Advancing the Theory of Nuclear Data Evaluations

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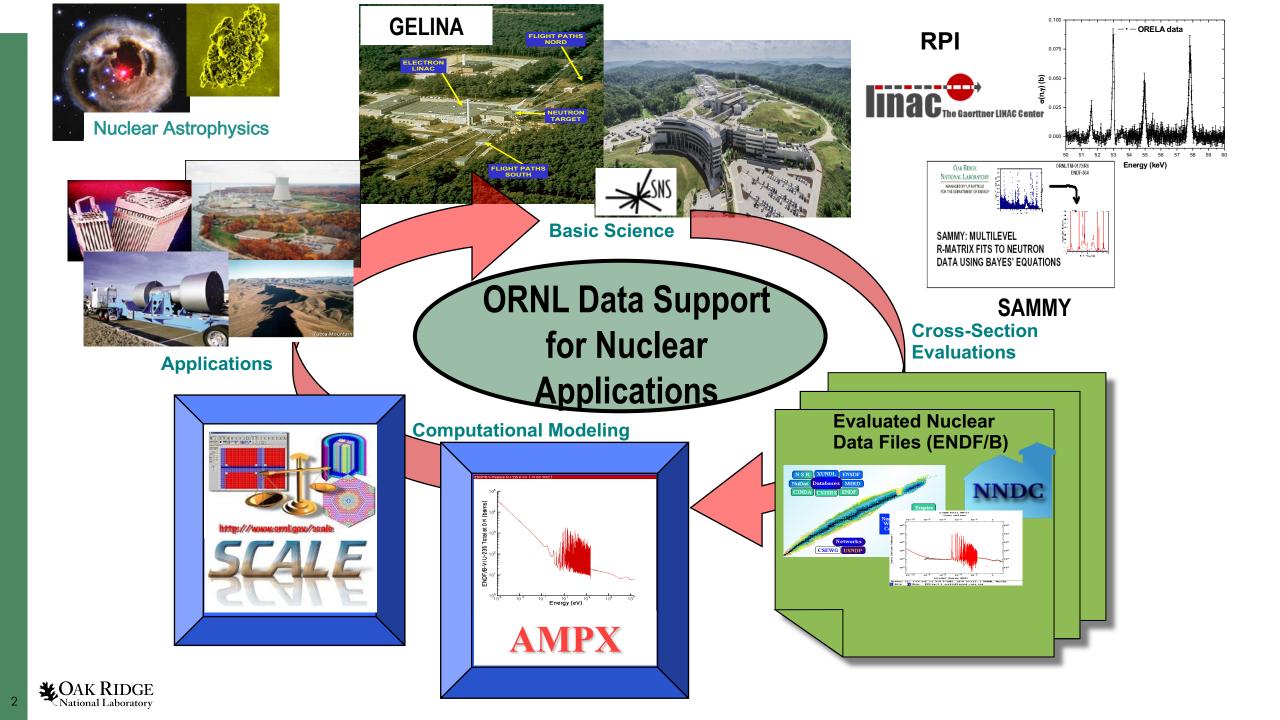
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WONDER-2023



International Workshop On Nuclear Data Evaluation for Reactor Applications

Aix-en-Provence, France, June 5-9, 2023.



Nuclear Criticality Safety Program (NCSP) Motivation:

- NCSP Mission & Vision, 5-year plans:
 - <u>https://ncsp.llnl.gov/sites/ncsp/files/2021-04/ncsp_mission_vision.pdf</u>
 - <u>https://ncsp.llnl.gov/program-management/ncsp-five-year-execution-plan</u>
 - Uncertainty Quantification
 - Differential Nuclear Data (resolved resonance region (RRR) by SAMMY)
 - Integral Benchmark Experiments
- II. SAMMY Modernization
 - R-matrix formalism



Part I. Theory of evaluated nuclear data (ND) uncertainties

Common assumptions used in Bayesian ND evaluations:

1. Linearity:

- all models are linear

2. Normality:

- all probability distribution functions are normal, i.e., Gaussian

3. Perfection:

a) The model provides a perfect description of the measured datab) The data are perfect and complete (*including* the covariances)



