Neurosymbolic Learning for Science

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Abstract

Neurosymbolic Programming (NP) techniques have the potential to accelerate scientific discovery across fields. These models are able to interface with symbolic domain knowledge from scientists, such as prior knowledge and experimental context, to produce interpretable outputs. NP models combine neural and symbolic components to learn complex patterns and representations from data, using high-level concepts or known constraints. Here, we identify opportunities and challenges between current NP models and scientific workflows, with real-world examples from behavior analysis in science. We define concrete next steps to move the NP for science field forward, to enable its use broadly for scientists across the natural and social sciences.

1 Introduction

One of the grand challenges in the artificial intelligence and scientific communities is to find an AI scientist: an artificial agent that can automatically design, test, and infer scientific hypotheses from data. This application poses a number of distinct challenges for existing machine learning techniques because of the need to ensure that any new theories that are formulated from data are consistent with our prior scientific knowledge, as well as to enable scientists to reason about the implications of new hypotheses and experimental designs.

The distinct requirements to satisfy scientific discovery in a variety of diciplines have pushed the community to explore techniques such as symbolic regression [Cranmer, 2020] a variety of approaches to interpretable machine learning [Ustun and Rudin, 2017, Doshi-Velez and Kim, 2017, Kleinberg et al., 2018, McGrath et al., 2021], as well as the use of program synthesis to understand biological systems [Koksal et al., 2013]. These techniques have helped the community make significant progress in a number of different applications, but we are still far from solving the grand challenge.

We focus on the opportunities behind an important class of learning techniques based on *Neurosymbolic Programming* (NP) [Chaudhuri et al., 2021]. This class of techniques combine neural and symbolic reasoning in order to build models that incorporate prior expert knowledge and strong constraints on model behavior and structure. NP is capable of producing symbolic representations of theories which can be analyzed and manipulated to answer rich counterfactuals.

We argue that NP has an important role to play in automating scientific discovery in a way that builds on existing knowledge and leads to finding new scientific insights. NP allows for a new line of attack on this challenge: represent scientific hypotheses as programs in a Domain Specific Language (DSL) and use neurosymbolic program synthesis to automatically discover these programs. An important benefit of this approach is that a user can incorporate complex forms of prior knowledge (for example, known features and constraints) into the design of the DSL. The NP learning algorithms can then follow principles of reasoning found in classical science. Also, models learned this way are often similar to the code that human domain experts write during manual scientific modeling. Collectively, these characteristics enable an interactive process through which a neurosymbolic AI system and a

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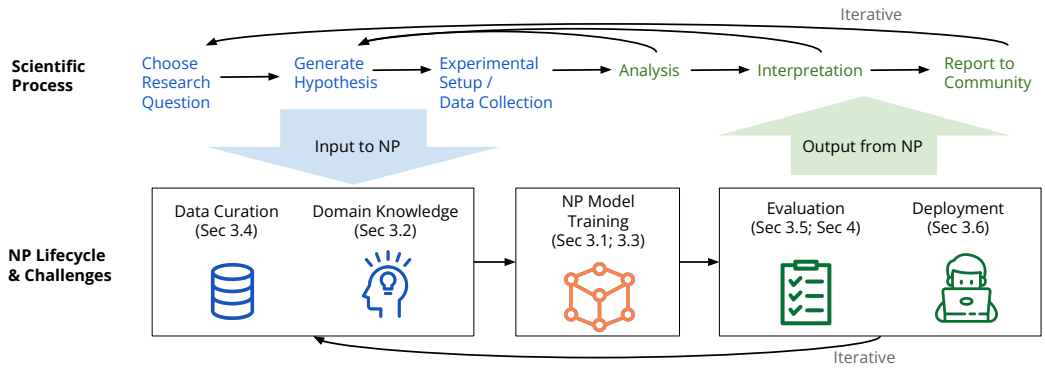

Figure 1: Neurosymbolic learning has the potential to improve the scientific process end-to-end. However, there are challenges that exist at the intersection of neurosymbolic learning and science, which we outline in Section 3, with a focus on evaluation and benchmarking in Section 4.

human expert collaborate on evidence-based reasoning and the discovery of new scientific facts. In this paper, we use behavior analysis in science as a concrete example to demonstrate the challenges and opportunities of NPs techniques and its contributions in the scientific context.

