

# The BAND Software Framework

Kyle Godbey

Slides with videos:

[https://docs.google.com/presentation/d/1XdE1vG8VEe97BCzEaW8fgwQkZimLL2gJcu\\_bDXwzL4XE/edit?usp=sharing](https://docs.google.com/presentation/d/1XdE1vG8VEe97BCzEaW8fgwQkZimLL2gJcu_bDXwzL4XE/edit?usp=sharing)



WANDA | 2024 WANDA Meeting  
Arlington, VA



# Introducing: BAND



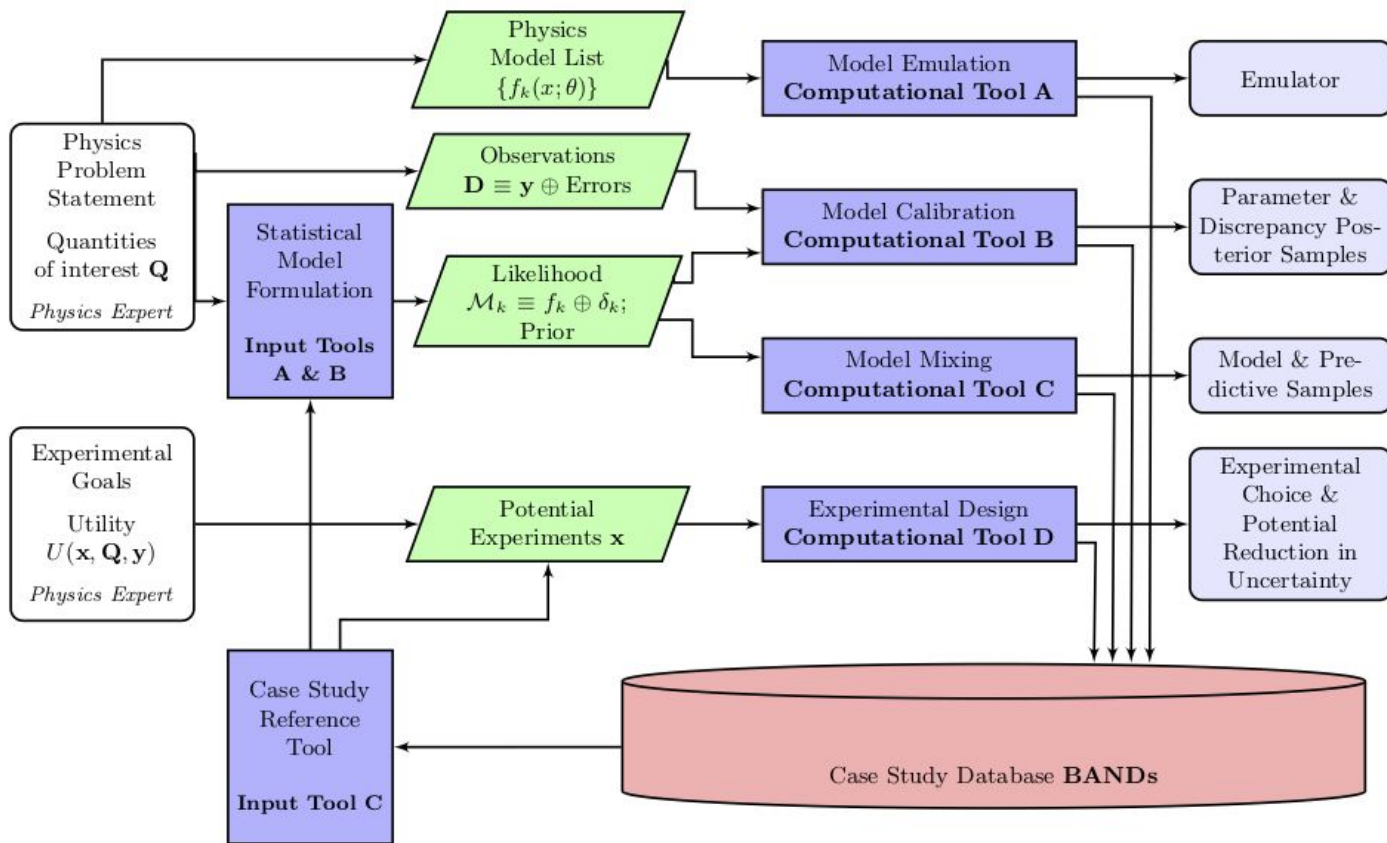
Supported by the NSF CSSI program under grant OAC-2004601



