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Machine learning approach to pattern recognition in nuclear dynamics from the *ab initio* symmetry-adapted no-core shell model

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A novel machine learning approach is used to provide further insight into atomic nuclei and to detect orderly patterns amidst a vast data of large-scale calculations. The method utilizes a neural network that is trained on ab initio results from the symmetry-adapted no-core shell model (SA-NCSM) for light nuclei. We show that the SA-NCSM, which expands ab initio applications up to medium-mass nuclei by using dominant symmetries of nuclear dynamics, can reach heavier nuclei when coupled with the machine learning approach. In particular, we find that a neural network trained on probability amplitudes for s-and p-shell nuclear wave functions not only predicts dominant configurations for heavier nuclei but in addition, when tested for the ²⁰Ne ground state, it accurately reproduces the probability distribution. The nonnegligible configurations predicted by the network provide an important input to the SA-NCSM for reducing ultra-large model spaces to manageable sizes that can be, in turn, utilized in SA-NCSM calculations to obtain accurate observables. The neural network is capable of describing nuclear deformation and is used to track the shape evolution along the ²⁰⁻⁴²Mg isotopic chain, suggesting a shape-coexistence that is more pronounced toward the very neutron-rich isotopes. We provide first descriptions of the structure and deformation of ²⁴Si and ⁴⁰Mg of interest to x-ray burst nucleosynthesis, and even of the extremely heavy nuclei such as ^{166,168}Er and ²³⁶U, that build upon first principles considerations.

I. INTRODUCTION

Modeling atomic nuclei from first principles (or ab initio) is a computationally demanding task. Ab initio approaches use controlled approximations and interactions informed by the few-nucleon physics only (including but not limited to two-nucleon scattering data). Therefore, they are suitable to determine reaction rates used in simulations of astrophysical processes, which involve shortlived nuclei that are impossible or difficult to be measured. The ab initio symmetry-adapted no-core shell model (SA-NCSM) [1–3] utilizes symmetries that are inherent to atomic nuclei, which allows the approach to provide ab initio descriptions of alpha clustering [4, 5] and of nuclear structure and reactions as far as the calcium region [6, 7]. While recent progress has been made in ab initio theory for applications to medium-mass nuclei [8–10], reaching heavier nuclei across the nuclear chart remains a challenge. The reason is that the number of basis states (model space size) in no-core shell models exponentially increases with the number of particles and the space they occupy. While the SA-NCSM drastically reduces the model space size based on an established symmetry-based prescription, this selection is validated through multiple simulations that ensure convergence of results. Therefore, novel computational approaches are needed to study challenging nuclear systems, such as ⁴⁰Mg that has been suggested to have an effect on xray burst nucleosynthesis modeling [11], as well as nuclei in the lanthanide and actinide regions of interest to rprocess simulations [12].

In this paper, we use a machine learning (ML) approach based on a neural network with the goal to pre-

dict nonnegligible configurations to be used in SA-NCSM model spaces in a much less computationally demanding way. Indeed, a multilayer feedforward neural network can be used to approximate complicated functions by adding hidden units until it reaches a desired accuracy [13]. Deep learning algorithms have emerged as a promising alternative to physics approaches [14] and have already been applied to nuclear physics, including extrapolations of nuclear observables [15–17], machine-learning-based inversion of nuclear responses [18], studies of the unitary limit [19], as well as optimization of experimental techniques and data analysis [20]. Here, we use data from abinitio SA-NCSM wave functions to train a network on selected light nuclei, in order to find general patterns in nuclear dynamics and use these patterns to make predictions for heavier nuclei.

Earlier SA-NCSM results for various nuclei have shown the emergence from first principles of highly ordered patterns that relate to the dominance of only a few nuclear shapes in low-lying states that vibrate and rotate (with each shape tracking with a symplectic Sp(3,R) irreducible representation) [1]. This suggests that a neural network can indeed be beneficial and by detecting these patterns it can inform us about nuclei that have not yet been modeled. The universality of these patterns ensures that the network can be applied across the nuclear chart, while retaining the properties of the training data. With a moderate size of training data, the network can identify the collection of negligible configurations that are eliminated from the SA-NCSM model space. This results in significantly fewer calculations needed to achieve convergence of results, thereby increasing the applicability of the model. Furthermore, for data training sets that

are sufficiently large, we show that the network provides practically accurate predictions of dominant configurations in light and intermediate-mass nuclei, with probability amplitudes being in a close agreement with those calculated in the SA-NCSM, as evident in calculations of $^4{\rm He}$ and $^{20}{\rm Ne}$. Remarkably, this is achieved in a tiny fraction of the time, without the need for full ab initio calculations.

Specifically, we explore the capabilities of the neural network and its efficacy in using information rooted in first principles to provide descriptions in nuclear regions where large-scale calculations are computationally demanding or practically impossible. We use the illustrative example of ⁴He to demonstrate that such a network is able to train on data from smaller model spaces to make accurate predictions for the same nucleus in larger model spaces. We compare the network predictions for the intermediate-mass nuclei, ²⁰Ne, ²⁴Si, and ²⁸Mg, to available SA-NCSM calculations, and use the network to predict results in larger model spaces. In addition, we study Mg isotopes between the proton and neutron driplines with a focus on the ground state evolution with neutron number, and provide insights into the shape coexistence phenomenon Finally, in an application of the network to extreme cases of heavy nuclei, we provide predictions for dominant configurations in ¹⁶⁶Er, ¹⁶⁸Er, and ^{236}U .

