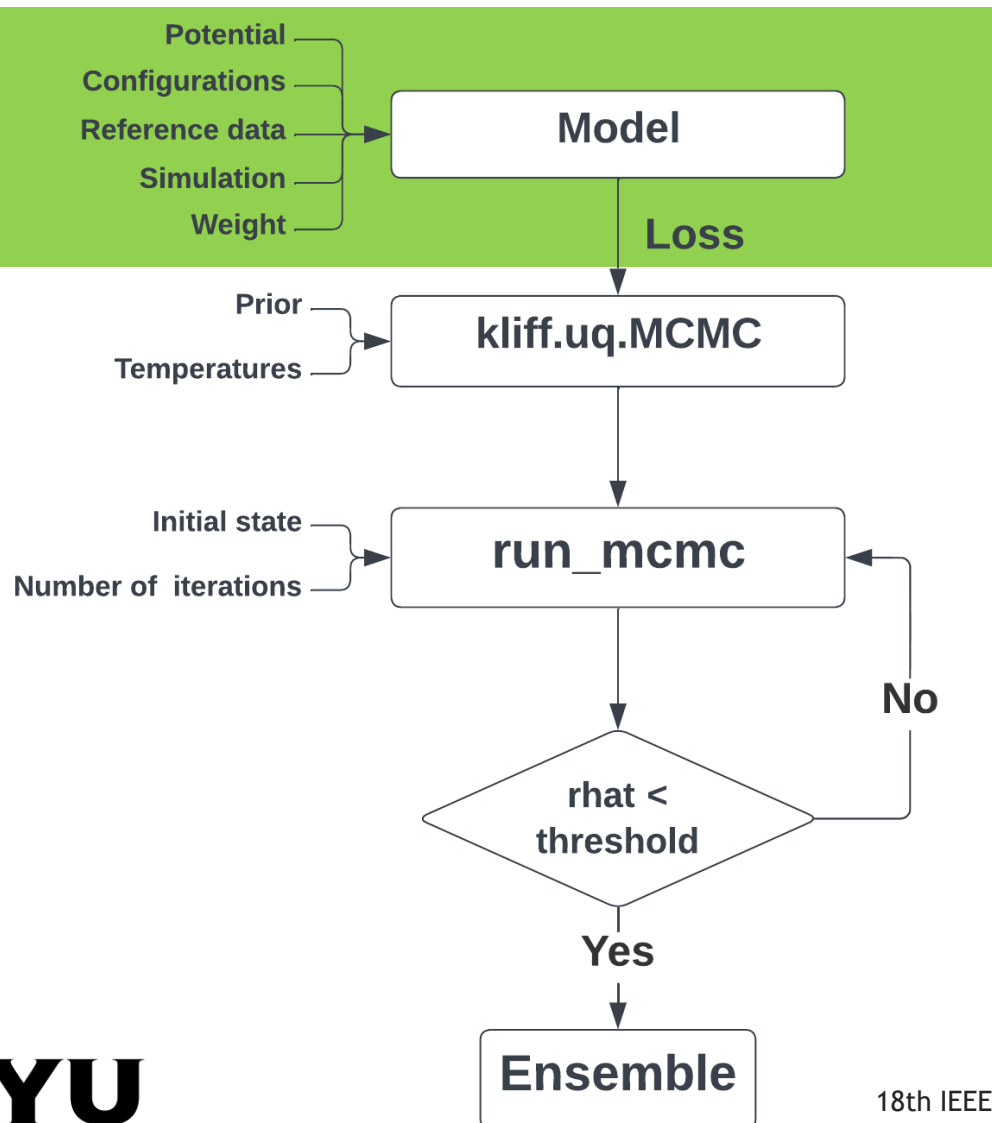
Implementation and workflow



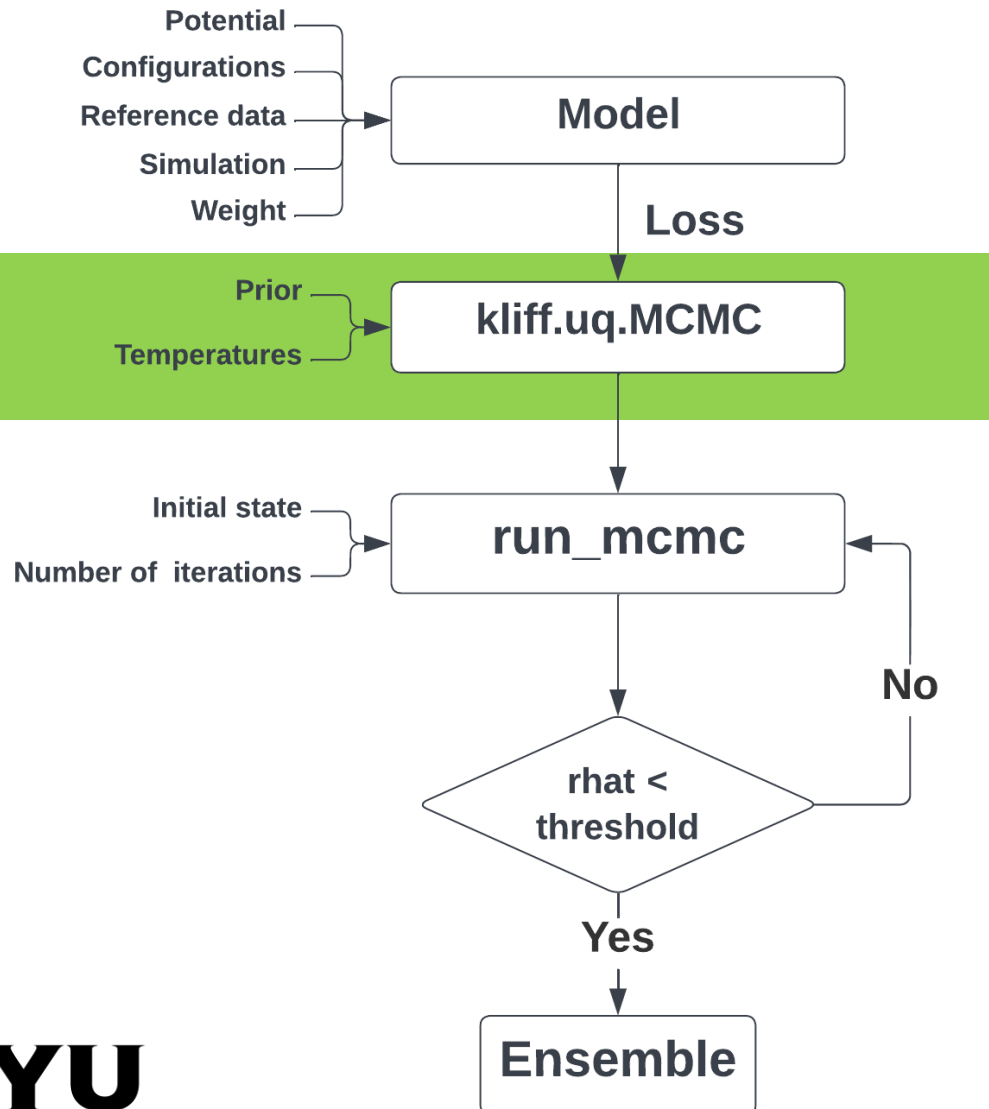
- We extend KLIFF to include uncertainty quantification functionality.
- This integration can:
 - Facilitate UQ studies for IPs.
 - Lead to more transparent and reproducible UQ analysis for IPs.
- KLIFF uses MCMC method.
- Other UQ methods will be implemented in the future.