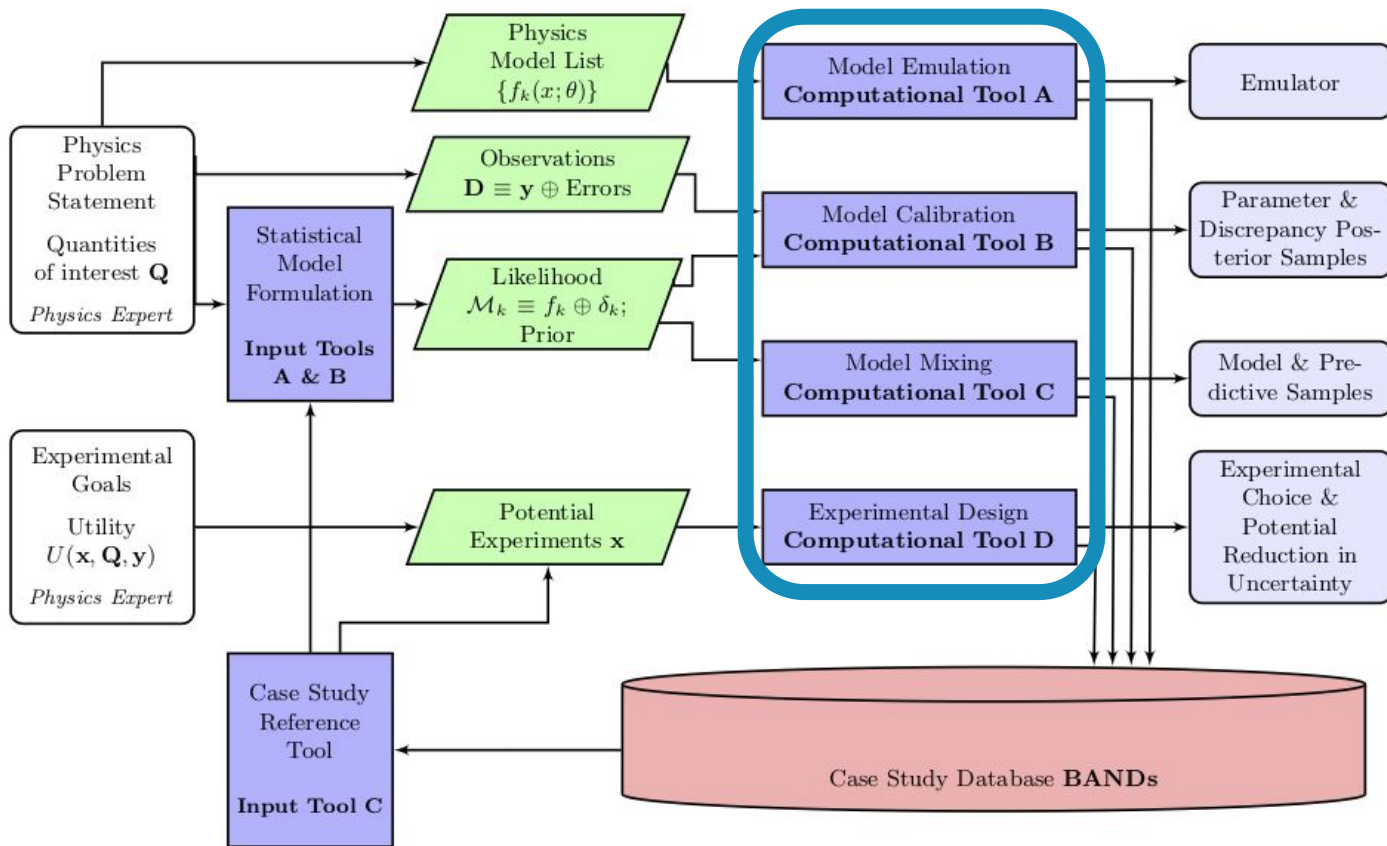
WANDA | 2024 WANDA Meeting  
Arlington, VA