II. THEORETICAL FRAMEWORK

A. Symmetry-adapted no-core shell model

The SA-NCSM provides ab initio descriptions of nuclei [2] by taking advantage of inherent symmetries of the nuclear dynamics. The approach is able to eliminate negligible nuclear configurations based on their deformations and to achieve model spaces with a manageable size, which has allowed ab initio descriptions for intermediate and medium-mass nuclei [1, 2, 6, 21, 22]. The SA-NCSM takes as input the interaction between the nucleons. In this paper, we utilize wave functions obtained with the NNLO_{opt} nucleon-nucleon (NN) potential [23] derived in the chiral effective-field-theory framework.

The SA-NCSM many-body basis states build upon harmonic oscillator (HO) single-particle states and are labeled schematically as:

$$|\mathfrak{a}; N(\lambda \mu)\kappa L; (S_p S_n)S; JM\rangle,$$
 (1)

where S_p , S_n , and S denote proton, neutron, and total intrinsic spins, respectively. N is the total number of HO excitation quanta. The deformation is represented by λ and μ quantum numbers, which inform how prolate and oblate a state is, that is, $(\lambda 0)$ indicates a prolate deformation, (0μ) indicates an oblate deformation, whereas (0 0) denotes a spherical shape [2, 6]. The label κ distinguishes multiple occurrences of the same orbital momentum L in

a given $(\lambda \mu)$. The L is coupled with S to the total angular momentum J and its projection M. The symbol $\mathfrak a$ schematically denotes the additional quantum numbers needed to specify the basis.

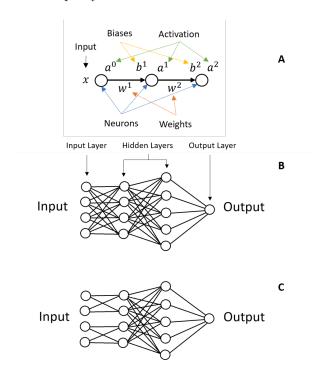


FIG. 1. (A) Feedforward neural network with interconnected neurons. Layers are labeled by index i; neurons are connected by weights w^i ; each neuron has a bias b^i and is assigned an activation function a^i . (B) A basic feedforward neural network with an input layer, two hidden layers, and an output layer with one neuron. In this network, every neuron in a layer is connected to every neuron in the previous layer. (C) A slightly more complicated network with segmented layers. Here, there are two groupings where two input neurons are exclusively connected to two hidden layer neurons but not the others.

Furthermore, the model space of the SA-NCSM is kept finite in size by introducing a maximum total number of HO excitation quanta $N_{\rm max}$ above the valence-shell configuration, that is, $N=0,\ldots N_{\rm max},$ which defines the overall size of the model space. It also has an energy resolution of $\hbar\Omega$, defined by the energy of a single HO excitation. Both $N_{\rm max}$ and $\hbar\Omega$ are parameters of the basis, with converged results coinciding with those in the infinite model-space size $(N_{\text{max}} \to \infty)$ that are independent of $\hbar\Omega$. Calculations in a model space with a limited size (low N_{max}) may result in enhanced deformations not fully developed. In this case, important basis states are not taken into account. Large enough $N_{\rm max}$ includes all necessary basis states at the cost of computational resources. In the SA-NCSM, the basis states are additionally selected above a given N cutoff (labeled as $\langle N \rangle N_{\text{max}}$). In this way, model subspaces above N

in the SA-NCSM include only non negligible configurations, and those are necessary to develop collective and clustering correlations.

B. Network Structure

A basic feedforward neural network, as shown in Fig. 1a & b, takes in inputs $\mathbf{x} \equiv \mathbf{a}^0$ in the input layer, and produces output a in the output layer. In each hidden layer i, the activation function of a neuron k is determined from the activation functions of the previous layer's neurons j, $a_k^i = f(\sum_i w_{ik}^i a_i^{i-1} + b_k^i)$, where the function fis discussed below, and the network parameters W and **b** provide, respectively, the weights between two neurons and the bias associated with each neuron [24]. The value produced by the activation function is transmitted to the neurons in the next layer where this process happens again, until the output is reached. In this work, all layers except the output layer use the rectified linear unit (RLU) activation function for f. This RLU function returns the inputted value if it is positive, and it returns zero otherwise [24]. The output layer has a sigmoid activation function S(x), where:

$$S(x) = \frac{1}{1 + e^{-x}}. (2)$$

To train a network, a set of data with inputs and their corresponding outputs is used to determine network parameters \mathbf{W} and \mathbf{b} , for each connection and neuron, that minimize the difference between the results of the network (activation function of the output layer, $a_{\text{predicted}}$), and the expected results (data outputs, a_{true}). This difference defines a loss, which is thereby calculated by a loss function. In this work, we use the Poisson loss function for N_D number of training points,

$$L = \frac{1}{N_D} \sum_{i}^{N_D} \left[a_{\text{predicted},i} - a_{\text{true},i} \ln(a_{\text{predicted},i}) \right], \quad (3)$$

which would prevent strict fitting to the true values in favor for an expected probabilistic pattern. For our case, using this instead of a standard mean squared error loss function resulted in data points that are more closely clustered around the expected values and yielded a better fit when reproducing the training data (Fig. 2). Specifically, for given $a_{\text{true}} \in [0, 1]$, the loss function has a (nonzero) minimum for $a_{\text{predicted}} = a_{\text{true}}$, and this minimum increases with increasing a_{true} [as illustrated in Fig. 2c for $f(a_{\text{predicted}}) = a_{\text{predicted}} - a_{\text{true}} \ln(a_{\text{predicted}})$. Hence, larger probabilities a_{true} effectively enter with a larger weight, and the dominant configurations are predicted most accurately. In addition, the loss function is more sensitive to overestimates, especially for very small probabilities. This ensures that the network efficiently detects the negligible configurations that can be then

eliminated from the model space in SA-NCSM production runs and reduce computational resources. This loss function appears ideal for the purposes of this network, but should be used with caution if accurate estimates are needed for all orders of magnitude.