1. Defining the model and loss function

- This functionality has been implemented previously and is not part of this integration.
- For more detail, visit <https://kliff.readthedocs.io/>.



2. Instantiating the posterior sampler



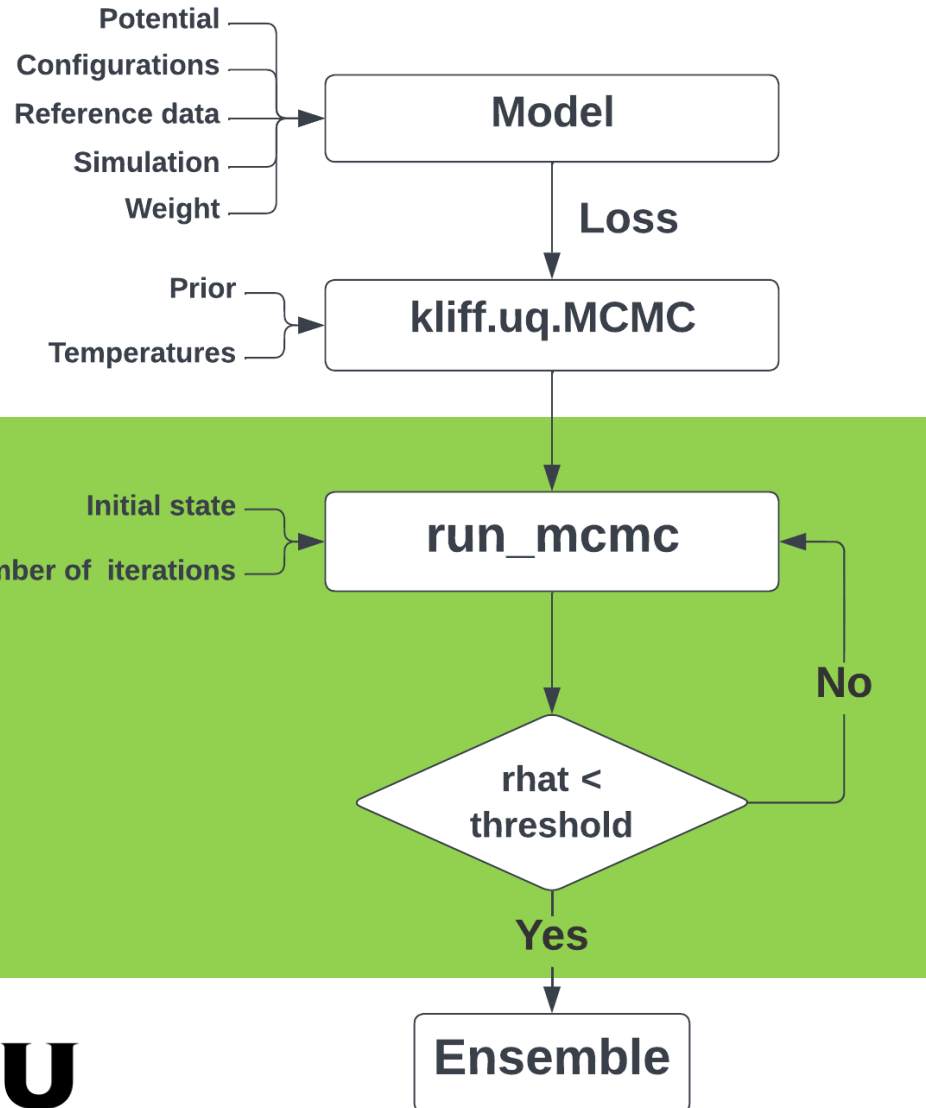
- We use ptemcee [3, 4] to perform parallel-tempered MCMC:
 - Simulating multiple different sampling temperatures, each with multiple chains/walkers.
- Parallel tempering improves convergence.
- Parallel tempering also allows us to explore how sampling results evolve with different scale of model error.
- Recommendation: Set the temperature ladder to be logarithmically spaced from 1.0 to few times larger than T_0 .

[3] W. Vousden, “Willvousden/ptemcee: A parallel-tempered version of emcee.” *GitHub*. [Online]. Available:

<https://github.com/willvousden/ptemcee>. [Accessed: 14-Sep-2022].

