

Cooling, Masses and Emulators

Sudhanva Lalit 10 December, 2024 INT, Washington









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Accreting Neutron Stars Observables

Accretion Outburst







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Accretion Outburst





Quiescence







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Accretion Outburst



Quiescence





KS 1731-260 observed with Chandra X-ray Satellite



Mass Model	Urca Pair
BCPM	⁵⁹ Ca ↔ ⁵⁹ Sc



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DZ31	⁵⁹ Sc ↔ ⁵⁹ Ti
FRDM-2012	⁵⁹ Sc ↔ ⁵⁹ Ti
WS4RBF	⁵⁹ Sc ↔ ⁵⁹ Ti



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DZ31	⁵⁹ Sc ↔ ⁵⁹ Ti
FRDM-2012	⁵⁹ Sc ↔ ⁵⁹ Ti
WS4RBF	⁵⁹ Sc ↔ ⁵⁹ Ti
D1M	⁵⁷ Sc ↔ ⁵⁷ Ti
HFB-24	${}^{57}\text{Sc} \leftrightarrow {}^{57}\text{Ti}$
UNEDF1	${}^{57}\text{Sc} \leftrightarrow {}^{57}\text{Ti}$
UNEDF2	${}^{57}Sc \leftrightarrow {}^{57}Ti$



Mass Model	Urca Pair
ВСРМ	⁵⁹ Ca ↔ ⁵⁹ Sc
DZ31	⁵⁹ Sc ↔ ⁵⁹ Ti
FRDM-2012	⁵⁹ Sc ↔ ⁵⁹ Ti
WS4RBF	${}^{59}\text{Sc} \leftrightarrow {}^{59}\text{Ti}$
D1M	⁵⁷ Sc ↔ ⁵⁷ Ti
HFB-24	⁵⁷ Sc ↔ ⁵⁷ Ti
UNEDF1	⁵⁷ Sc ↔ ⁵⁷ Ti
UNEDF2	${}^{57}\text{Sc} \leftrightarrow {}^{57}\text{Ti}$
SkM*	⁵⁷ Ca ↔ ⁵⁷ Sc
SkP	⁵⁷ Ca ↔ ⁵⁷ Sc
SLy4	⁵⁷ Ca ↔ ⁵⁷ Sc
SV-Min	⁵⁷ Ca ↔ ⁵⁷ Sc
UNEDF0	⁵⁷ Ca ↔ ⁵⁷ Sc



Bayesian Model Averaging – A Brief Introduction



Courtesy: P. Guiliani





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S. Lalit, 10 Dec, 2024, Slide 14

Neutrons



*Warning: Collinearity of models affects the BMA



Modeling Residuals

Modeling Residuals $y_i = y^{exp}(x_i) - y^{th}(x_i)$ $y_i = f(x_i, \theta) + \sigma \epsilon_i$

Formulation of a GP

 $f(x,\theta) \sim \mathcal{GP}(\mu, k_{\eta,\rho}(x, x'))$ $k_{\eta,\rho}(x, x') = \eta^2 e^{\sum_i -\frac{(x_i - x'_i)^2}{\rho_{x_i}}}$

Input features: $x_i := \{Z, N, p, \delta\}$

p is the promiscuity (shell effects) and $\pmb{\delta}$ is the pairing factor

Model Parameters: $\theta := {\mu, \eta, \rho_i}$



R. Jain et al., (to be submitted to PRC)



Normalized Weights

$$w_k = p(\mathcal{M}_k | y^*) = \frac{p(y^* | \mathcal{M}_k) \pi(\mathcal{M}_k)}{\sum_{\ell=1}^{11} p(y | \mathcal{M}_\ell) \pi(\mathcal{M}_\ell)}$$

Evidence Integrals $p(y|\mathcal{M}_k) = \int p(y|\theta_k, \mathcal{M}_k) \pi(\theta_k, \mathcal{M}_k) d\theta_k$



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Final Predictions

$$y(x) = \sum_{k} w_{k} y^{(k)}(x)$$

$$\sigma_{y}^{2}(x) = \sum_{k} w_{k} (y^{(k)}(x) - y(x))^{2} + \sum_{k} w_{k} \sigma_{y_{k}}^{2}(x)$$

R. Jain et al., (to be submitted to PRC)





Impact on Urca Cooling





Summary: Cooling + Masses

- Quantifying the strength of Urca Cooling in accreting neutron star crusts is essential to improve crust models.
- Urca cooling rates are sensitive to nuclear masses.
- A quantified global nuclear mass model developed in Bayesian Framework predicts a lower cooling at a higher depth.



Reduced Order Models

- Surrogate models that are simplified versions of high-dimensional models to reduce computational effort while preserving accuracy
- Have been used in fluid dynamics, structural mechanics, control systems, etc.
- Methods include:
 - Proper Orthogonalization Decomposition: Reduces dimensions by identifying dominant modes in data
 - Reduced Basis Methods: Builds reduced models from snapshots of high fidelity solutions
 - Balanced Truncation: Focuses on preserving system dynamics while reducing state space
- Advantages:
 - Reduces computational costs and times
 - Enables real-time analysis and control of systems
 - Facilitates understanding and decision-making in complex systems



Dynamic Mode Decomposition

- Dynamic Mode Decomposition (DMD) is a data-driven technique used to analyze the dynamics of complex systems by decomposing them into spatiotemporal modes. It bridges numerical simulation and experimental data, revealing key features of system evolution.
- Key Features:
 - Extracts dominant dynamic structures from time-series data.
 - Provides insight into the system's temporal behavior, including growth, decay, and oscillations.
 - Works with both linear and nonlinear systems (via linear approximations).
- How it Works:
 - Collect snapshots of the system's state at different time steps.
 - Represent the data as a matrix and approximate it using low-rank matrices.
 - Identify eigenvalues and eigenvectors to derive modes and their dynamics.



Dynamic Mode Decomposition

Advantages:

- Requires no prior knowledge of the governing equations.
- Captures dominant dynamics with reduced complexity.
- Limitations:
 - Assumes linear dynamics between snapshots.
 - May require modifications (e.g., Extended DMD) for highly nonlinear systems.



Star Log-extended eMulator (SLM)



- Logarithm-based emulator that overcomes the limitations of DMDs
 - Does not assume linear dynamics between snapshots
 - Includes Extended DMD formalism for nonlinear systems
- Advantages:
 - Understands the underlying dynamics of system
 - High accuracy, low computational cost
 - Uses greedy algorithms to make predictions



Applications: Tolmann-Oppenheimer-Volkoff Equations



Emulating the TOV equations: dots – emulations, curves – HF solutions



ArXiv: 2411.10556

Applications: Tolmann-Oppenheimer-Volkoff Equations



Emulating the pressure, mass and k2 curves: dots – data, curves – emulations



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Applications: Tolmann-Oppenheimer-Volkoff Equations



Parametric SLM (pSLM) predictions: the pressure, mass and k2 curves: dots – data, curves – emulations

- -- Requires only 15 data sets for training
- -- Speeds up the calculation by a factor of 30,000

ArXiv: 2411.10556



Applications: Reaction Networks, CNO cycle



Abundances for various elements in the CNO cycle using pSLM



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Summary: Emulators

- Emulators are a useful alternative to high fidelity solvers for uncertainty quantification
- They are computationally inexpensive
- SLM accuracy seems to be independent of the number of parameters used



Thank you!



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