The loss function L is minimized using an optimizer [25], which defines how the weights and biases are updated during training. The present approach utilizes Adam optimization, which is a stochastic gradient descent method with adaptive learning rates between parameters and carried momentum between updates [26]. In this study, the activation functions, loss function, and optimizer are chosen during test runs to ensure that the network fits the data well and have a capability of avoiding local minima. The training is run over 200 epochs and each epoch covers the entirety of the training set. In an effort to further avoid local minima, multiple networks may be individually trained to search for a network with nonzero and non-constant predictions for extreme cases, such as the heavy Er and U isotopes. The computational implementation of the network is achieved using the Keras deep learning application programming interfaces [27].

In this work, the neural network takes in a specific many-body basis configuration of a nuclear state and returns a value for the probability amplitude of that configuration. Since this network trains on data from SANCSM, it must have inputs relating to the parameters of the SA-NCSM model space. The presented results only use data with the NNLO $_{\rm opt}$ interaction to maintain consistency.

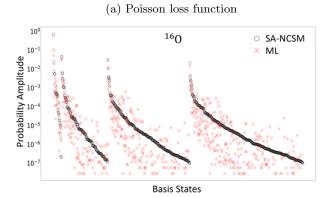
There are 11 input neurons. Each neuron has a single value as input. The inputs for the neurons are the number of protons Z, number of neutrons A-Z, total angular momentum J, excitations N, spin of protons S_p , spin of neutrons S_n , total spin S, λ , and μ according to Eq. (1), as well $N_{\rm max}$ and $\hbar\Omega$ (in MeV) basis parameters. However, for the purposes of reducing the scope of the data used by the network for this study, all of the cases have zero total angular momentum only. The inputs are entered as an array:

$$\left[Z, A - Z, N_{\text{max}}, \hbar\Omega, J, S_p, S_n, S, \tilde{\lambda}, \tilde{\mu}, N\right], \qquad (4)$$

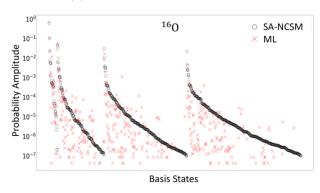
where $\tilde{\lambda} = \lambda - \lambda_0$ and $\tilde{\mu} = \mu - \mu_0$ with $(\lambda_0 \mu_0)$ being the configuration with the largest deformation and lowest intrinsic spin in the N=0 model subspace, also known as the leading SU(3) configuration, which is unique for a given nucleus. This is based on a pattern we notice in the existing SA-NCSM results across nuclei, namely, the most dominant configurations often lie in the set

$$N(\lambda \mu) = \{0(\lambda_0 \mu_0), 2(\lambda_0 + 2 \mu_0), 2(\lambda_0 - 2 \mu_0 + 2), 4(\lambda_0 + 4 \mu_0), 4(\lambda_0 \mu_0 + 2), 6(\lambda_0 + 6 \mu_0), 6(\lambda_0 + 2 \mu_0 + 2), \dots \}$$
 (5)

(see also Refs. [1, 2]). By subtracting $(\lambda_0 \mu_0)$, we expect the same pattern



(b) Mean squared error loss function



(c) Single-point Poisson loss function

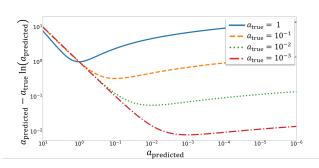


FIG. 2. Probability amplitudes across basis configurations for $^{16}{\rm O}$ in $N_{\rm max}=8$ with $\hbar\Omega=20$ MeV calculated in the large-scale many-body model (labeled as "SA-NCSM") and predicted by the network (labeled as "ML") trained on data for $^{4,8}{\rm He},\,^6{\rm Li},\,^8{\rm Be},\,^{10,12,14}{\rm C},\,$ and $^{16}{\rm O}.$ Only basis states that have SA-NCSM probability amplitudes above 10^{-7} are shown. For states where the network predicts zero, no point is plotted for "ML". (a) Predictions using Poisson loss function. (b) Predictions using the mean squared error as the loss function. (c) The Poisson loss function for a single training point shown for multiple values of $a_{\rm true}$.

 $\{0(0\ 0),\, 2(2\ 0),\, 2(-2\ 2),\, 4(4\ 0),\, 4(0\ 2),\, 6(6\ 0),\, 6(2\ 2),\, \ldots\}$ to be detected in all nuclei. For simplicity, we will drop the tilde notations from $(\lambda\,\mu)$ in further discussions of the neural network input. We emphasize that this pattern naturally emerges from calculations in complete model spaces without any a priori symmetry consideration. In addition, the training data set, as described

below, consists mostly of complete model spaces; the selected model spaces used include basis states that largely surpass in number the configurations listed in Eq. (5).