[4] W. D. Vousden, W. M. Farr, and I. Mandel, “Dynamic temperature selection for parallel tempering in Markov chain Monte Carlo simulations,” *Monthly Notices of the Royal Astronomical Society*, vol. 455, no. 2, pp. 1919-1937, Jan. 2016, doi: [10.1093/mnras/stv2422](https://doi.org/10.1093/mnras/stv2422).

3. Running MCMC & monitoring convergence



- Convergence is monitored by computing the Potential Scale Reduction Factor (PSRF) [5]:

$$\hat{R}^p = \frac{K-1}{K} + \frac{J+1}{J} \lambda_{\max}(W^{-1} B/K)$$

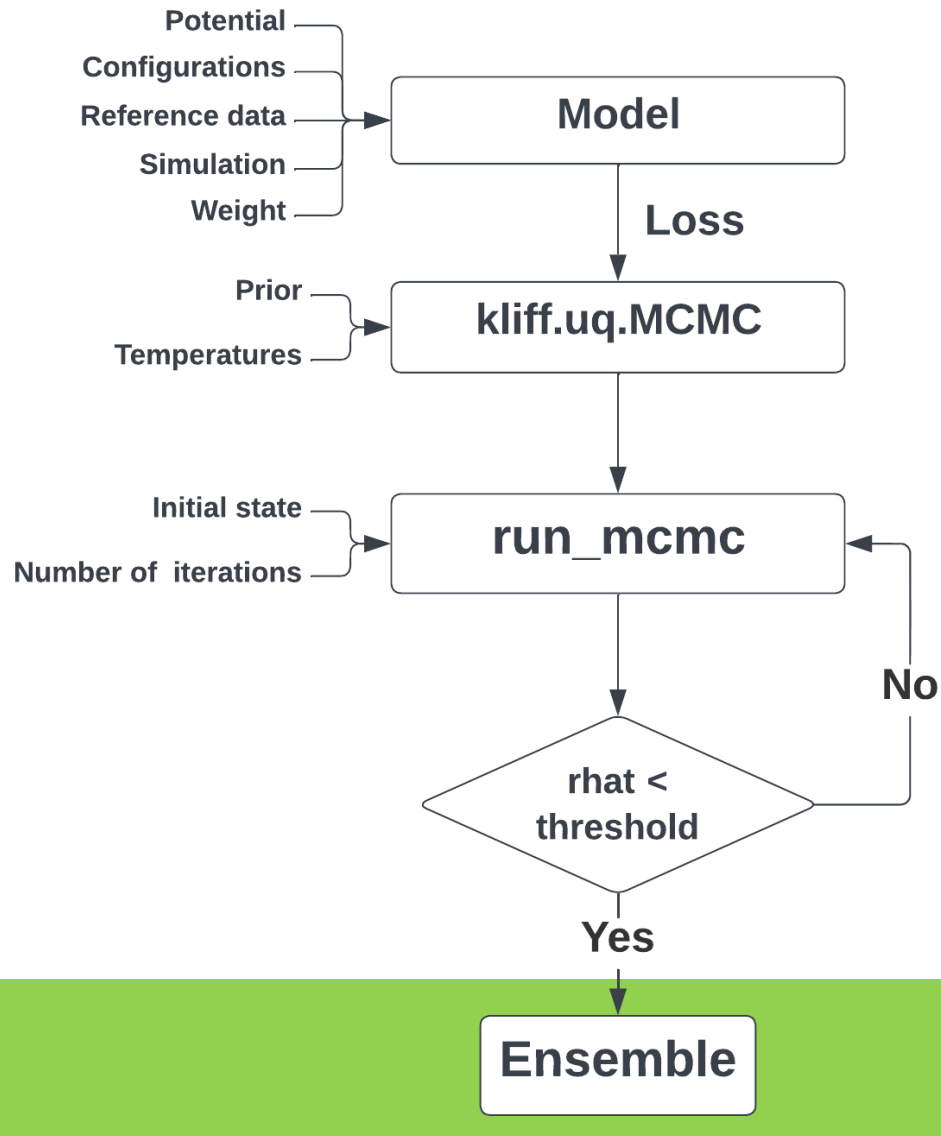
$$\frac{B}{K} = \frac{1}{J-1} \sum_{j=1}^J (\bar{\psi}_j - \bar{\psi})(\bar{\psi}_j - \bar{\psi})^T$$

$$W = \frac{1}{J(K-1)} \sum_{j=1}^J \sum_{k=1}^K (\psi_{jk} - \bar{\psi}_j)(\psi_{jk} - \bar{\psi}_j)^T$$