These assumptions can now be selectively removed or enforced

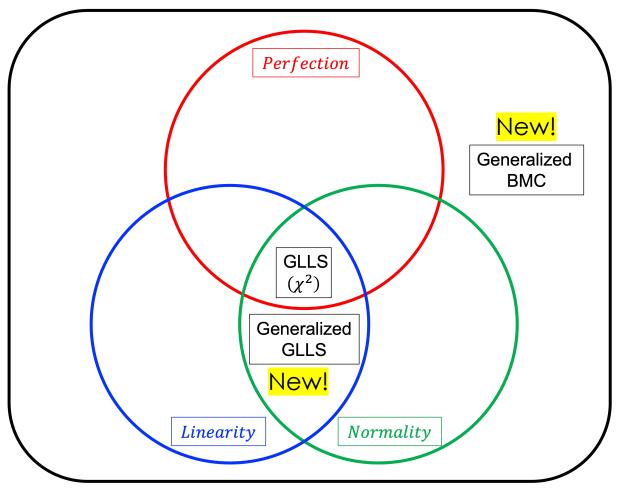
• Linearity and Normality:

 Removed by, e.g., Metropolis Hastings Monte Carlo (MHMC); known as Bayesian Monte Carlo (BMC)

• Perfection:

 Evaluator can remove this assumption by specifying posterior expectation values and covariance of *deviations* between the model and data

GLLS = Generalized Linear Least Squares BMC = Bayesian Monte Carlo

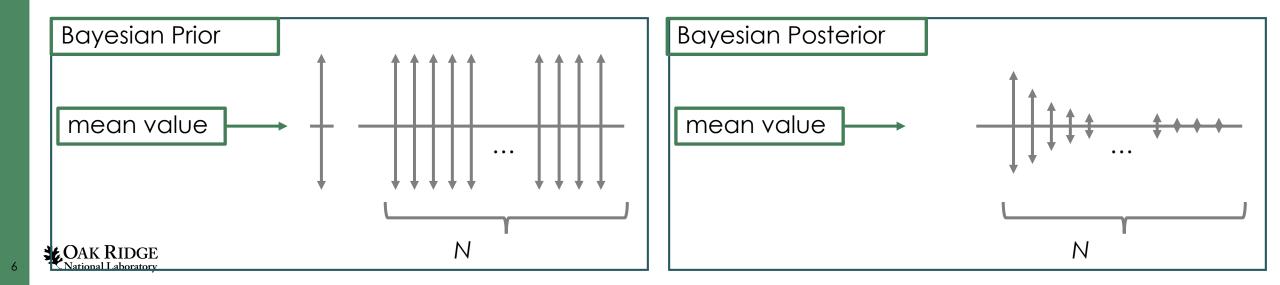


Note: everything else being the same, consistency with Bayes' theorem in this framework improves the likelihood of success. Reliability of a given evaluation still depends on evaluator's skill/expertise.



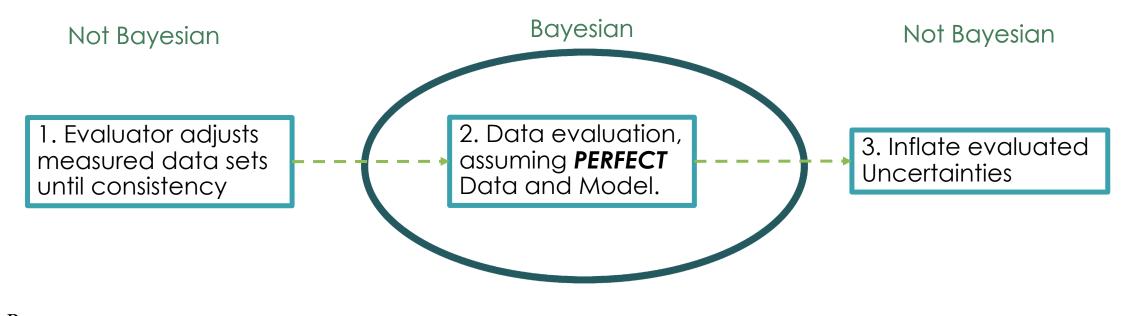
Illustrating a mechanism behind **small** evaluated uncertainties

- Example: suppose a large number ("N") of identical measurements
 - Suppose measurements are *identical* in value as well as uncertainty
 - This enables focus on evaluated covariance/uncertainty since the mean values are unaffected
 - Suppose that the correlation among measurements is set to 0
 - Bayes' theorem then yields <u>uncertainty</u> \rightarrow 0 as N \rightarrow infinity (illustrated below)
 - Unrealistically small evaluated uncertainties are rectified by inflating them until reasonable
 - The uncertainties are underestimated less apparently for any value of N
- Our Bayesian framework provides tools to address this problem.