Behavior Analysis as Running Example. Many of the underlying challenges in behavior analysis are shared in other scientific domains, from monitoring vital signs, to modeling physical systems, to dynamics of chemical reactions. Fundamentally, behavioral data is a type of time-series data, and this data type is common across the sciences as many phenomenon is spatiotemporal in nature. Models that are developed for behavioral data have the potential to be applied broadly in science. Furthermore, behavior analysis illustrates common challenges with scientific data analysis. Scientists generally study behaviors that are rare with imbalanced data. Datasets vary across labs, organisms, and experimental setups, and are often noisy and imperfect. Finally, automatic behavior quantification is becoming increasingly crucial in a range of scientific fields, such as in neuroscience, ecology, biology, and healthcare monitoring. As automatic behavior analysis and neurosymbolic learning are both developing fields, there are many exciting opportunities to explore at their intersection.

Background of behavior analysis. An important objective of behavior analysis is to quantify behavior from video using continuous or discrete representations. We focus on the case of animal behavior analysis in science [Anderson and Perona, 2014, Datta et al., 2019], where there are diverse organisms and naturalistic behaviors. In this setting, a common analysis method is first to perform animal pose tracking from video [Mathis et al., 2018, Pereira et al., 2022], then categorize behaviors of interest from animal pose [Segalin et al., 2021]. We need methods that facilitates scientists' extraction of biological insights from pose trajectories or directly from recorded videos of animals.

Existing challenges in behavior analysis. Similar to other fields of science, data collection and annotation is expensive for behavioral experiments, and raw data is often imperfect and noisy. The process of data analysis is time-consuming since specialized domain expertise is generally required for identifying behaviors-of-interest and extracting knowledge. Models need to efficiently interface scientists and data at both the inputs and outputs from the scientific process (Figure 1). For NP models, leveraging domain expertise in the form of behavioral attributes has been demonstrated to improve data efficiency [Sun et al., 2021] and interpretability [Tjandrasuwita et al., 2021].

Beyond existing works, there is a variety of domain expertise that requires new algorithmic designs to integrate into the NP workflow, such as experimental context, ethograms, and scientific spatiotemporal constraints. Incorporating such domain knowledge has the potential to enable NP models to be more robust to noisy and imperfect data. Furthermore, when black-box models are used for studying behavior, it is difficult to diagnose errors and interpret model output. NP models have the potential to produce symbolic descriptions of behavior (Figure 3), which enables experts to connect model interpretations with other parts of the science analysis workflow. For example, for describing behavioral differences across different strains of mice. Finally, to enable the use of NP models in real-world science pipelines, these models must be scalable and produce robustly reproducible interpretations, which is a shared challenge across science.

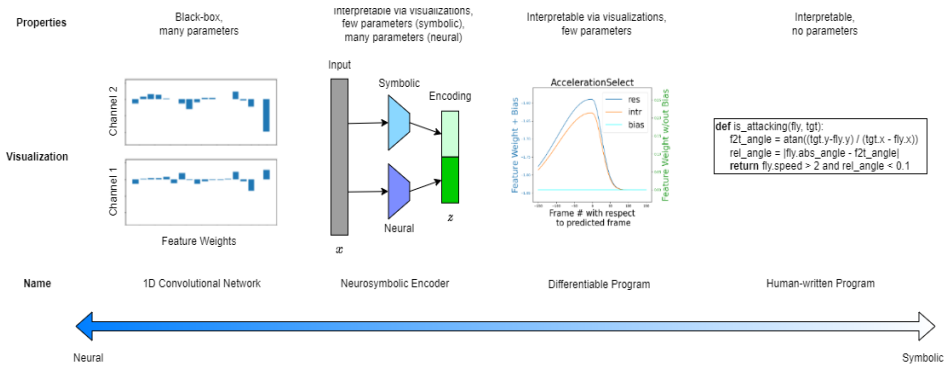

Figure 2: Examples from behavior analysis that illustrate the space of neurosymbolic models.