# The Framework



# The Framework



# The Framework

Online:

<https://bandframework.github.io/>

On GitHub:

<https://github.com/bandframework/bandframework>

## Software

External code delivery will be from the [bandframework github repository](#)



### surmise

A Python package designed to provide a surrogate model interface for calibration, uncertainty quantification, and other tools.

*O. Surer, M. Plumlee, S.M. Wild, M. Y-H. Chan*  
[surmise Read the Docs](#)



### Taweret

A versatile Python package containing multiple model mixing techniques for a variety of use cases.

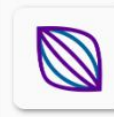
*K. Ingles, D. Liyanage, A. C. Semposki, J. C. Yannotty*  
[Taweret documentation](#)



### SAMBA

The SAndbox for Mixing using Bayesian Analysis, developed as a testing ground for multivariate model mixing on a toy model setup.

*A. C. Semposki, R. J. Furnstahl, D. R. Phillips*  
[SAMBA repository](#)



### ParMOO

ParMOO is a parallel multiobjective optimization solver that seeks to exploit simulation-based structure in objective and constraint functions.

*T.H. Chang, S.M. Wild, H. Dickinson*  
[parmoo Read the Docs](#)



### BMEX

The Bayesian Mass Explorer (BMEX) is a user-focused web application that provides a one-stop-shop for quantified theoretical model predictions of nuclear masses and related quantities.

*K. Godbey, L. Buskirk, P. Giuliani*  
[BMEX Web Application](#)



### ROSE

The Reduced Order Scattering Emulator (ROSE) is a Python package for building emulators using reduced basis methods for calculating nuclear scattering observables for user-defined interactions, including optical potentials.

*D. Odell, P. Giuliani, K. Godbey, K. Beyer, M. Y. Chan*  
[ROSE Github](#)



WANDA | 2024 WANDA Meeting  
Arlington, VA



# The Focus

ROSE – A scattering emulator for fast theoretical simulation without sacrificing accuracy

Use cases include model calibration, quick evaluation, efficient model mixing, experimental design, science gateways, and more (presumably)