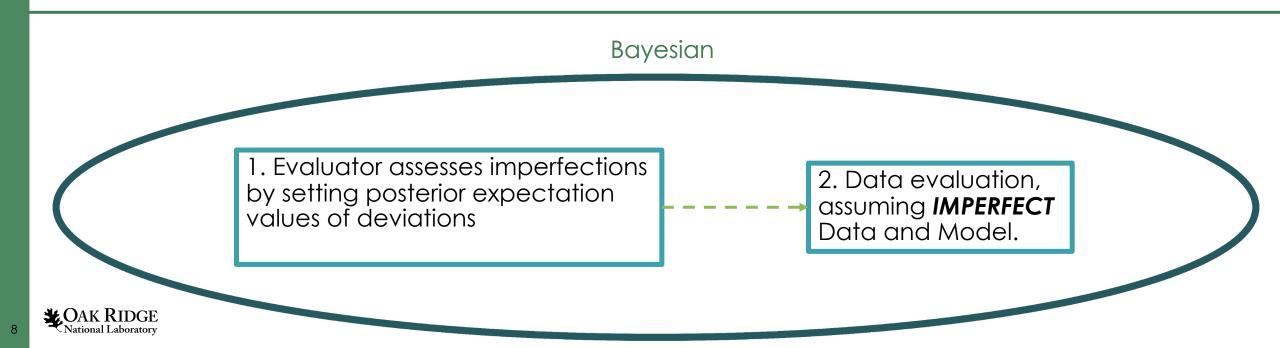
Conventional evaluation workflow is not completely Bayesian:

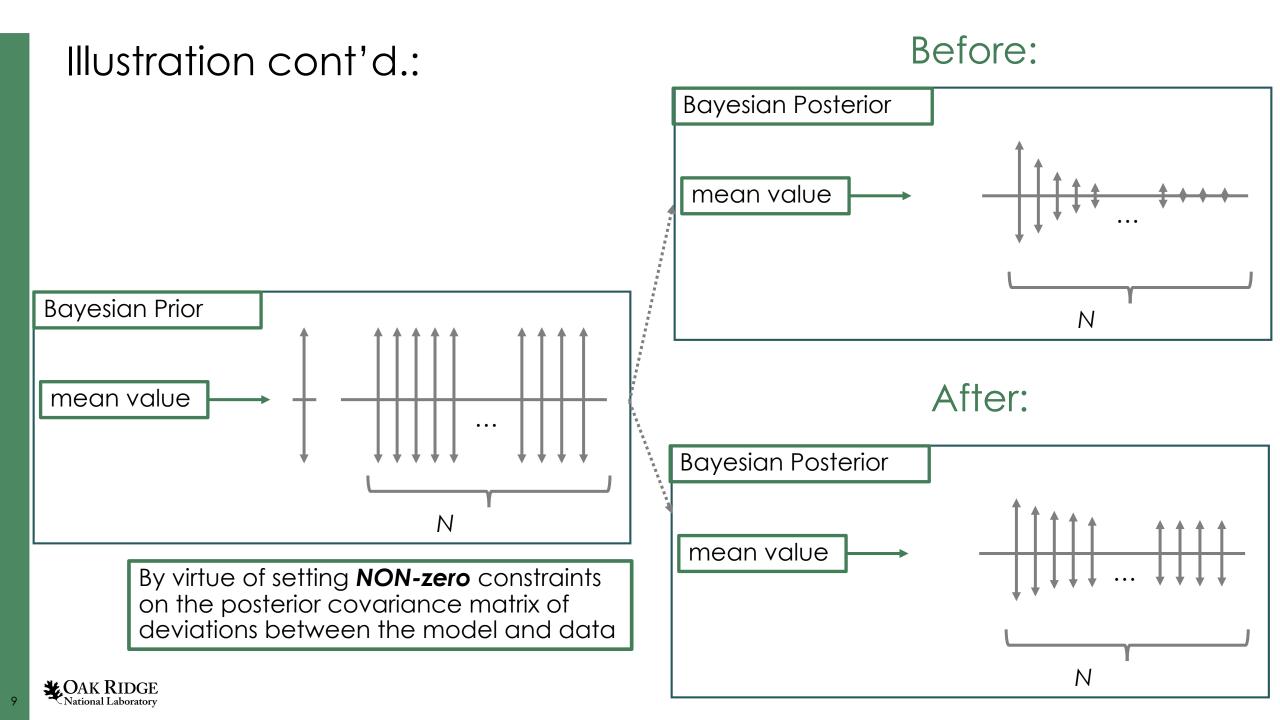
- 1. Evaluator uses expert judgment to align measured data sets before the evaluation
- 2. Bayesian evaluation is performed (implicitly) assuming perfect data and model
- 3. Unrealistically small uncertainties are inflated manually afterwards