As mentioned above, in a basic neural network, when one layer serves as input to a second layer, it is common for each neuron in the second layer to be connected to every single neuron in the first layer by some set of weights (Fig. 1b). For applications to SA-NCSM, we find that a network with a segmented first hidden layer, as schematically shown in Fig. 1c, provides important improvements. Since the input layer includes different kinds of information about the nucleus, including particle numbers (Z, A - Z), basis parameters (N_{max}) and $\hbar\Omega$), spins of a nuclear state $(J, S_p, S_n, \text{ and } S)$, and its deformation, $(N, \lambda, \text{ and } \mu)$, it is beneficial to group these inputs into different sets (Fig. 3). Each of these sets will then have their own segment of the first hidden layer that they connect to. This segment will not use any values from the other sets as input. For each of these sets we know that information within has strong relationships that affects the output. Segmenting the first layer, therefore, forces the network to detect correlations between these inputs independently from other inputs.

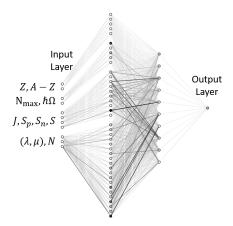


FIG. 3. Neural network structure. Circles represent neurons and lines represent weights. Darker circles signify a larger absolute value of the bias. Darker lines signify a larger absolute value of the weight. Input neurons are in the left-most layer. There are two hidden layers. The input layer and first hidden layer have their connections segmented into groups. The input labels are shown next to the input neurons in downward order. The second hidden layer and output layer connect to all neurons before them. The output is the right-most neuron, and it yields the probability amplitude for a given input basis state. The normalization of the entire wavefunction is performed afterwards. The network has 581 parameters and is trained over 15626 data points.

As illustrated in Fig. 3, the first hidden layer has some segments with more neurons to reflect the increased complexity expected for the relationship between the output and the spins, and more so, the deformation $N(\lambda \mu)$. The second hidden layer has 10 neurons, but connects to all of

the first hidden layer. Finally, there is one output neuron giving the probability of the configuration. The network is used to give a prediction for the probability amplitudes for all possible configurations for a nucleus with given J and for given $\hbar\Omega$ and $N_{\rm max}$, which are normalized afterwards. The number of neurons was chosen in an effort to have the number of parameters be about 0.01 of the number of data points in the training set (Table I). For the training data, we consider 14 complete model spaces, and 4 selected model spaces that are sufficiently larger compared to the number of configurations listed in Eq. (5).

TABLE I. Training and validation data sets. A data set for given $N_{\rm max}$ and $\hbar\Omega$ includes tens through hundreds to thousands of configurations. Complete model spaces are denoted by $N_{\rm max}$, whereas selected model spaces are denoted as $\langle N \rangle N_{\rm max}$ (e.g., for ²⁰Ne, $\langle 2 \rangle 10$ indicates the use of all basis states up to N=2 and selected basis states for N=4, 6, 8, and 10). The number of inputs used in the training data set shows the total number of $N(\lambda \mu)S_pS_nS$ configurations in all model spaces listed for each nucleus. Also shown is the dimension (number of all basis states) of the largest J=0 model space listed for each nucleus.

| Nucleus | $N_{ m max}$ | $\hbar\Omega~({\rm MeV})$ | Number of | Dimension |
|-------------------|------------------------------------------------------------------|---------------------------|-----------|-----------------------|
| | | | Inputs | |
| Training data | | | | |
| ⁴ He | 6,8,10,12,14 | 22 | 696 | 5.80×10^4 |
| ⁸ He | 6 | 15 | 218 | 3.18×10^4 |
| ⁶ Li | 8 | 15 | 208 | 6.78×10^4 |
| ⁸ Be | $8,10,12,\langle 8 \rangle 14$ | 15 | 3417 | 1.53×10^{8} |
| $^{10}\mathrm{C}$ | 8 | 15 | 1037 | 4.44×10^{6} |
| $^{12}\mathrm{C}$ | $6,\langle 6\rangle 10,\langle 6\rangle 12$ | 15 | 3211 | 1.90×10^{9} |
| $^{14}\mathrm{C}$ | $6,\langle 6\rangle 10$ | 15 | 2113 | 4.03×10^{8} |
| ¹⁶ O | 8 | 20,25 | 4726 | 3.01×10^{7} |
| Validation data | | | | |
| ²⁰ Ne | $\langle 4 \rangle 6, \langle 4 \rangle 8, \langle 2 \rangle 10$ | 15 | | 7.18×10^{10} |
| $^{28}{ m Mg}$ | $\langle 0 \rangle 6$ | 15 | | 9.77×10^9 |
| ²⁴ Si | 4 | 20 | | 2.59×10^{7} |

III. RESULTS AND DISCUSSION

Results discussed in this paper, except those for ⁴He, are presented for a training set that includes only the s-and p-shell nuclei. This includes the s-shell nucleus ⁴He and p-shell nuclei ⁶Li, ⁸He, ⁸Be, ^{10,12,14}C, and ¹⁶O, as listed in Table I. The neural network predicts probability amplitudes for the lowest 0⁺ wave function of each nucleus, which are subsequently normalized to one. In this section, we present network validation (for ⁴He and ²⁰Ne), as well as network predictions in heavier nuclei (²⁴Si, ²⁸Mg, ^{166,168}Er and ²³⁶U), together with a shape evolution along the Mg isotopic chain.