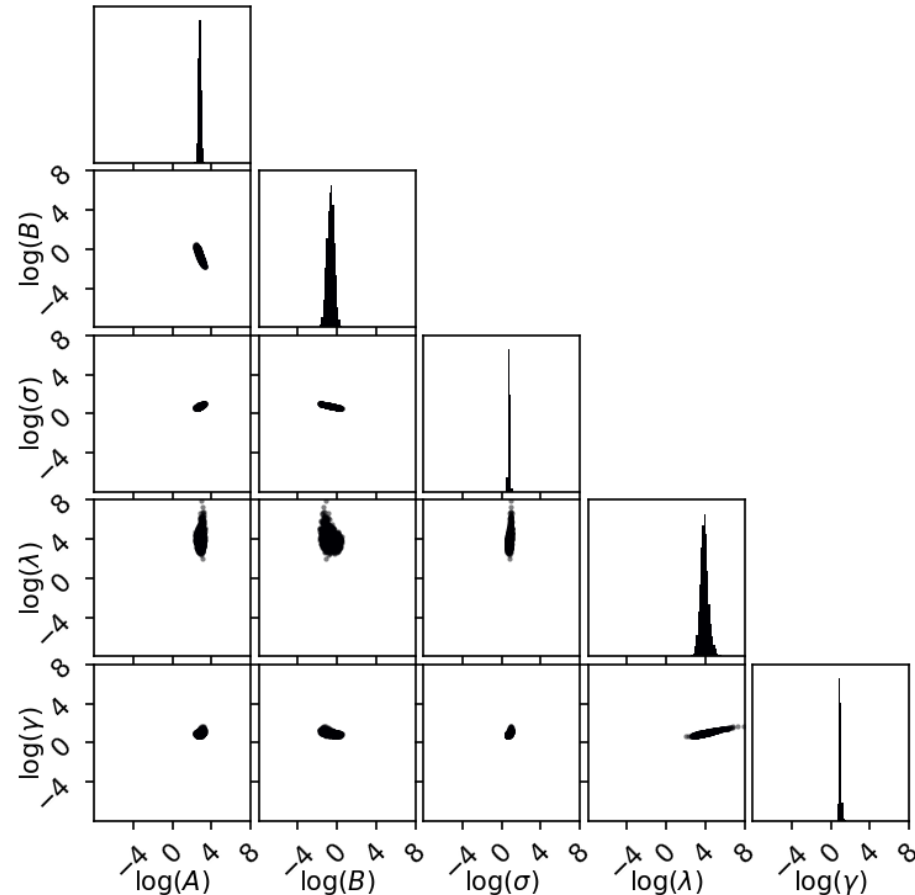
- When samples have converged, $\hat{R}^p \rightarrow 1$ (common threshold is 1.1).

[5] S. P. Brooks and A. Gelman, "General Methods for Monitoring Convergence of Iterative Simulations," *Journal of Computational and Graphical Statistics*, vol. 7, no. 4, pp. 434-455, Dec. 1998, doi: [10.1080/10618600.1998.10474787](https://doi.org/10.1080/10618600.1998.10474787).