2 Neurosymbolic Programming techniques

In NP, the objective is to learn *neurosymbolic programs*, which incorporate latent representations computed by neural networks and symbols that explicitly capture pre-existing human knowledge, and connect these elements using rich architectures that mirror classical models of computation. The programs, assumed to belong to a DSL, are learned using a combination of gradient-based optimization, probabilistic methods, and symbolic techniques.

The range of NP methods vary in the degree to which they use neural vs. symbolic reasoning (Figure 2). The two ends of the spectrum correspond to purely neural—a 1D convolutional network—and purely symbolic techniques—a human written-program used for weak supervision [Tseng et al., 2022] —respectively, whereas the techniques close to the center are both neurosymbolic. Specifically, the model in the center-left is a neurosymbolic encoder [Zhan et al., 2021], while the model in the center-right is a program with differentiable parameters [Tjandrasuwita et al., 2021].

For an example of the strengths and weaknesses of each model, assume that we have a scientific hypothesis to test on a dataset. On one side, the fully symbolic model would involve an expert-written program that encodes the hypothesis in a general programming language. This program requires no learnable parameters and is fully interpretable and, if needed, can be iteratively improved. However, this method is also brittle, and the program must be engineered to handle all the dynamics of the dataset. This is intractable for models with complex dynamics. On the other side, the purely neural model would model the hypothesis directly using the dataset. Such models generalize well to the dataset, but offer limited interpretability and control over the generated hypothesis.

Differentiable programs. While deep learning models are powerful choices when given sufficient high-quality data, they are generally prone to overfitting and difficult to interpret. To address some of these weaknesses, NP methods have been applied in a variety of machine learning paradigms to leverage the advantages of neural networks and programs. For example, in [Shah et al., 2020] introduce a Neural Admissible Relaxation (NEAR) technique in a supervised behavior classification setting. Their goal is to synthesize differentiable programs, a powerful way of engineering systems with both discrete architectures and continuous parameters. On a high level, a differentiable program α_θ for a programming language L is defined as a composition of functions $\sigma \in L$ such that the parameters of the function $\sigma_\theta \in \mathbb{R}$ are differentiable. A differentiable program follows the syntax defined by a DSL which can consist of parametric functions (multi-layered perceptrons, linear transformations), algebraic functions (`add`, `multiply`), and higher-order functions (`map`, `fold`). Furthermore, the composition of these differentiable functions is also differentiable through the chain rule. This property enables resulting programs from Shah *et al.*'s framework to be fully differentiable. The programs synthesized by NEAR compete with the performance of a black-box recurrent neural network while being much more human-interpretable. Fig 3 is an example program discovered by NEAR for behavior analysis.

Though NEAR and follow-up work [Cui and Zhu, 2021] demonstrate great promise in terms of accuracy on behavior classification tasks, their programs are still written in a fairly generic DSL, but the results may be difficult to interpret by domain experts. [Tjandrasuwita et al., 2021] directly addresses this challenge in the context of explaining the difference between different sets of behavior

$$
\begin{aligned}
&\textbf{map}\ (\textbf{fun}\ x_t. \\
&\quad \textbf{if}\ \mathit{DistAffine}_{[.0217];-.2785}(x_t) \\
&\quad\ \textbf{then}\ \mathit{AccAffine}_{[-.0007,.0055,.0051,-.0025];3.7426}(x_t)\ \textbf{else}\ \mathit{DistAffine}_{[-.2143];1.822)}(x_t))\ x
\end{aligned}
$$

Figure 3: A program generated with NP techniques [Shah et al., 2020]. The above program classifies the “sniff" action between two mice. An interpretation is that if the distance between two mice is small, they are doing a “sniff"; otherwise, they are only doing a “sniff" if the accelerations are small.

expert annotations. Their primary contribution is to replace generic higher-order functions over recursive data structures, e.g. `map` and `fold`, with a differentiable temporal filter operation, the Morlet Filter. The filter models temporal information in a highly data-efficient manner, and can be interpreted as a human's impulse response to a given behavioral feature for classification.

3 Opportunities and Challenges at the Intersection of Neurosymbolic Learning and Science

The techniques outlined above demonstrate some benefits of the NP framework. However, there are still major gaps between current NP approaches and practical use cases for scientists. There exist challenges across the life-cycle of NP algorithm development and deployment, from dealing with noisy input data, to scalable model training, to evaluation and tool building an deployment (see Figure 1). We draw attention to these challenges to encourage the research community to collaborate in the development of new NP methods that can connect analysis efforts and broadly accelerate progress across science.