We note that the network results are for the specific interaction used in the training data set, namely, for the NNLO_{opt} NN chiral potential, which has been found to reproduce various observables, such as 3- and 4-nucleon binding energy [23], electric dipole polarizability [28], analyzing power [29], and B(E2) transition strengths [21], without the three-nucleon (3N) forces. In addition, limited number of SA-NCSM calculations have been performed with other chiral potentials, and we find that contribution percentages slightly vary, but dominant features remain [1, 30]. This also holds for the NNLO_{sat} [31], for which we include the corresponding 3N forces as averages, leading to large improvement in binding energy with no effect on the dominant pattern in the wave functions [6]. Hence, the present neural network approach could be applied to pattern recognition in calculations from any NN+3N chiral potentials, given a sufficiently large training data set.

A. Network validation

We study the capability of the neural network to predict larger model spaces by training on smaller ones, and show an example for the $^4{\rm He}$ ground state (Fig. 4). In this case, we expect that $^4{\rm He}$ develops its most significant structures in lower $N_{\rm max}$ spaces and hence, no new patterns are expected as one goes from $N_{\rm max}=12$ with 22,716 basis states to $N_{\rm max}=14$ with 58,080 basis states. However, probability amplitudes for low-N basis states often vary with an increase in the size of the model space. Indeed, the results show that the network has the capability to make close predictions for the probability amplitudes of basis states for data similar to the training data.

The most important objective of using machine learning in the present approach is to predict information about heavier nuclei using ab initio calculations for light nuclei. As an illustrative example we validate the network with results for the intermediate-mass nucleus $^{20}\mathrm{Ne}$ from calculations of s- and p-shell light nuclei for $N_{\text{max}} =$ 6, 8, and 10 (Table I). We show predictions for the largest model space $N_{\rm max}=10$ (Fig. 5). What we predict is an sd-shell nucleus, ²⁰Ne, which is not included in the training set. We utilize a $N_{\rm max}=10$ model space that consists of about 10¹² basis states that is currently infeasible, but can be reduced to several million SA basis states with a solution in the SA-NCSM [1]. Remarkably, we find that the network predictions for the most significant configurations in the ²⁰Ne ground state follow the same pattern as the one revealed by the SA-NCSM and is associated with an important and widely spread feature of nuclear dynamics, namely, vibrations of equilibrium shapes [1]. Note that for ²⁰Ne, $(\lambda_0 \mu_0)$ is (80), and the dominant configurations of Eq. (5), 0(80), 2(100), 2(62), 4(120), ... are clearly revealed, as shown in Fig. 5. Even beyond, the network results are in a very close agreement with the SA-NCSM probability amplitudes. This implies that once the network is trained on light nuclei, it can efficiently provide probability amplitudes

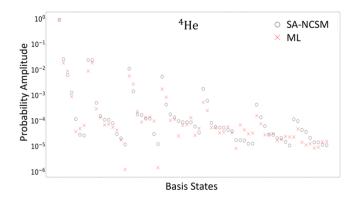


FIG. 4. Probability amplitudes across basis configurations for 4 He in $N_{\rm max}=16$ calculated in the large-scale many-body model (labeled as "SA-NCSM") and predicted by the network (labeled as "ML") trained on data for 4 He in $N_{\rm max}=8$, 10, 12, and 14. Only basis states with SA-NCSM probability amplitudes above 10^{-5} are shown. For states where the network predicted zero, no point is plotted for "ML".

for a deformation-based wave function and related information, e.g., shape dominance and coexistence along with moments of inertia, without the need for large-scale computationally intensive calculations. It can also provide an upper estimate for E2 transition strengths. This result is important as it shows that it is not necessary to include sd-shell nuclei in the training set to get good predictions in the sd-shell region. However, we find that it is imperative to use sufficiently large training data sets, as tests with networks trained only on a few nuclei result in larger deviations, but increasing the volume of training data remedies this.

It is interesting to examine the weights and biases of the network itself (Fig. 3). From the proton and neutron inputs, the outgoing weights are relatively small. This is expected since the dominant configuration $(\lambda_0 \mu_0)$ is subtracted from the $(\lambda \mu)$ before it enters as an input, as discussed above. This dominant configuration depends heavily on the numbers of these particles. If this subtraction was not implemented, then we would expect more heavy weights from those neurons. In our case, the network is building on top of the dominant $(\lambda_0 \mu_0)$, so while the proton and neutron numbers still affect the pattern, the network uses them for fine tuning since the pattern is indeed similar across different nuclei. The largest deformation generally comes from a zero spin state, so this is likely reflected in how the total spin S neuron has the heaviest weight within the spin segment of the network. The pattern previously noted in Eq. (5) also depends on N, thus the network has multiple relatively heavier weights from the N input neuron. The heaviest weights in the network are between the two hidden layers, most notably from the spins and the deformation segments in the first hidden layer, showing that the network reflects the complexity associated with these quantum numbers. The biases mainly vary in the first hidden layer. Both

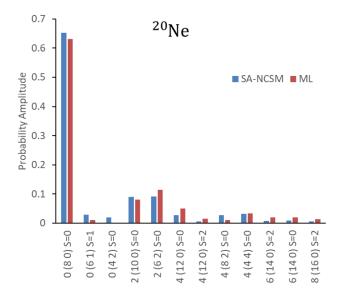
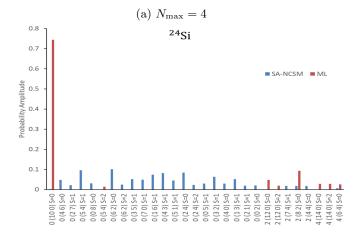