New framework enables completely Bayesian evaluations:

- 1. Evaluator *estimates* the effect of *imperfections* by *setting* Bayesian *posterior* expectation values of *deviations* as well as their *covariances*
 - **Deviation** is defined as a **difference** between the evaluated data and model
 - Evaluators' expert judgment (or intuition) now formally recognized within Bayes' theorem!
- 2. Bayesian evaluation is now determined by the *deviations* defined in 1.
 - No need to manually inflate evaluated uncertainties as in Step 3. previously





Assumptions or approximations used with Bayes' Theorem

- 1. The model and the prior PDF of data are assumed to be perfect
- 2. The model is approximated by its 1st order (*linear*) expansion
- 3. Prior and posterior PDFs are approximated by <u>normal</u> PDFs.
- We recognize that 1. is equivalent to constraining the posterior expectation values of $\,\delta$, and of its covariance matrix, to 0:

$$\langle \delta \rangle' = 0$$
 $\Delta' \equiv \langle (\delta - \langle \delta \rangle')(\delta - \langle \delta \rangle')^{\mathsf{T}} \rangle' = \mathbb{O}$ $\delta \equiv \delta(z, T(P)) \equiv T(P) - D$ $z \equiv \begin{pmatrix} P \\ D \end{pmatrix}$

(-)

• We remove 1. by letting evaluator choose values of $\langle\delta
angle'$ and Δ'

• 2. and 3. can be removed by Metropolis-Hastings Monte Carlo

Overview of approximations used by ORNL codes

Algorithm / Code	< δ>'	Δ'	Prior/Post PDF	Cost Function	Minimization
GLLS / SAMMY	0	0	Normal/Normal	Chi ²	Linear (iterative)
GLLS /TSURFER	0	0	Normal/Normal	Chi2	Linear (single step)
Generalized GLLS	Any	Any	Normal/Normal	Generlized Chi ²	Linear
ВМС	Any	Any	Any/any	Posterior PDF	MHMC

• Bayes' theorem with arbitrary constraints:

 $p'(z|\beta\gamma) = p'(z|\gamma\beta) = N'\mathcal{L}(\beta|z\gamma) \times p(z|\gamma)$

 $\mathcal{L}(\beta|z,\gamma) \leftarrow \mathcal{L}(\langle \delta \rangle', \mathbf{\Delta}'|z,\gamma) = e^{-\frac{1}{2}(\delta-\lambda)^{\intercal} \mathbf{\Lambda}^{-1}(\delta-\lambda)}$

 $\beta \leftarrow \{ \text{any constraints on posteriors imposed by evaluator} \}, \\ \gamma \leftarrow \{ \text{any parameters needed to define the prior PDF}, p(z|\gamma) \}$