3.1 Scalability challenge in NP

Inductive synthesis. A large body of works on program synthesis has focused on *inductive synthesis*, or synthesizing programs from examples [Lau and Weld, 1998, Gulwani, 2011, Devlin et al., 2017]. While such a goal is on the surface-level similar to performing machine learning (ML) with programs as models, a key difference is that ML approaches hinge on defining a clear space of models (i.e. neural networks, support vector machines, decision trees) and generalizing to unseen data. In contrast, much work in inductive synthesis considers an arbitrary space of programs and spends significant effort on sample engineering, treating them as noiseless specifications. As a result, inductive synthesis scales poorly with increase in program length and number of examples.

Scaling NP in science. To tackle scalability in science, models need to handle large and potentially noisy datasets, high-dimensional input space, and a variety of analysis tasks. Recently, NP research [Shah et al., 2020, Cui and Zhu, 2021] propose frameworks that scale to large datasets given an expressive DSL. These works are instantiated in behavior analysis: learning programs on temporal trajectory data to reproduce expert annotations of behavior, and similar to other scientific data, contains noisy labels. These works tackle the challenge of discovering programs with parameters, which can be directly optimized through popular gradient optimization techniques. While NP methods provide a means of scaling inductive synthesis to scientific datasets, these techniques often involve combining a discrete search over an exponential space of programs with continuous optimization.

Challenges for enabling scalability. While the above works are able to leverage the strengths of programs, such as data efficiency and interpretability, there still remain challenges in scalability. The differentiable program synthesis library (dPads) [Cui and Zhu, 2021] addresses the computational efficiency challenge of having to synthesize discrete program architectures and perform gradient optimization at runtime. To do so, dPads trades off computation with the cost of storing many more parameters in memory, with heuristics to mitigate memory usage. However, training fully neural models is often more efficient than NP models, which requires searching through an exponentially large space of symbolic architectures. Furthermore, scalability has not been broadly explored for different types of scientific data, such as video recordings, which is much higher dimensional than trajectory data. Finally, the effectiveness of program synthesis may still be limited by the expressivity of the DSL, which requires experts to spend time to encode domain knowledge, such as expert-designed behavior attributes [Sun et al., 2021] and temporal filters [Tjandrasuwita et al., 2021] (further discussed in Section 3.2).

Scalability in other domains. Such scalability challenges also arise in symbolic regression [Cranmer et al., 2020], where the goal is to learn exact mathematical relations between variables by searching the space of mathematical expressions. In this case, the regression scales poorly with the number of interactions between variables. [Cranmer et al., 2020] posit a solution that separately models the variable-variable interaction and the variable transformation through a graph neural network. Distilling the neural networks to symbolic functions is then tractable via symbolic regression. This domain sufers from the same problems.

3.2 Encoding and Learning Domain Knowledge

The success of NP techniques often depends on how a DSL is defined. However, it may be not only time-consuming but also unclear about how to handcraft domain-specific components that work best in a scientific context. As such, works in *library learning* propose algorithms that consolidate common patterns in successful programs generated in program synthesis and add them to the current DSL, enabling the program search to discover high-performing programs with little effort.

Library learning for science. In behavior analysis, humans are capable of writing short programs, such as by designing features and heuristics [Segalin et al., 2021, Tseng et al., 2022, Eyjolfsdottir et al., 2014]. However, these programs are greatly limited by their simplicity and may not capture complex behavior. Library learning has the potential to augment human feature design, by not only synthesizing interpretable programs but also inducing high-level DSL's, given low-level, generic primitives. These models have been applied to tasks such as studying the relationship between variables in physics equations [Ellis et al., 2021]. Another example that demonstrates the potential of library learning is generative modeling in molecular chemistry [Guo et al., 2022]. The proposed library learning method is capable of handling highly data-limited settings, often found in science due to the cost of obtaining high-quality labels from experts. Library learning methods also have the advantage of enabling greater possibilities for including domain-specific knowledge, as all components in the library are written in terms of human-interpretable primitives.