FIG. 5. Probability amplitudes of dominant configurations predicted for the $^{20}\mathrm{Ne}$ ground state in $N_{\mathrm{max}}=10$ by the network trained on $^{4.8}\mathrm{He},~^{6}\mathrm{Li},~^{8}\mathrm{Be},~^{10,12,14}\mathrm{C},$ and $^{16}\mathrm{O},$ as compared to the SA-NCSM calculations. The configurations are labeled by $N(\lambda\,\mu)$ and shown for the largest SA-NCSM probability amplitudes $\geq\,1\%.$ The configurations are labeled by $N(\lambda\,\mu)S$ and, for each N, listed left to right with decreasing deformation.

the biases in the second hidden and the weights connecting to the output neuron have a smaller range. This suggests that the network calculates the final probability by taking into account a number of similar contributions rather than arising from a single main pattern stored in one neuron in the last hidden layer.

B. Predictions for sd-shell nuclei

We present another example for an sd-shell nucleus, the ground state of ²⁴Si, where we compare no-core shellmodel results in small model spaces to the network prediction (Fig. 6). The results largely differ when the SA-NCSM calculations are limited to $N_{\text{max}} = 4$. In this very small model space, the dominant deformations cannot fully develop. However, larger model spaces are very computationally intensive and prohibitive for many models (the complete space in $N_{\rm max}=6$ has dimension of $\sim 10^{11}$). The network predicts (100) to be the most dominant configuration even within a limited model space, and in addition recognizes large configurations in N=2and N=4 from the pattern of Eq. (5) (Fig. 6a). In contrast, the limited $N_{\text{max}} = 4 \text{ SA-NCSM}$ calculations, as expected, fail to account for such collective correlations. The reason is that, within an N_{max} model space, the no-core shell model minimizes the ground state energy and to achieve this, introduces spurious (less de-



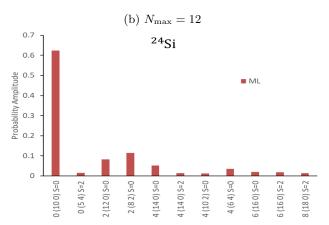
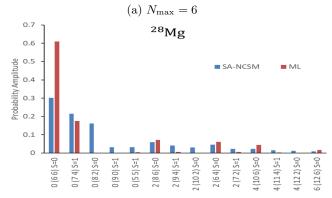


FIG. 6. Probability amplitudes of dominant configurations predicted for the $^{24}{\rm Si}$ ground state by the network ("ML") trained on $^{4,8}{\rm He},\,^6{\rm Li},\,^8{\rm Be},\,^{10,12,14}{\rm C},\,$ and $^{16}{\rm O},\,$ and compared to available SA-NCSM calculations. (a) $N_{\rm max}=4$ model space; (b) $N_{\rm max}=12$ model space, with configurations with probability amplitudes $\geq 1\%.$ The configurations are labeled by $N(\lambda\,\mu)S$ and, for each N, listed left to right with decreasing deformation.

formed) configurations that favor lower energies at the given model space cutoff. The network, however, builds upon the collective correlations and clearly reveals some of them even in relatively small model spaces. In larger model spaces, the network suggests a slightly smaller contribution of the predominant shape as other shapes become slightly more important (Fig. 6b). The network results point to physics that is similar to neighboring nuclei and clearly omitted in no-core shell-model calculations in limited model spaces. This information is critical for the construction of a selected SA-NCSM model space for ²⁴Si, which, as a next step, will be used as input to large-scale SA-NCSM calculations for first ab initio predictions of various observables, such as energy spectrum, radii, and reaction rates, for low-lying states in ²⁴Si that currently cannot be measured directly.

For an example of predicting an even heavier nuclei,



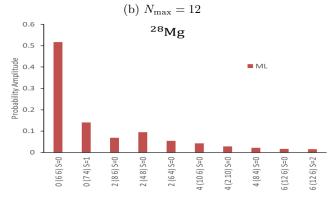


FIG. 7. Probability amplitudes of dominant configurations predicted for the $^{28}{\rm Mg}$ ground state by the network ("ML") trained on $^{4,8}{\rm He},\,^6{\rm Li},\,^8{\rm Be},\,^{10,12,14}{\rm C},\,$ and $^{16}{\rm O},\,$ and compared to available SA-NCSM calculations. (a) $N_{\rm max}=6$ model space; (b) $N_{\rm max}=12$ model space, with configurations with probability amplitudes that are $\geq 1\%.$ The configurations are labeled by $N(\lambda\mu)S$ and, for each N, listed left to right with decreasing deformation.