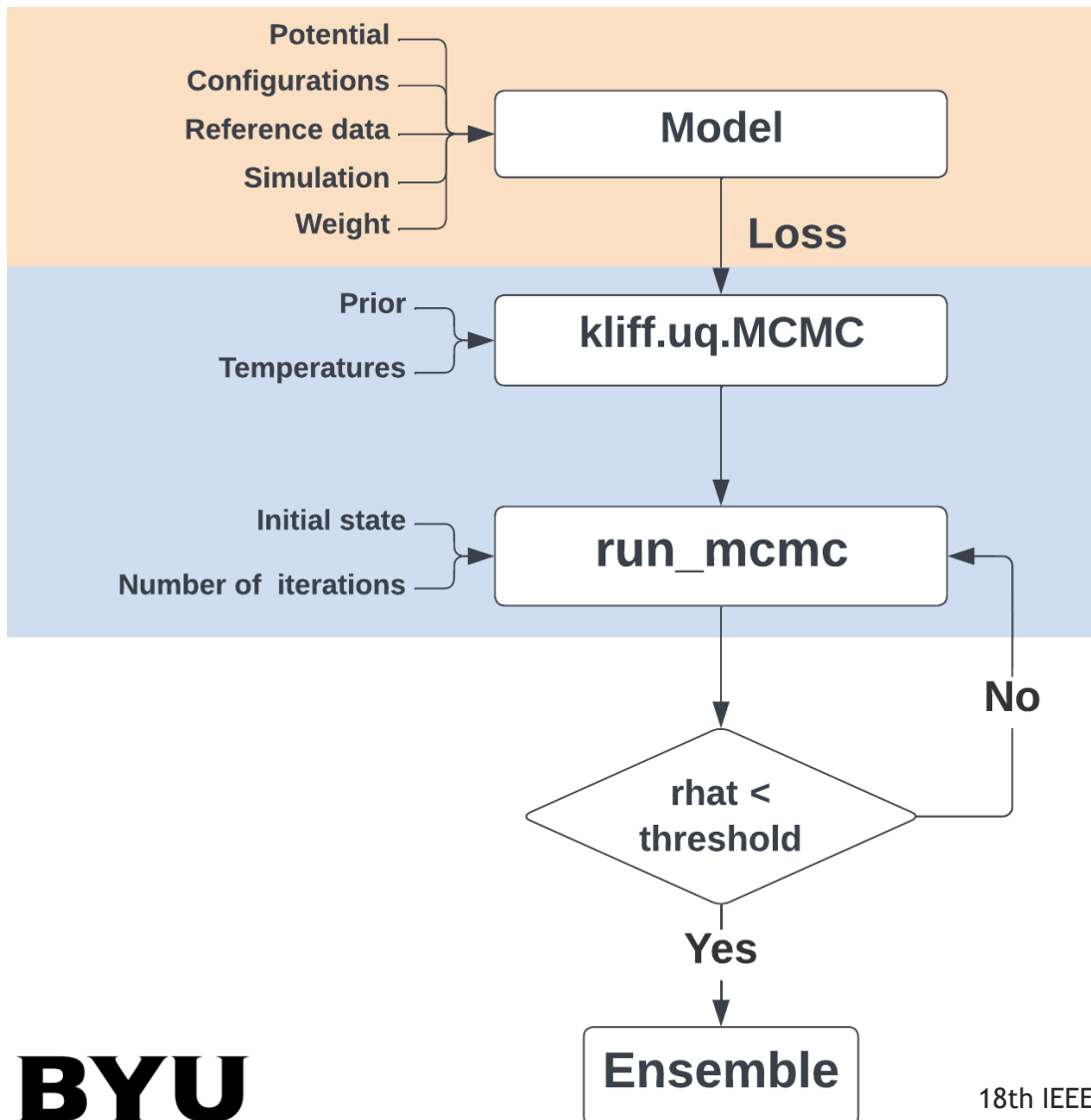
4. Retrieving the ensemble



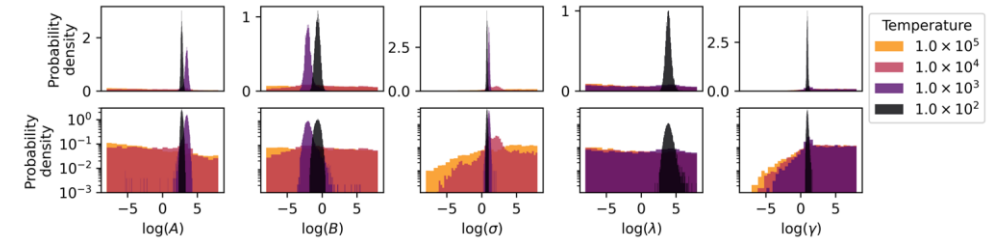
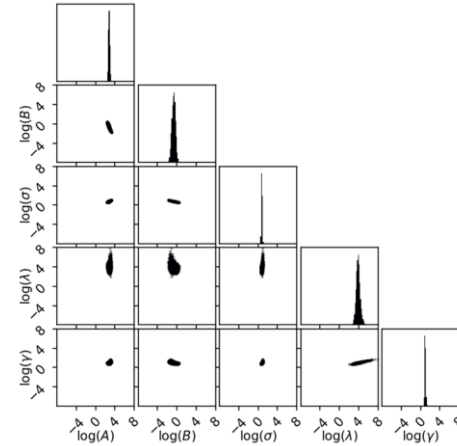
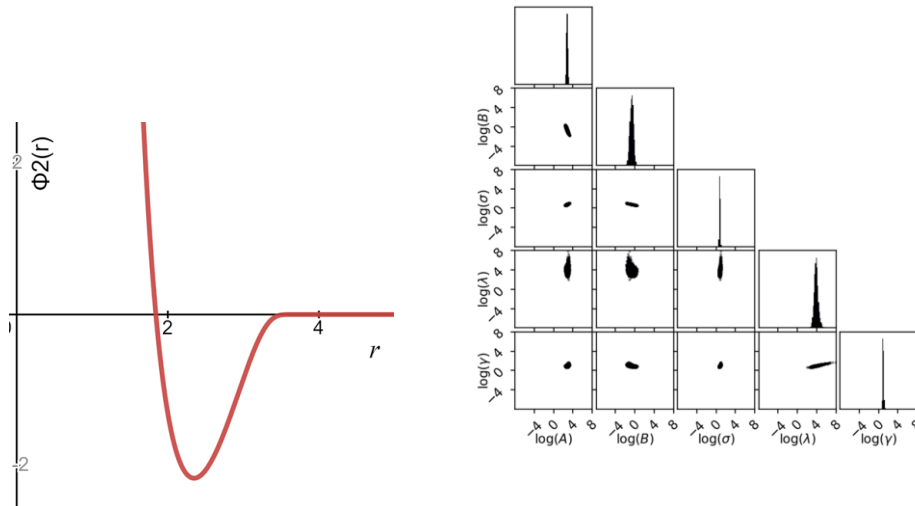
- The distribution of the parameters is inferred from the ensemble.



Parallel computing



- Parallelization can be done in 2 places:
 - Loss evaluation (over configurations)
 - MCMC sampling (over walkers)
- Suggestion:
 - Use OpenMP-style parallelization for loss evaluation.
 - Use MPI-style parallelization for MCMC sampling.
- For more detail, visit <https://kliff.readthedocs.io/>.



Demonstration: Study of Stillinger-Weber potential



Access to the
example
scripts.

Stillinger-Weber potential

- Model: Stillinger-Weber potential [6]

$$\phi_2(r_{ij}) = A \left[B \left(\frac{\sigma}{r_{ij}} \right)^p - \left(\frac{\sigma}{r_{ij}} \right)^q \right] \exp \left(\frac{\sigma}{r_{ij} - r^{cut}} \right)$$

$$\phi_3(r_{ij}, r_{ik}, \beta_{jik}) = \lambda [\cos(\beta_{jik}) - \cos(\beta^0)]^2 \times \exp \left(\frac{\gamma}{r_{ij} - r^{cut}} + \frac{\gamma}{r_{ik} - r^{cut}} \right)$$

- Parameters: $\log(A)$, $\log(B)$, $\log(\sigma)$, $\log(\lambda)$, $\log(\gamma)$
- Training data: energy and force of Silicon atoms in several configurations (weights \propto data values).
- Best fit:

$$A = 15.2792223 \text{ eV}$$

$$B = 0.6032372$$

$$\sigma = 2.09420085 \text{ \AA}$$

$$\lambda = 45.47927476 \text{ eV}$$

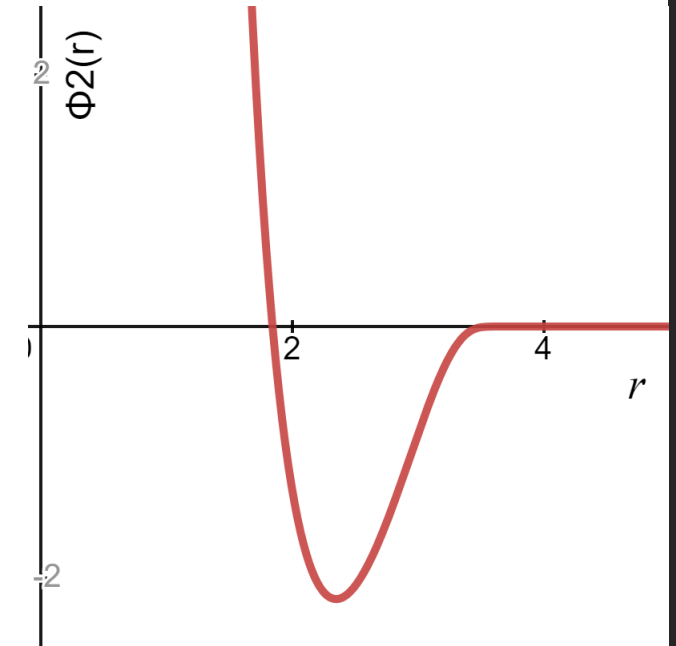
$$\gamma = 2.51306949 \text{ \AA}$$

$$p = 4$$

$$q = 0$$

$$\cos(\beta^0) = -0.33333333$$

$$r^{cut} = 3.77118 \text{ \AA}$$





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scripts.