## ROSE

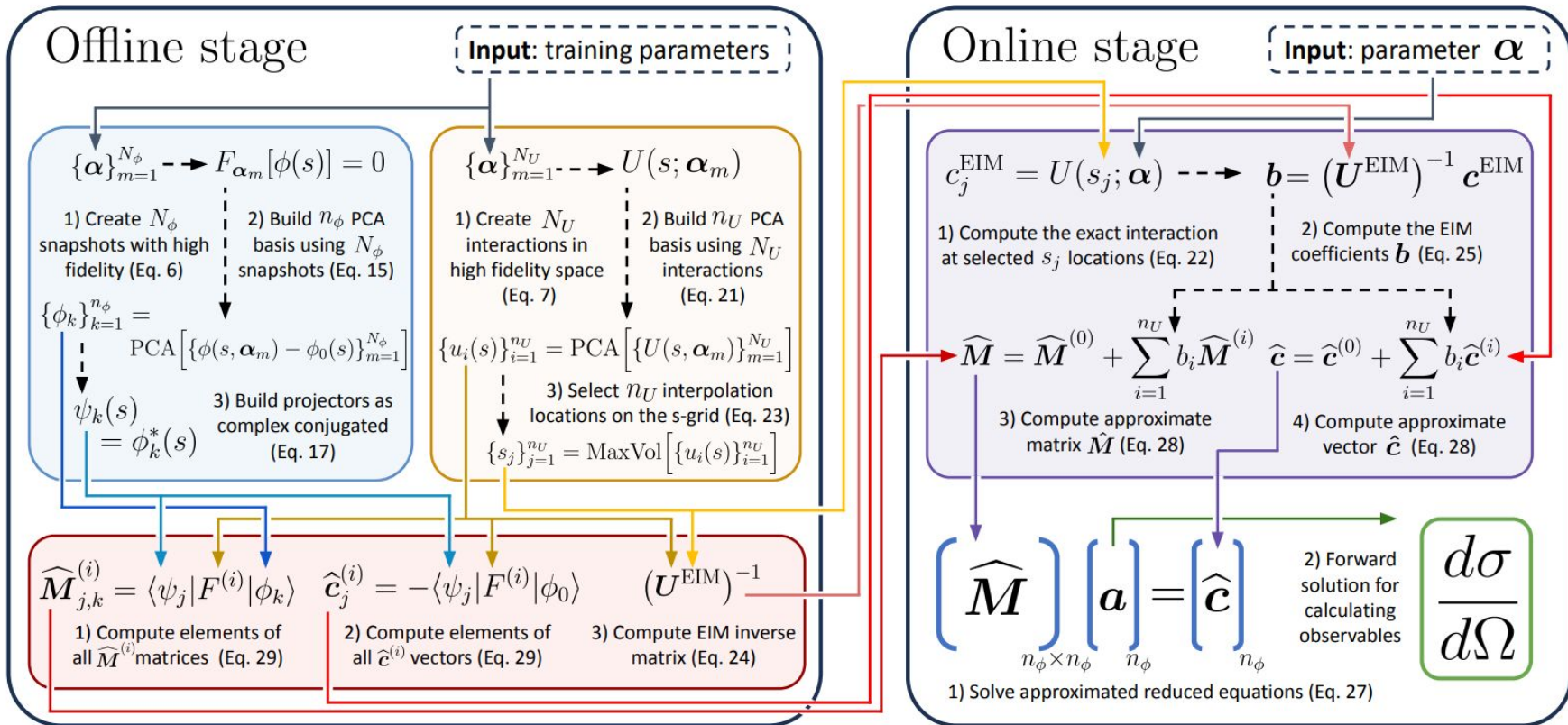
The Reduced Order Scattering Emulator (ROSE) is a Python package for building emulators using reduced basis methods for calculating nuclear scattering observables for user-defined interactions, including optical potentials.

*D. Odell, P. Giuliani, K. Godbey, K. Beyer, M. Y. Chan*

[ROSE Github](#)