Generalized GLLS Framework in TSURFER notation

<u>Using TSURFER-like notation:</u>

$$\begin{split} z &\equiv \begin{pmatrix} \alpha \\ m \end{pmatrix} \\ \tilde{C}_{ZZ} &= \begin{pmatrix} \tilde{C}_{\alpha\alpha} & \tilde{C}_{\alpham} \\ \tilde{C}_{m\alpha} & \tilde{C}_{mm} \end{pmatrix} \\ \tilde{S} &\equiv \nabla_Z \tilde{d}^T = \begin{pmatrix} \nabla_\alpha \\ \nabla_m \end{pmatrix} (k(\alpha) - m)^T \\ &= \begin{pmatrix} \nabla_\alpha k(\alpha)^T \\ -\nabla_m m^T \end{pmatrix} = \begin{pmatrix} \tilde{S}^T_{k\alpha} \\ -\mathbb{I} \end{pmatrix} , \text{ (I = identity matrix)} \\ \tilde{C}_{dd} &= \tilde{S} \tilde{C}_{ZZ} \tilde{S} \end{split}$$

$$\alpha$$
 = nuclear data
 m = measured response
 $d = k(\alpha) - m$



$$\tilde{C}_{z'z'}^{-1} = \tilde{S} \left[\tilde{C}_{d'd'}^{-1} - \tilde{C}_{dd}^{-1} \right] \tilde{S}^T + \tilde{C}_{zz}^{-1}$$
$$z' = z + \tilde{C}_{zz} \tilde{S} \tilde{C}_{dd}^{-1} (\tilde{d}' - \tilde{d})$$

Note: SAMMY uses the GLLS *iteratively*, by re-computing sensitivities in each iteration, as a way to approximately account for non-linearity of the R-matrix formalism.



Benefits of a generalized form of the Bayes' Theorem (BT):

- It could improve evaluations of any data, separately or jointly
 - differential cross sections (SAMMY),
 - integral benchmarks (TSURFER/SAMPLER), ...
- Enables <u>Bayesian Monte Carlo</u> evaluation of <u>large data sets</u>
 - Useful for, e.g., TSL evaluations of SNS data, RRR, ...
- Enables <u>Bayesian</u> evaluation of:
 - *inconsistent* data sets, and/or
 - <u>defective</u> model
- Conventional BT is recovered when imperfections made to vanish
 A seamless connection to the BT in SAMMY/TSURFER/SAMPLER
- API implementation in the SCALE code system
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For more information on uncertainty evaluation consider:

- ORNL/TM-2022/2448 Technical Report
 - Brown, Jesse M., Arbanas, Goran, Wiarda, Dorothea, and Holcomb, Andrew, "<u>Bayesian Optimization Framework for Imperfect Data or</u> <u>Models</u>". United States: 2022. <u>https://doi.org/10.2172/1874643</u>.
- ANS Winter Meeting 2022:
 - Jesse M. Brown, Goran Arbanas, Hany Abdel-Khalik, Ugur Mertyurek, William B. Marshall, William A. Wieselquist, "<u>Generalized Bayesian</u> <u>Framework for Evaluation of Integral Benchmark Experiments</u>", Transactions, Volume 125, Number 1, December 2021, Pages 691-694.
- ND 2022:
 - Jesse M. Brown, Goran Arbanas, Andrew Holcomb, Dorothea Wiarda, "<u>Bayesian Evaluation Framework for Imperfect ²³³U Data and Models",</u> Proceedings of the 15. Int.'I Conference on Nuclear Data for Science and Technology (ND2022), Sacramento, California, 21-29 Jul 2022.



Part II. R-matrix formalism: Background and Outline

- Classes of nuclear reaction processes, by increasing complexity:
 - 1. Direct: direct channel coupling, direct capture, ...
 - 2. Doorway: Giant Dipole Resonance, Isobar Analogue Resonance, 2p1h, ...
 - 3. Compound: narrow resonances of the Resolved Resonance Region
- Only the compound kind parameterized in the extant R-matrix
- New *R*-matrix parameterization of direct and doorway processes
 - **Direct**: parameterized exactly, using an idea suggested by Wigner [8]
 - **Doorway**: parameterized analogously to a corresponding K-matrix way [1]
- *R*-matrix formalism for evaluations rediagonalizes parameterization of any many body Hamiltonian, obviating the need for doorways:
 – Elimination of capture channels yields complex resonance energies, widths



Non-resonant (or external) direct reactions in *R*-matrix

• Direct channel coupling is introduced by mixing among channels of the incoming, I, and the outgoing, O, formerly diagonal matrices of wave functions: $\Psi = Iy + Ox$

$$x = -Uy$$

 $y \leftarrow \mathcal{M} y$ $x \leftarrow \mathcal{M}^* x$

where \mathcal{M} is a unitary matrix parameterizing direct reactions. A corresponding scattering matrix U is then found to be:

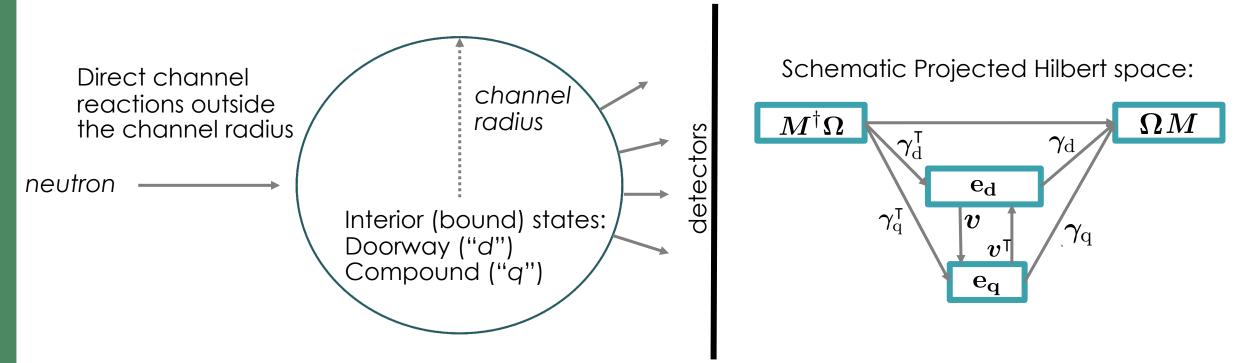
$$U_{\mathcal{M}} = \mathcal{M}^{\mathsf{T}} U \mathcal{M}, \qquad \qquad \mathcal{M}^{-1} = \mathcal{M}^{\dagger} \equiv (\mathcal{M}^{*})^{\mathsf{T}}$$

Suggested by Wigner in a paper on generalized Euler angles [8].

 Its analogue in the T-matrix by H. Feshbach in Eq. (III.2.26) of [7] is an example of similarities among R-, T-, K-matrix formalisms.
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Introducing doorway states into R-matrix formalism

- Generalized R-matrix parameterization of doorway states ("d"):
 - Comparison of formal expressions for doorway state K-matrix (derived by Feshbach's projection operator formalism [1,2]) to the Brune's alternative R-matrix [9] has helped uncover an analogous R-matrix parameterization (using a simple derivation shown on the next slide):



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(This is another example of leveraging similarities among the R-, T-, K-matrix formalisms.)

Doorway state parameterization in conventional R-matrix [3]

- Q^{-1} : a 2x2 block matrix of expectation values of projected H-E1– The projected R-matrix is simpler than the projected T-,K-matrix in [1, 2]
- $m{v}\equiv \langle d|m{H}|q
 angle$ is a doorway-compound coupling parameter matrix
- Doorway states are far fewer and wider than compound states
 - The new doorway R-matrix parameters are: $m{e}_{
 m d}$, $m{\gamma}_{
 m d}$, and $m{v}$
 - Extensible to multistep reactions by adding new subspaces into $\langle \lambda |$
 - Single-particle resonances for direct processes, 3p2h for "hallway" states, ..., CN.

National Laboratory Note: Re-diagonalization of the R-matrix returns to the original form of the R-matrix formalism!

Elimination of capture channels yields an effective R-matrix:

Eqs. (X.1.7-9a) of Lane and Thomas (LT) express a reduced *R*-matrix as

$$\mathcal{R} = \boldsymbol{\gamma}_r^{\mathsf{T}} \boldsymbol{A}^{-1} \boldsymbol{\gamma}_r \tag{1}$$

$$= \boldsymbol{\gamma}_r^{\mathsf{T}} [\boldsymbol{e} - \boldsymbol{\xi} - E \boldsymbol{1}]^{-1} \boldsymbol{\gamma}_r, \qquad (2)$$

where¹

$$\boldsymbol{\xi} \equiv \boldsymbol{\gamma}_e \boldsymbol{L}_e' \boldsymbol{\gamma}_e^{\mathsf{T}},\tag{3}$$