Challenges of library learning for science. While DreamCoder and molecular generation have shown promise in their respective domains, it is unclear how library learning can scale to more complex scientific domains, such as behavior analysis, which often consist of thousands of frames of videos with noisy data. In addition, it is highly expensive to collect behavior annotations across a large number of behaviors, which is needed to perform traditional library learning over a representative task corpus for a given domain. In contrast, current library learning methods have been applied to contexts where each task consists of very few examples, not exceeding hundreds of data points. In addition, the labels are noiseless as opposed to situations found in behavior analysis where even experts may disagree on behavior annotations [Leng et al., 2020, Segalin et al., 2021].

Another challenge is that domain experts still need to interpret library components learned by the system. One promising approach leverages natural language to impose a stronger prior on the program search and the library learning [Wong et al., 2021], resulting in a more interpretable DSL. However, these experiments have only demonstrated the method's strength in domains that differ significantly from real-world scientific problems, such as behavior analysis, in terms of dataset size, task corpus size, and the presence of noise, similar to the above discussion. In addition to the challenge of scaling library learning to real behavior analysis settings, building a smooth interface between expertise in science, program synthesis and library learning methods, found in NP, would likely require significant engineering and research efforts (Section 3.6).

3.3 Challenges of optimization of discrete and continuous space in neurosymbolic programs

NP relies on techniques from symbolic program synthesis to facilitate interpretable and verifiable searches over the scientific hypothesis space. However, programs are inherently discontinuous, owing to their roots in mathematical logic. This makes modeling phenomena in the continuous domain challenging without modifying the way we interpret programs.

For instance, consider a simple program that is modeled by an if-then-else statement (`if condition do expr1 else do expr2`). The possible behaviors of `condition` are partitioned into two sets –

True (`1`) or False (`0`). These sets evaluate to either `expr1` or `expr2` respectively. However, scientific applications require reasoning about behaviors over a *gradient* of possibilities to be differentiable. Discrete programs are inaccurate models for these applications. Specifically, in behavior classification, modeling the "attack" action using a symbolic if-then-else expression would partition the mouse's aggression into two sets: either always attacking or not attacking at all.

We approach the continuous program optimization problem by changing the semantics of the programming language. Specifically, work on *Smooth Interpretation* [Chaudhuri and Solar-Lezama, 2010] rewrites discrete functions using their closest smooth mathematical functions. Consecutively, an if-then-else statement would be rewritten as a shifted sigmoid function with a high temperature (Figure 5). This smoothening is not restricted to a one dimension input space and specialized functions. In general, in higher dimensions, we can use Gaussian smoothing to smooth discontinuities. Such relaxations, in conjunction with other program analysis tools, allow gradient descent-based optimizers to converge to optimal programmatic models.

Continuous relaxations enable an approximate interface between neural networks and programming languages, that are essential for NP. For example, in Houdini [Valkov et al., 2018], continuous relaxations enabled constructing a functional programming language that admits neural networks and higher order functions. This construction facilitated high-level transfer of learned concepts across tasks in a life-long learning setting. In NEAR [Shah et al., 2020], the interface between neural networks and differentiable programs allowed measuring the performance of partial programs. This proved to be an $\epsilon-$admissible heuristic for synthesizing differentiable programs. However, in both of these methods, the *degree of smoothening* introduced an additional hyperparameter that significantly affected the model output. On the one hand, a high degree of smoothening introduces a smoothening error into the output of the program. This error increases with program complexity and can be hard to overlook, especially in scientific domains where the program output directly models measurements instead of classification error. On the other hand, a low degree of smoothening impedes the gradient descent optimizer to converge to an optimal value and, hence, makes training challenging.

3.4 Dealing with raw, noisy, and imperfect data

Scientific data provides an opportunity to study NP models with imperfect data in real-world conditions, such as with missing data, experiment noise, and distribution shifts.

Current NP models are not typically developed or evaluated in these settings, but by incorporating prior knowledge and known constraints, they have the potential to perform well in the presence of imperfect data. For example, for mouse social interactions, neurosymbolic models can automatically learn weak labels from a small amount of annotated examples and apply these learned models to generate weak supervision for the full dataset [Tseng et al., 2022]. Further explorations into NP models have the potential to systematically address imperfect datasets across domains.