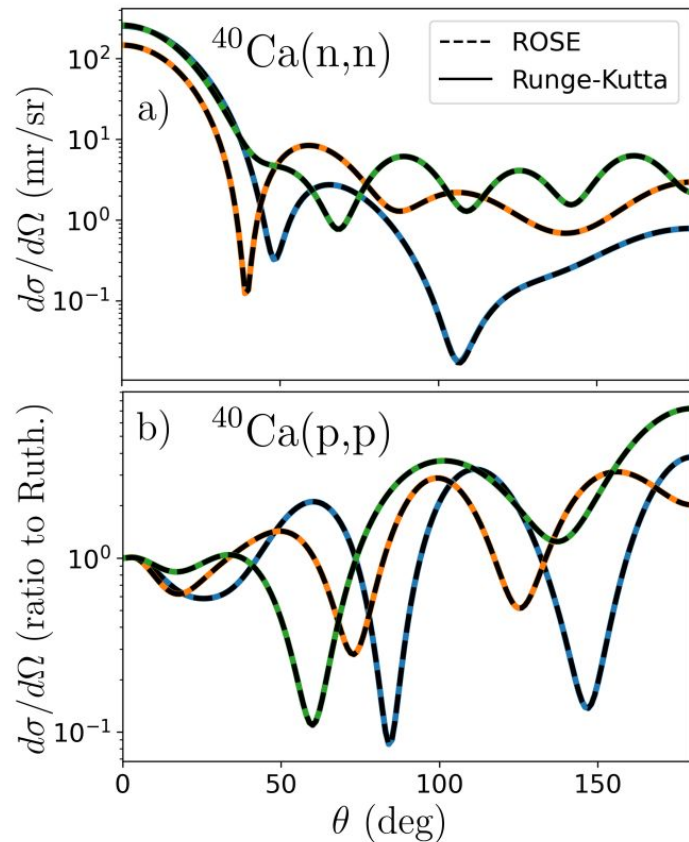


# The Focus



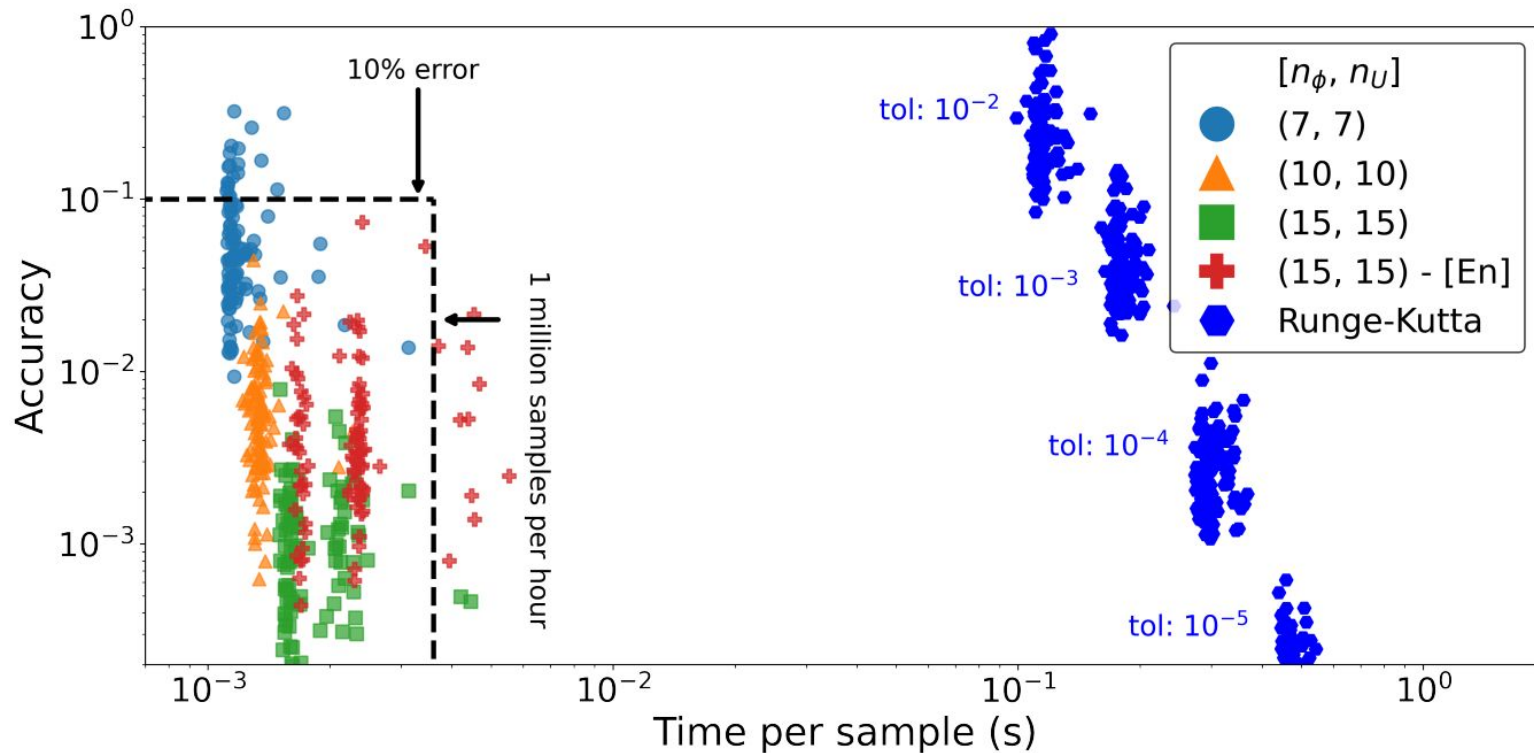
# The Goal

Supplementing high-fidelity simulations with 'almost as good' replacements



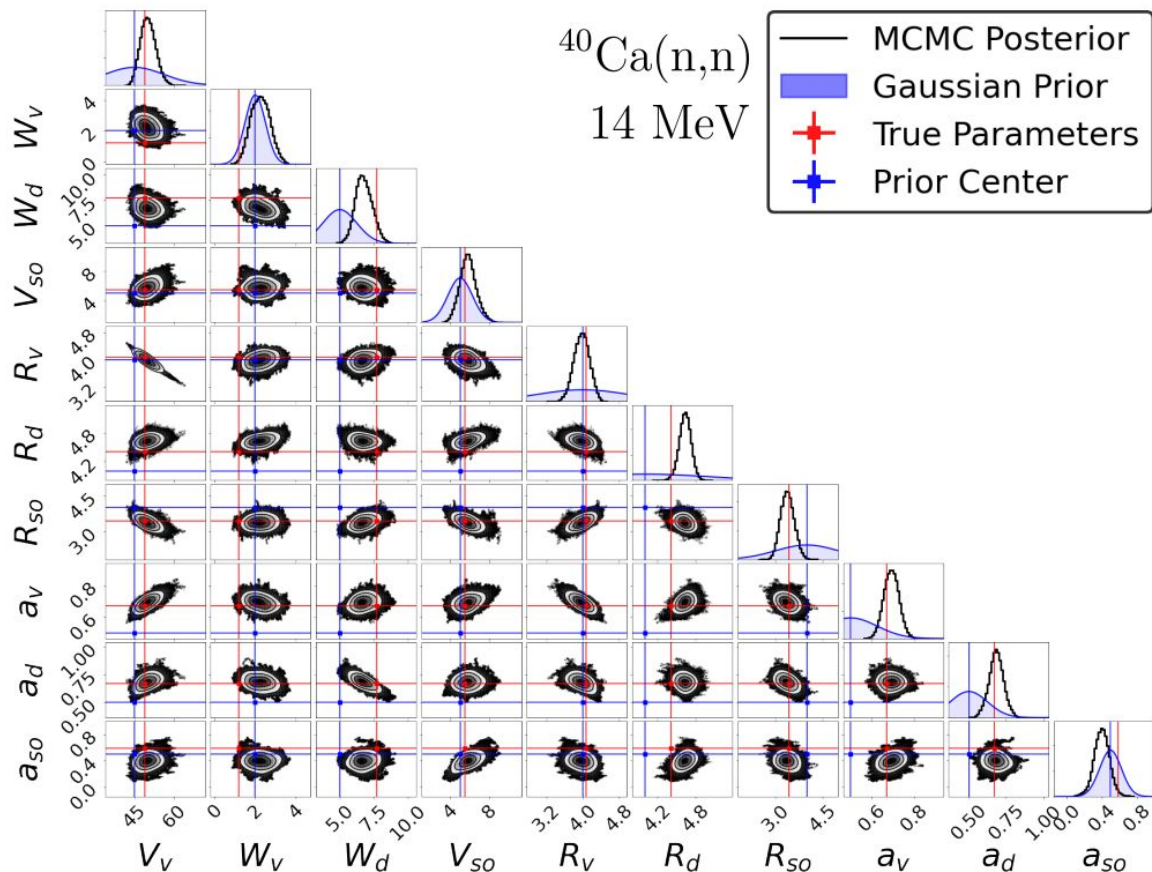


# The Goal



# The Outcome

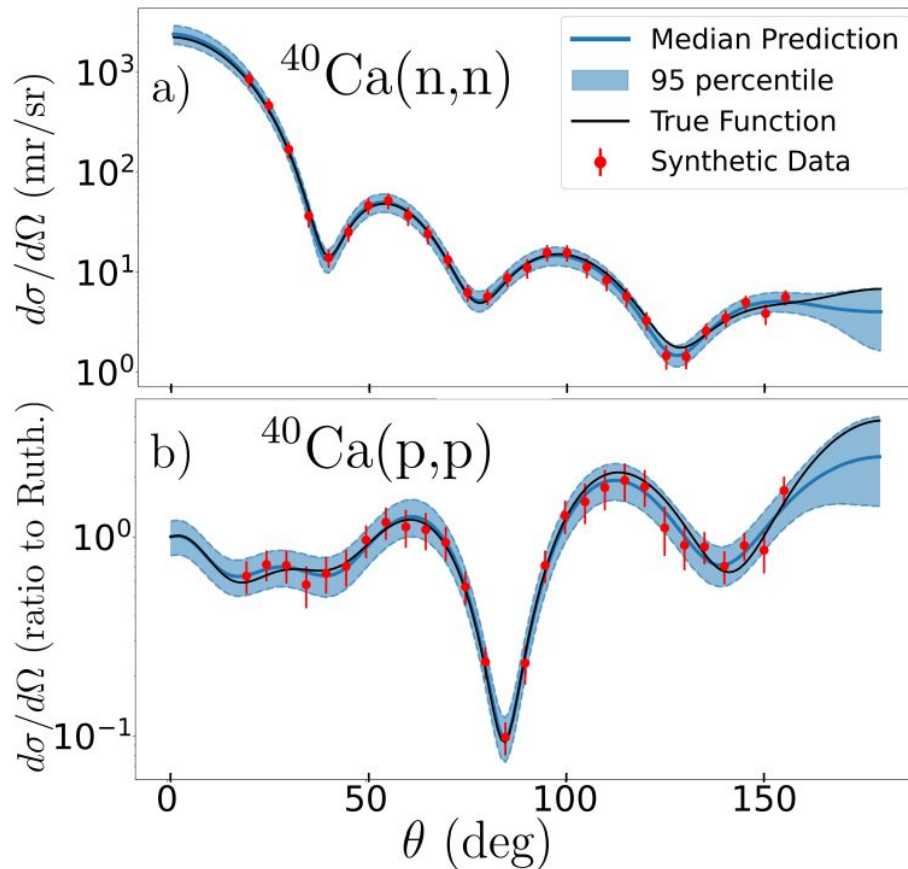
Principled Bayesian calibration is more accessible than ever before



# The Outcome

The point is to get predictions, uncertainties, and covariances quickly