is a symmetric matrix, and where $L'_e = L^0_e \equiv L_e - B_e$ for $R^0_{ee} = 0$, as in Section X.1 of LT. In order to show how an *effective* set of reduced *R*-matrix parameters may become complex-valued, a singular value decomposition (SVD) of a symmetric² matrix $e - \xi$ is assumed to exist, *i.e.*,

$$\boldsymbol{e} - \boldsymbol{\xi} = \boldsymbol{v}\boldsymbol{\epsilon}\boldsymbol{v}^{\mathsf{T}},\tag{4}$$

where $vv^{\dagger} = 1$ and ϵ is diagonal³. Utilizing this SVD in Eq. (2) yields an equivalent expression for the reduced *R*-matrix of LT,

$$\mathcal{R} = \gamma_r^{\mathsf{T}} [\boldsymbol{v} \boldsymbol{\epsilon} \boldsymbol{v}^{\mathsf{T}} - E \mathbf{1}]^{-1} \boldsymbol{\gamma}_r$$
 (5)

$$= \boldsymbol{\gamma}_r^{\mathsf{T}} [\boldsymbol{v}(\boldsymbol{\epsilon} - E\mathbf{1})\boldsymbol{v}^{\mathsf{T}}]^{-1} \boldsymbol{\gamma}_r \tag{6}$$

$$= \boldsymbol{\gamma}_r^{\mathsf{T}} (\boldsymbol{v}^{\mathsf{T}})^{-1} [\boldsymbol{\epsilon} - E \mathbf{1}]^{-1} \boldsymbol{v}^{-1} \boldsymbol{\gamma}_r$$
(7)

$$= (\boldsymbol{\gamma}_r^{\mathsf{T}} \boldsymbol{v}) [\boldsymbol{\epsilon} - E \mathbf{1}]^{-1} (\boldsymbol{v}^{\mathsf{T}} \boldsymbol{\gamma}_r)$$
(8)

$$= \boldsymbol{g}_r^{\mathsf{T}} [\boldsymbol{\epsilon} - E \boldsymbol{1}]^{-1} \boldsymbol{g}_r \tag{9}$$

$$\boldsymbol{\gamma}_r^{\mathsf{T}}(\boldsymbol{v}^{\mathsf{T}})^{-1} [\boldsymbol{\epsilon} - E \mathbf{1}]^{-1} \boldsymbol{v}^{-1} \boldsymbol{\gamma}_r \tag{7}$$

$$= (\boldsymbol{\gamma}_r^{\mathsf{T}} \boldsymbol{v}) [\boldsymbol{\epsilon} - E \mathbf{1}]^{-1} (\boldsymbol{v}^{\mathsf{T}} \boldsymbol{\gamma}_r)$$
(8)

$$= \boldsymbol{g}_r^{\mathsf{T}} [\boldsymbol{\epsilon} - E \boldsymbol{1}]^{-1} \boldsymbol{g}_r \tag{9}$$

of our WONDER 2023 proceedings. Note: the form of the

Taken from a draft

the R-matrix formalism remains the same; the effective parameters become complex-valued.

Unlike the conventional Reich-Moore approx. [5], the total capture cross section is exact, as in [6].

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$$oldsymbol{g}_r\equivoldsymbol{v}^{\intercal}oldsymbol{\gamma}_r;$$

 (ϵ, q_r) may be viewed as an *effective* reduced *R*-matrix parameter set which could be optimized independently of the formal reduced parameter set (e, γ_r) ;

(10)

where

¹Labels r and e refer to "retained" and "eliminated" subset of channels, respectively. ²The same form of SVD holds for *complex*-valued symmetric matrix, for which v and ϵ may be complex-valued.

³Note that a symmetric matrix e need not be diagonal.