These types of imperfect data exist throughout science: missing data in neural recordings due to hardware issues, noise in pose estimators for tracking animal movements, and distribution shifts. An additional source of noise in data is the considerable variability that exists in the label generation process, such as annotator subjectivity and ambiguity in category definitions. Furthermore, scientists are often interested in studying rare categories, such as behaviors that may occur in less than 1% of frames. NEAR and dPads [Shah et al., 2020, Cui and Zhu, 2021] addresses this by leveraging the flexibility of neural networks with symbolic domain knowledge; however, there remains challenges in improving performance and scalability before widespread deployment.

Structural discovery. In many scientific workflows, meaningful categories and structures in raw data may not be clear ahead of time, and requires unsupervised or self-supervised learning from data. For example, there are many tools for discovering new behavior categories from data without expert supervision [Pereira et al., 2020]. Recent work [Zhan et al., 2021] has tackled this using NP for behavior analysis and demonstrated that integrating domain knowledge results in more meaningful discovered categories compared to fully neural methods. However, these methods have not been widely developed. For broader impact across science, in addition to the algorithmic challenges discussed in previous sections, future research work needs to be robust to variations in experimental noise and produce reproducible interpretations of discovered structures in the data that are useful for scientists.

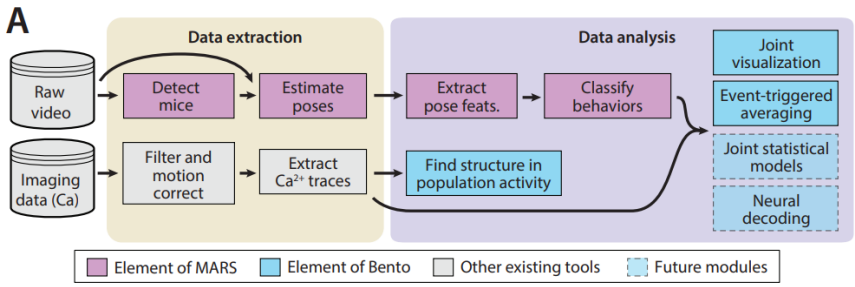

Figure 4: Functionalities of MARS and Bento [Segalin et al., 2021] in the behavior analysis pipeline.

Distribution Shifts. Distribution shifts are common in real-world scientific applications [Koh et al., 2021]. For typical black-box machine learning models, it is difficult to diagnose and address these errors. NP approaches generally learn interpretable and modular programs, which have the promise to tackle this challenge. For example, in behavior analysis, when the physical behavioral area changes in size or shape, the relative size of mice also changes. This causes errors in behavior classifiers trained in the previous arena, but we can scale and adjust the programs from NP models. Future work to improve model interpretability and distributional robustness of NP approaches is important for facilitating the integration of these models in real-world science workflows.

3.5 Evaluation of NP models

Similar to other ML models, evaluation is crucial for measuring progress in model development and understanding gaps in model performance. This is also important before real-world deployment in scientific discovery pipelines. As an important next step in reducing the gap between NP models and science, we discuss evaluation and benchmarking in more detail in Section 4, and focus on interpretability in this subsection.

The main goal of model interpretability is to obtain model insights that are understandable and actionable to humans, in order to assist scientists in the data analysis workflow. The following are commonly described properties of explanations that can be found in machine learning algorithms [Murphy, 2023], that have the potential to improve the interpretability and evaluation of NP workflows: **Compactness or sparsity:** Sparsity generally corresponds to some notion of smallness (a few features or a few parameters) ; **Completeness:** To measure if the explanation include all the relevant elements, higher-level concepts; **Stability:** To measure the extent that there are explanations similar for similar input; **Actionability:** To allow focusing on only aspects of the model that the user might be able to intervene on; **Modularity:** Explanation can be broken down into understandable parts. To study interpretability for NP models for science, we need datasets and benchmarks that is able to quantify these different dimensions in a range of scientific contexts.