we study the challenging ²⁸Mg (Fig. 7), which has been suggested to lie in the so-called island of inversion largely affected by the higher pf shell [22]. In this case, the network is also capable of detecting the significant configurations, and even reproduce the probability amplitudes to a good degree. In particular, for $N_{\text{max}} = 6$, the most dominant configuration is predicted by the network, but with larger probability as compared to the SA-NCSM calculations, while another important configuration, 0(82)S = 0, is drastically underestimated (with probability amplitude of only 0.084%). The second dominant shape 0(74)S = 1, when combined across proton and neutron spins, is also in a reasonable agreement. We note that here, as is the case with the results of Fig. 6a, for comparatively small N_{max} values (Fig. 7a), the network that is trained on collective features predicts a more pronounced collective pattern compared to the SA-NCSM results, whereas the latter reveals a larger shape mixing as a result of the ground-state energy minimization in the finite model space. It is interesting that the

network recognizes the most dominant S=0 mode, and especially the most dominant S=1 modes which are often suppressed within the light nuclei in the training set. The dominant configurations for N=2 and N=4 are also found to be well reproduced, but here again some slightly less significant modes are neglected entirely according to the network output. The network is then applied to the larger $N_{\rm max}=12$ model space (Fig. 7b), where the results reveal a very similar pattern as the one predicted for $N_{\rm max}=6$, with a slightly smaller $(6\,6)S=0$ contribution. Hence, the network suggests a predominance of a triaxial shape in $^{28}{\rm Mg}$, the same one detected at the smaller model space, and some admixture with an oblate shape.

We expect that including more isotopes in the training set with different numbers of protons and neutrons would further improve the network accuracy for both ²⁴Si and ²⁸Mg, since a large part of the current training set consists of nuclei with equal numbers of protons and neutrons.

C. Shape evolution in Mg isotopes

As an important outcome, we utilize the neural network to study shape evolution in Mg isotopes, from the proton-rich $^{20}\mathrm{Mg}$ to the $^{40,42}\mathrm{Mg}$ near the neutron drip line. Since these are open-shell nuclei in the intermediateand medium-mass region, these systems pose a challenge to ab initio theory. We consider even-mass nuclei only, since all of the training data is for even-mass nuclei. For each isotope, the $(\lambda \mu)$ quantum numbers of the basis states can be directly linked to the deformation β and triaxiality γ shape parameters [32, 33] (Fig. 8). All of the isotopes favor prolate shapes except ²⁸Mg, which is triaxial, and ³⁰Mg, which is oblate. For example, the ²⁶Mg deformation distribution predicted by the network closely resembles the E2 transition density previously calculated in the constrained Hartree-Fock-Bogoliubov plus local quasiparticle random-phase approximation method [34]. In all cases, there is a predominant shape, which has the expected $(\lambda_0 \mu_0)$ deformation and intrinsic spin S=0, followed by another shape with $(\lambda_0 + 1 \mu_0 - 2) S = 1$. In general, a large ratio of the probability amplitudes for the secondary shape relative to the dominant shape indicates a strong interplay of the two shapes, pointing to a shape coexistence in the low-lying energy spectrum. It is interesting that the network suggests multiple cases where the shape coexistence becomes pronounced, namely, the neutron-rich $^{28}\mathrm{Mg},~^{30}\mathrm{Mg},~^{36}\mathrm{Mg},~^{40}\mathrm{Mg},~\mathrm{and}~^{42}\mathrm{Mg}.$ This may be a result of reduced spin-orbit interaction toward the drip line. In addition to these two shapes, there are other S=0 configurations that appear important, however, they follow the pattern of Eq. (5) and are associated with vibrations of the dominant shape rather than the existence of a new equilibrium shape (cf. Ref. [1]).

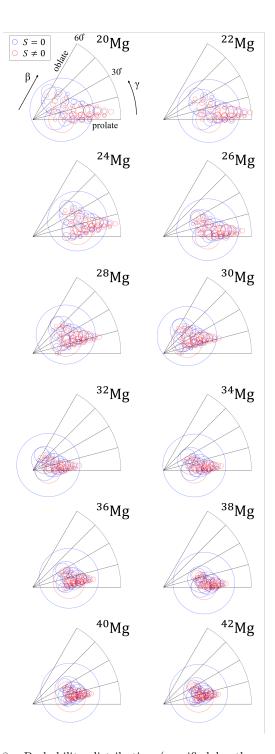
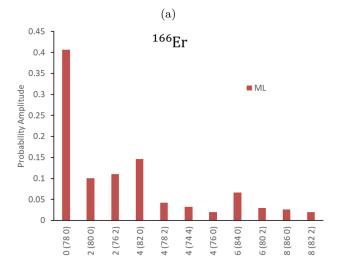
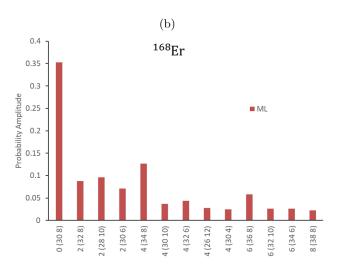


FIG. 8. Probability distribution (specified by the area of the circles) across the deformation β and triaxiality γ shape parameters for various isotopes of Mg, for $N_{\rm max}=12$ and $\hbar\Omega=15$ MeV, trained on ^{4,8}He, ⁶Li, ⁸Be, ^{10,12,14}C, and ¹⁶O. The total spin that contributes most to each configuration is also shown for S=0 (blue) and nonzero S (red).





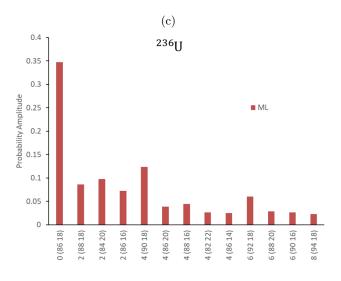


FIG. 9. Largest probability amplitudes ($\geq 1\%$) across basis configurations predicted by the network for (a) $^{166}{\rm Er}$ in $N_{\rm max}=12$, (b) $^{168}{\rm Er}$ in $N_{\rm max}=12$, and (c) $^{236}{\rm U}$ in $N_{\rm max}=12$, and for $\hbar\Omega=15$ MeV. Network trained on $^{4,8}{\rm He}$, $^{6}{\rm Li}$, $^{8}{\rm Be}$, $^{10,12,14}{\rm C}$, and $^{16}{\rm O}$.