MCMC setup

MCMC Setup

- Posterior distribution:

$$P(\theta|\vec{d}) \propto \mathcal{L}(\theta|\vec{d}) \times \pi(\theta),$$

$$\mathcal{L}(\theta|\vec{d}) \propto \exp(-L(\theta)/T)$$

- Temperatures:

- $T_0 = 1.324$
- $T \in [1, 10^7]$

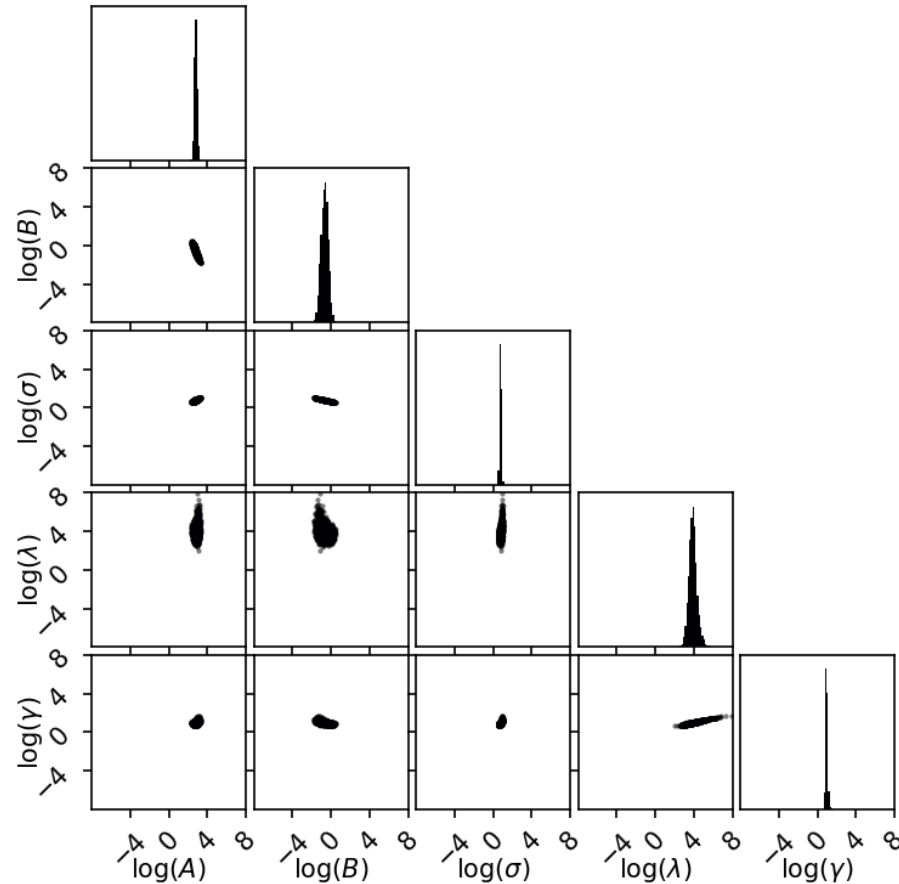
- Prior: $\log(\theta) \sim \mathcal{U}(-8, 8)$

- Run MCMC for 150,000 steps
 - Burn-in: 10,000
 - Thinning factor: 200
- Convergence test: $\hat{R}^p \leq 1.046$



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scripts.

Presenting the samples

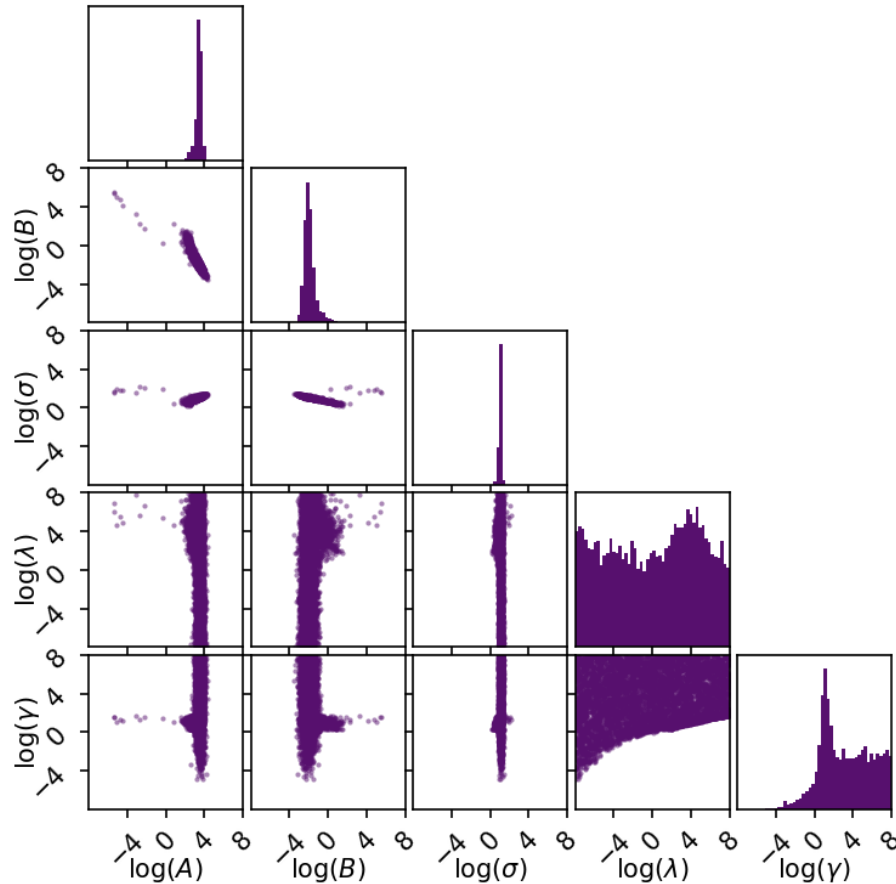


- Sampling temperature $T = 10^2$
- What's plotted:
 - Main diagonal: Marginal distribution for each parameter.
 - Below diagonal: 2D projection of the samples in parameter space.
- At lower temperature, the distributions are concentrated.