3.6 Cross-Domain Analysis Science Tools

Importance of tools in science. User-friendly tools are important for facilitating the integration of ML models in real-world workflows; but have not been well-explored for NP approaches. For example, numerous tools, based on statistical analysis and ML, have been developed to interface with scientists and facilitate behavior analysis from videos [Pereira et al., 2020]. These tools assist with much of the computational pipeline for behavior classification as outlined in Figure 4, and will often provide visual interfaces that visualize relevant raw data such as video, with model outputs, such as pose data and behavior [Segalin et al., 2021]. Enabling similar tools for NP approaches has the potential to benefit existing scientific workflows. For instance, integrating NEAR into a visual interface could not only provide scientists with a user-friendly way of generating differentiable programs but also greater means of understanding the programs that NP pipelines generate. The parameters associated with programmatic primitives are likely to have a much more human interpretation [Tjandrasuwita et al., 2021] than those found in black-box neural networks.

Challenges of building NP tools. Though we foresee various benefits of integrating NP algorithms into scientific pipelines, following through with this vision will take nontrivial engineering efforts

to design a suitable interface between scientists and the algorithmic backends. Domain expertise vary in behavior analysis, for example to know behavioral attributes, or visual or textual descriptions, to known dynamics of movement, to knowledge graphs, and differs across labs and domains. Also, combining NP with more traditional human-in-the-loop machine learning methods, such as active learning offer promise to move the field forward.

4 Benchmarking

Much of the innovations in ML algorithms for scientific applications have focused on single domains, for example behavioral neuroscience [Sun et al., 2021], chemistry [Guo et al., 2022], or physics [Cranmer et al., 2020]. While many individual fields of science have seen successes through NP, consolidating underlying insights that are generalizable across domains in science remains a significant challenge. To encourage the development of new methods that work across a diverse set of scientific problems and enable comparisons between these methods, we propose to build a benchmark featuring low-dimensional spatio-temporal data, a setting where NP methods have previously demonstrated successful applications [Shah et al., 2020, Verma et al., 2018]. We believe that there are several benefits to gain from developing this benchmark for the ML and scientific communities: (1) systematic improvements across broad scientific use cases, instead of piece-wise, (2) model improvements across a variety of metrics, instead of in domain-specific dimensions, (3) increased awareness of important scientific applications that have not received as much attention from the ML community.

Challenges of benchmarking NP for science. Constructing such a benchmark presents unique challenges. First, interpreting programmatic structures requires expert domain knowledge, which can be expensive and time-consuming to obtain. In behavior analysis, evaluating learned programmatic structures requires interactions with experts in the behavioral science community to interpret these structures. This imposes a physical and manual bottleneck on evaluating outputs. A standardized benchmark will make it easier for the community to convene and interact with a panel of experts. We believe that developing a benchmark for NP pipelines is integral to moving the NP field forward.

Second, the space of NP models is broad. Each algorithm presents a different methodology for encoding expert knowledge into the machine learning lifecycle. This requires comparing models on multiple evaluation metrics. However, not all algorithms can be systematically evaluated on the same set of metrics. For instance, certain classes of NP models use stochastic search to discover the programmatic structure and the programs found by such models may not be exactly reproducible. Additionally, NP models might exhibit properties that do not have concrete evaluation metrics. For instance, classes of NP algorithms exhibit robust reproducibility. That is, the model's outputs are reproducible with small perturbations to the input data. However, to the best of our knowledge, defining such a metric quantitatively remains an open problem.

Finally, the hardware requirements for learning neural representations and symbolic functions are orthogonal. Neural network training is GPU intensive, while program synthesis is CPU intensive. This increases the cost of computation and imposes a barrier to entry for aspiring NP researchers. We need benchmarks that takes training and inference efficiency and performance into account.

5 Conclusion

Neurosymbolic programming offers clear promise to accelerate scientific discovery. The benefits are in its ability to incorporate prior knowledge and the symbolic nature of the solutions. This approach is compatible with current scientific workflows and thus has the potential to optimize scientific discovery end-to-end. However, challenges still remain in scalability and optimization stability of these approaches, comprehensive evaluations, and deployment in the form of tools. In this paper, we have demonstrated opportunities and challenges of NP in a concrete scientific application, behavior analysis. A key promise of NP is to provide a set of unifying principles that connect and generalize solutions in interpretable machine learning and prior scientific literature. We invite the science and computer science communities to adopt these methods in their scientific workflow and to contribute to the research to advance NP techniques for science.

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