D. Neural network results for the actinide and lanthanide region

Using the neural network, it is even possible to make predictions for extremely heavy nuclei, as shown in Fig. 9 for ^{166,168}Er and ²³⁶U. For such nuclei, full *ab initio* calculations are definitely not feasible. Yet, the network is able to suggest a very similar pattern in these nuclei as the one observed in light and medium-mass nuclei. There is only a small difference, namely, vibrations of the nuclear shape along the x and y axes are comparable to those in the z direction at N=2. Nonetheless, the network clearly recognizes a set of dominant configurations that resembles the one suggested by earlier studies with schematic interactions [35–37]. For ¹⁶⁶Er, Ref. [35] suggests the choice of $(\lambda_0 \mu_0) = (780)$ with all spins assumed to be zero, which has reproduced E2 experimental data. For ¹⁶⁸Er and ²³⁶U, we use the leading $0(\lambda_0 \mu_0)$ configurations as done for the lighter nuclei presented above, namely, 0(30.8) for 168 Er and 0(86.18) for 236 U. We note that a different $(\lambda_0 \mu_0)$ is suggested in Ref. [37] based on the use of pseudo-SU(3). Comparing the networks predictions with the results of these earlier papers, we find that the network reproduces the dominant configurations and the overall pattern, while future studies of B(E2) strengths based on the neural network results will provide insight on the interplay of vibrations in the z, x, and y directions in these nuclei. While the earlier models [35, 36] assume symmetries, the network results are especially important to provide information on the degree of symmetry breaking. If more than one dominant shape is suggested by the networks, these shapes can be then studied by models with schematic interactions that admit symmetry-breaking terms as done, e.g., in Ref. [38]. As mentioned above, the network is trained on nuclei that are very light compared to the Er and U isotopes, so we expect that adding significantly more data from much heavier nuclei, e.g. SA-NCSM calculations around mass A = 50, will improve the network predictions at this scale, since increasing the volume and diversity of training data is indeed beneficial.

IV. CONCLUSION

In this study, we demonstrate that a neural network provides an efficient mechanism – statistical in its nature – to describe dominant features of nuclei, from light to heavy mass, by using first principles input. In particular, we showed that the network was able to train on data from a single nucleus to make accurate predictions of dominant configurations for larger $N_{\rm max}$ model spaces, as in the case of ⁴He. Training on multiple s- and p-shell nuclei allowed the network to accurately predict the dominant configurations for the sd-shell nucleus of ²⁰Ne. This validates the network and suggests that the network is not limited to predicting extremely similar nuclei.

In addition, the network results reasonably agreed with

existing *ab initio* SA-NCSM calculations for ²⁸Mg, and we found that the network was especially suitable for identifying the nonnegligible configurations, which, in turn, could be used to inform the selection of model spaces for large-scale SA-NCSM calculations of various observables, including energies, radii, and electromagnetic moments and transitions. We further showed, for the illustrative example of ²⁴Si, that the network was capable of detecting the important collective correlations even in smaller model spaces where shell-model calculations fail to account for those.

As a notable outcome, the neural network was capable of detecting a common feature of nuclear dynamics, namely, vibrations of equilibrium shapes, thereby applicable across the nuclear chart. This is important, since the network is not used to extrapolate to regions of different physics compared to the training data set.

The neural network was utilized to study the interesting phenomenon of shape coexistence and even to reach heavy nuclei. In particular, the network results for different Mg isotopes were used to study the deformation distribution within each isotope, suggesting an interplay of two shapes as recognized in the neutron-rich ^{28,30}Mg, ³⁶Mg, and ^{40,42}Mg. Furthermore, the nature of the network allowed us to apply the approach to extreme cases that may never be accessible by ab initio modeling, such as ^{166,168}Er and ²³⁶U. Remarkably, this was accomplished with training on s- and p-shell nuclei only. While adding sd-shell nuclei to the training set will help the network achieve more accurate predictions, the lighter nuclei appear sufficient for the network to detect the patterns that can be used for these significantly heavier nuclei cases. We note that the network results remain to be tested against other observables, such as E2 transitions.

We emphasize two important features of the network. We found that substantially increasing the volume of the training data set led to better predictive capability, which suggests that, as more SA-NCSM calculations are performed and added to the training set, the training data will become richer and hence, will increase the network

predictive power. Another feature of the neural network is that the minimization procedure depends on a seed for a random number generator used to update base weights and biases. Hence, the loss function can reach a local minimum, preventing the network from finding the best fit to the patterns presented. Within a given run, there is an innate variability of how the weights and biases are chosen. Hence, the results could be different even with two networks trained on the same data. To resolve this. we create multiple networks that are trained on the same data subset, and select the one with the lowest loss. This network is then used to continue training with the complete data set. Alternatively, multiple iterations may be performed, with the goal to reach the global minimum. Such a procedure will result in improved predictions and is part of ongoing work.

In short, we construct a novel machine learning approach that, coupled with large-scale *ab initio* SA-NCSM calculations, provides further insight into atomic nuclei, and is capable of detecting orderly patterns amidst a vast data of large-scale calculations. This approach is ideal for studies and predictions of dominant shapes across the nuclear chart.

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