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scripts.

Parameter evaporation



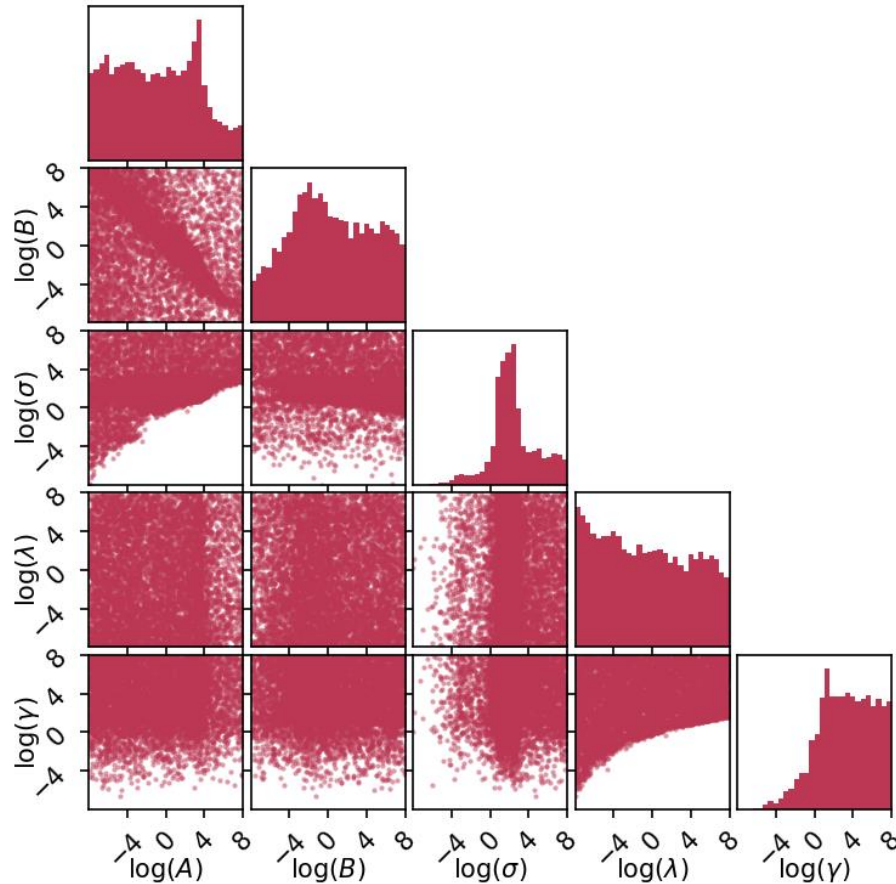
- Sampling temperature $T = 10^3$
- The distribution becomes wider as we increase the temperature.
- Parameter evaporation occurs: the walkers tend to run to sub-optimal parameter values [7, 8].
- Evaporated parameters are unconstrained by the data.

[7] M. K. Transtrum, B. B. Machta, and J. P. Sethna, “Geometry of nonlinear least squares with applications to sloppy models and optimization,” *Phys. Rev. E*, vol. 83, no. 3, p. 036701, Mar. 2011, doi: [10.1103/PhysRevE.83.036701](https://doi.org/10.1103/PhysRevE.83.036701).
[8] R. Gutenkunst, “Sloppiness, Modeling, and Evolution in Biochemical Networks,” Cornell University, Ithaca, New York, 2007. Accessed: May 14, 2021. [Online]. Available: <https://ecommons.cornell.edu/handle/1813/8206>



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example
scripts.

Parameter evaporation



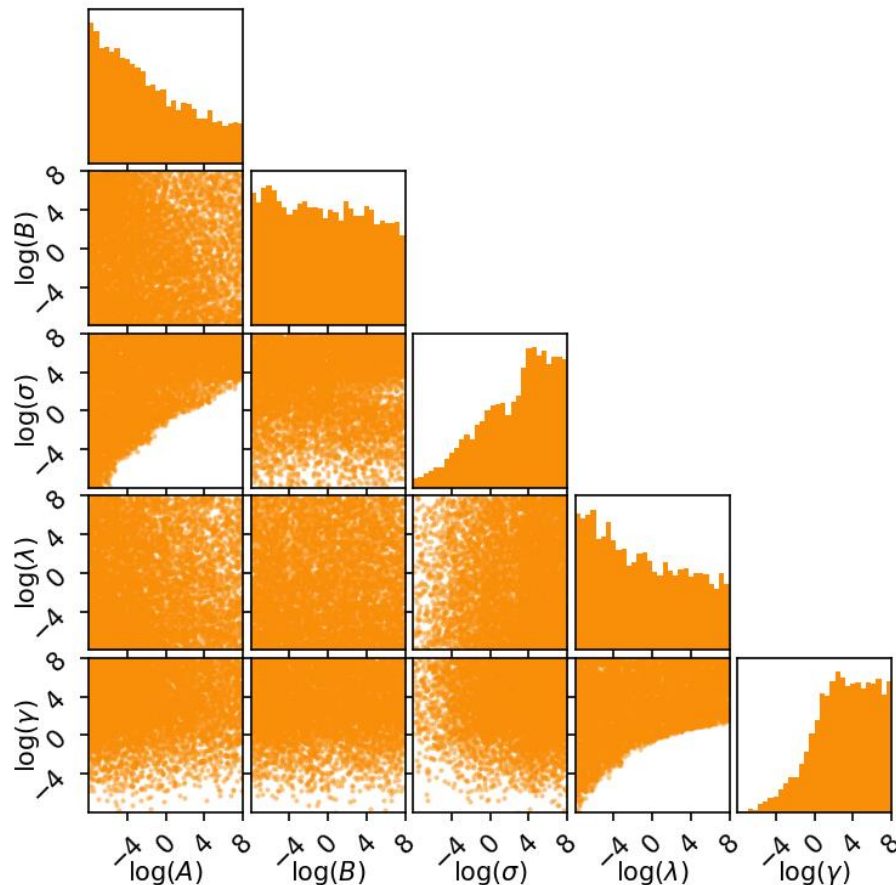
- Sampling temperature $T = 10^4$
- The distribution becomes wider as we increase the temperature.
- Parameter evaporation occurs: the walkers tend to run to sub-optimal parameter values [7, 8].
- Evaporated parameters are unconstrained by the data.
- Parameter evaporation becomes more apparent at higher temperatures.