Integration with model mixing tools will help consider the wisdom of many models



# Outlook

Continued feature development in both physics and emulation

Coupled-channels emulation

Black-box dimensionality reduction

Expanded integration within and without the BAND framework

Bayesian model mixing

Explore novel applications and user interaction modality

Cloud-enabled backends

Web applications



1 2 3 +

Dimension:  
1D Chains

1D Chain:  
Isotonic Chain

Select Quantity:  
Single-Proton Energy Splitting

$$\Delta E_p(N,Z) = S_p(N,Z) - S_p(N,Z+2)$$

- N=60 | AME2020
- N=60 | UNEDF1
- N=60 | SV
- N=60 | HFB24
- N=60 | FRDM12**
- +

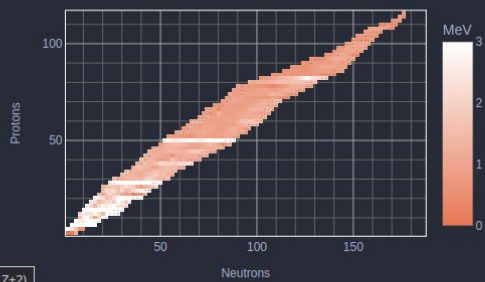
Neutrons:  
60

Select Dataset:  
FRDM12

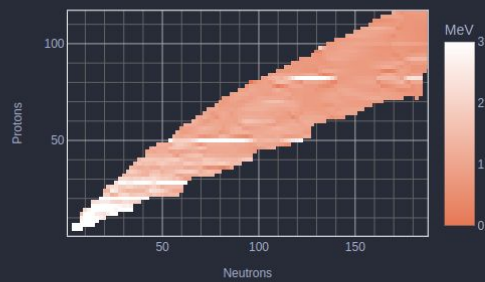
Wigner Adjustment:  
None

DELETE SERIES

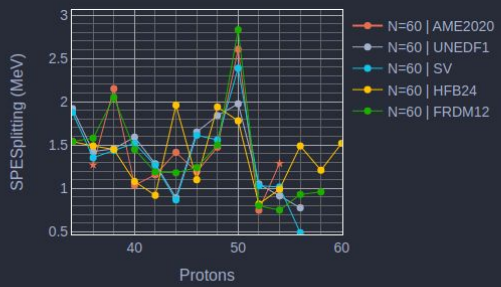
Single Proton Energy Splitting | AME2020