Outlook:

- Other potential applications of this *R*-matrix parameterization:
 - Refining the assumptions of the Random Matrix Theory in nuclear physics by investigating the effect of direct and doorway states on statistical distributions of resonance widths (Porter-Thomas), energy spacing (Wigner)
 - A new way for modeling *multistep* (3p2h "hallway", ...) nuclear reactions
- Look for other ways to exploit similarities among various formalisms
- To be implemented into SAMMY [4] after code modernization
 - See the presentation by Doro Wiarda (ORNL) on SAMMY modernization
 - <u>https://ncsp.llnl.gov/sites/ncsp/files/2023-03/60_sammy_tpr_2023.pdf</u>



SAMMY and all dependencies are freely available

SAMMY () https://code.ornl.gov/RNSD/SAMMY



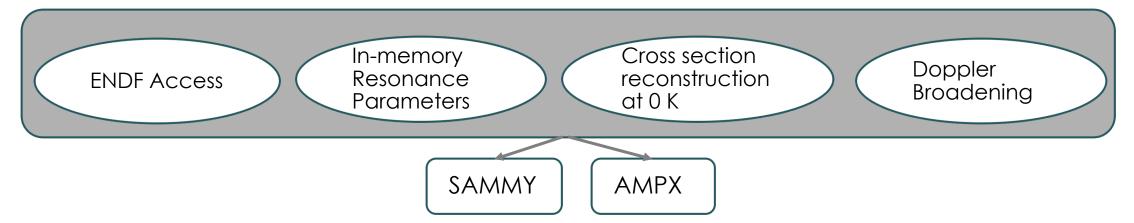
- The build system has been updated to automatically retrieve and build all dependencies.
- The README has been updated.



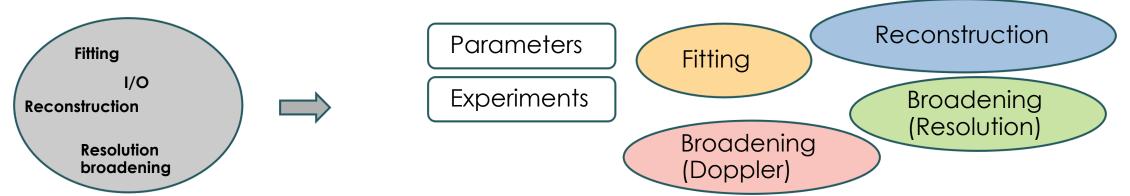
SAMMY Modernization and Maintenance goals

We plan to modernize as follows:

• Write code once and reuse



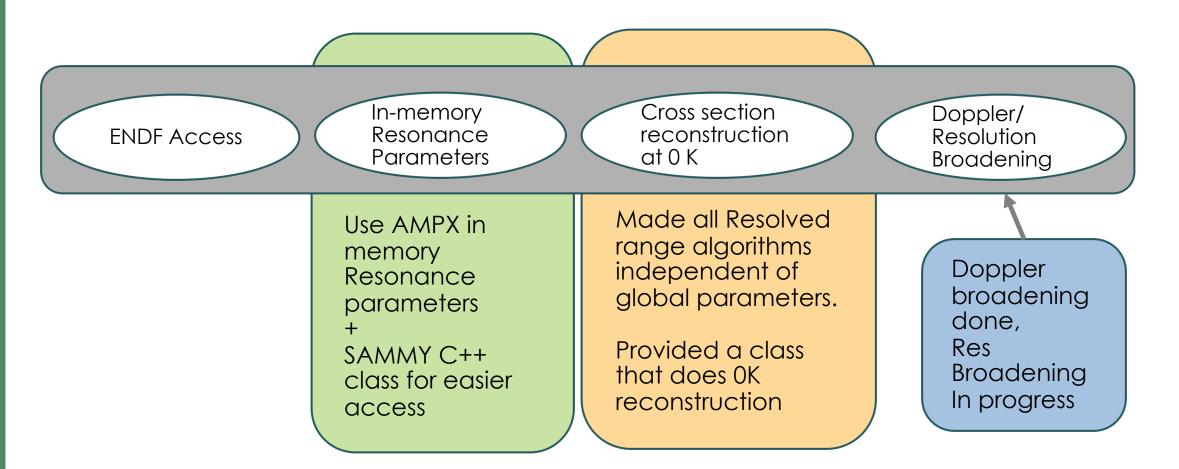
• Transform SAMMY into a modular code, with independent modules with clear interfaces.



• Add new features, which is now easier as only the desired module needs to be changed.



SAMMY Modernization Accomplishments



These modernizations will make adding new R-Matrix features easier as changes will be localized to the relevant part of the code as no global parameters are used. In addition, this will allow more parallelization of the code in the future.



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