[7] M. K. Transtrum, B. B. Machta, and J. P. Sethna, “Geometry of nonlinear least squares with applications to sloppy models and optimization,” *Phys. Rev. E*, vol. 83, no. 3, p. 036701, Mar. 2011, doi: [10.1103/PhysRevE.83.036701](https://doi.org/10.1103/PhysRevE.83.036701).
[8] R. Gutenkunst, “Sloppiness, Modeling, and Evolution in Biochemical Networks,” Cornell University, Ithaca, New York, 2007. Accessed: May 14, 2021. [Online]. Available: <https://ecommons.cornell.edu/handle/1813/8206>



Access to the
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scripts.

Parameter evaporation



- Sampling temperature $T = 10^5$
- The distribution becomes wider as we increase the temperature.
- Parameter evaporation occurs: the walkers tend to run to sub-optimal parameter values [7, 8].
- Evaporated parameters are unconstrained by the data.
- Parameter evaporation becomes more apparent at higher temperatures.
- We can use this result as a guide to collect more training data.

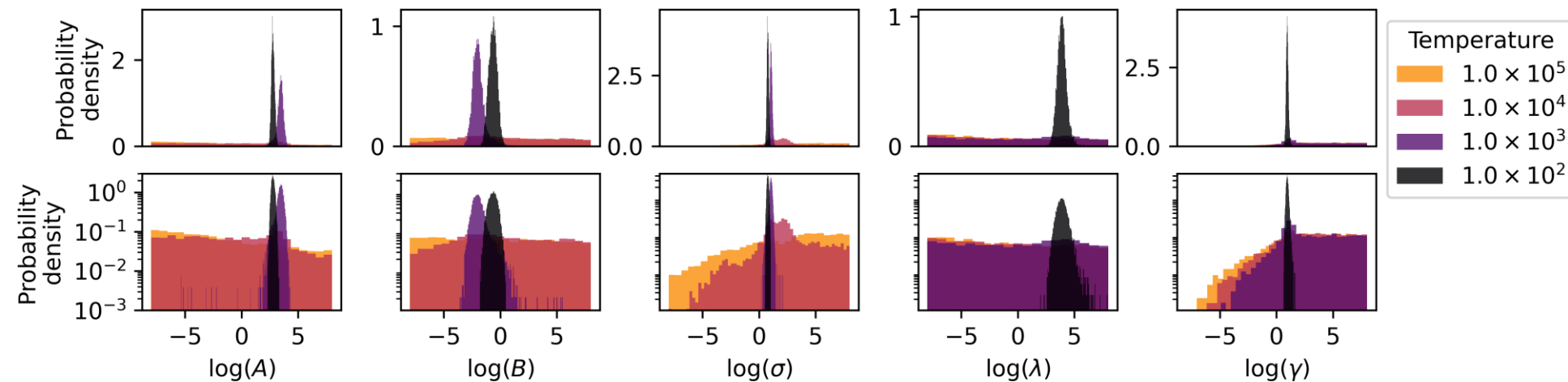
[7] M. K. Transtrum, B. B. Machta, and J. P. Sethna, “Geometry of nonlinear least squares with applications to sloppy models and optimization,” *Phys. Rev. E*, vol. 83, no. 3, p. 036701, Mar. 2011, doi: [10.1103/PhysRevE.83.036701](https://doi.org/10.1103/PhysRevE.83.036701).
[8] R. Gutenkunst, “Sloppiness, Modeling, and Evolution in Biochemical Networks,” Cornell University, Ithaca, New York, 2007. Accessed: May 14, 2021. [Online]. Available: <https://ecommons.cornell.edu/handle/1813/8206>



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example
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Comparison of the marginal distributions

Marginal distribution of the parameters at several sampling temperatures



- Compare the distributions at $T = 10^2$ and $T = 10^3$:
 - λ and γ evaporate.
 - The expectation value of A , B , and σ are shifted away from the best fit.
- How we should treat parameter evaporation is an open question.

Conclusion

- We enhance KLIFF with UQ framework.
- This implementation can facilitate more UQ studies and lead to more transparent and reproducible UQ analysis for IPs.
- We demonstrate it to study SW potential for silicon system.
- The result indicates parameter evaporation.
 - The data cannot constrain the evaporated parameters and future predictions.
 - The sampling result is highly dependent on the sampling temperature and prior.
- Suggestions:
 - Check for robustness of the result to several choice of prior.
 - Use the result to inform what other training data are needed.
- Future work:
 - Integrate other UQ methods.
 - Work on accelerating MCMC.



Access to the
example scripts.

Acknowledgement

- This work has been funded by the NSF under grant CMMT-1834332
- OpenKIM
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Access to the
example scripts.