Single Proton Energy Splitting | UNEDF1



Isotonic Chain



Share View

EXPORT PUB. PDFS

LINK VIEWS  
 1  2  3

Even-Even Nuclei

RESCALE COLORBAR

RESET PAGE



WANDA | 2024 WANDA Meeting  
Arlington, VA

<https://bmex.dev>

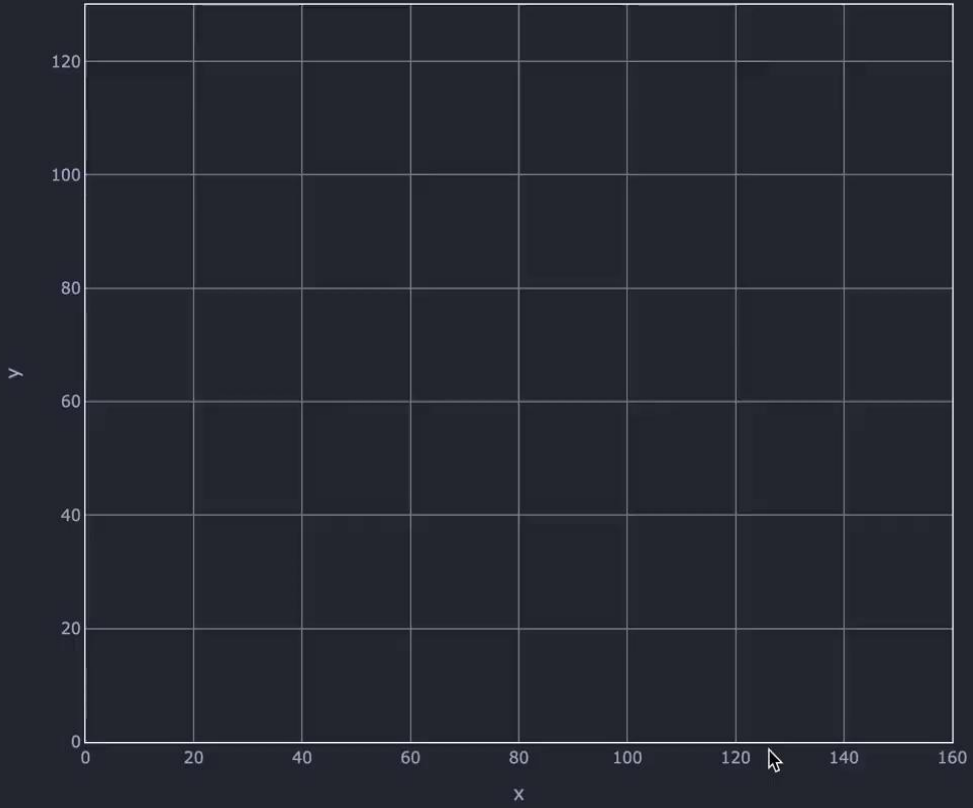


**BAND**  
Bayesian Analysis of Nuclear Dynamics

FIT

CLEAR

	x	y	$\bar{y}$
x			



# Our Request

Continued feature development in both physics and emulation

Expanded integration within and without the BAND framework

Explore novel applications and user interaction modality

What features are needed? What models are highest impact?

What does harmonious integration look like?

What **tools** have the highest potential for impact across the pipeline?



# Conclusion

We want to build a broadly useful, open-source framework that meets the needs of the community

To that end we welcome suggestions, feedback, critiques, requests, code, etc.

