NeuroDiff: Scalable Differential Verification of Neural Networks using Fine-Grained Approximation

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ABSTRACT

As neural networks make their way into safety-critical systems, where misbehavior can lead to catastrophes, there is a growing interest in certifying the equivalence of two structurally similar neural networks – a problem known as differential verification. For example, compression techniques are often used in practice for deploying trained neural networks on computationally- and energy-constrained devices, which raises the question of how faithfully the compressed network mimics the original network. Unfortunately, existing methods either focus on verifying a single network or rely on loose approximations to prove the equivalence of two networks. Due to overly conservative approximation, differential verification lacks scalability in terms of both accuracy and computational cost. To overcome these problems, we propose NeuroDiff, a symbolic and fine-grained approximation technique that drastically increases the accuracy of differential verification on feed-forward ReLU networks while achieving many orders-of-magnitude speedup. NeuroDiff has two key contributions. The first one is new convex approximations that more accurately bound the difference of two networks under all possible inputs. The second one is judicious use of symbolic variables to represent neurons whose difference bounds have accumulated significant error. We find that these two techniques are complementary, i.e., when combined, the benefit is greater than the sum of their individual benefits. We have evaluated NeuroDiff on a variety of differential verification tasks. Our results show that NeuroDiff is up to 1000X faster and 5X more accurate than the state-of-the-art tool.

1 INTRODUCTION

There is a growing need for rigorous analysis techniques that can compare the behaviors of two or more neural networks trained for the same task. For example, such techniques have applications in better understanding the representations learned by different networks [46], and finding inputs where networks disagree [52]. The need is further motivated by the increasing use of neural network compression [14] – a technique that alters the network’s parameters to reduce its energy and computational cost – where we expect the compressed network to be functionally equivalent to the original network. In safety-critical systems where a single instance of misbehavior can lead to catastrophe, having formal guarantees on the equivalence of the original and compressed networks is highly desirable.

Unfortunately, most work aimed at verifying or testing neural networks does not provide formal guarantees on their equivalence. For example, testing techniques geared toward refutation can provide inputs where a single network misbehaves [22, 31, 42, 44, 51] or multiple networks disagree [23, 34, 52], but they do not guarantee the absence of misbehaviors or disagreements. While techniques geared toward verification can prove safety or robustness properties of a single network [7–9, 15, 18, 25, 38, 41, 47], they lack crucial information needed to prove the equivalence of multiple networks. One exception is the ReluDiff tool of Paulsen et al. [33], which computes a sound approximation of the difference of two neural networks, a problem known as differential verification. While ReluDiff performs better than other techniques, the overly conservative approximation it computes often causes both accuracy and efficiency to suffer.

To overcome these problems, we propose NeuroDiff, a new symbolic and fine-grained approximation technique that significantly increases the accuracy of differential verification while achieving many orders-of-magnitude speedup. NeuroDiff has two key contributions. The first contribution is the development of convex approximations, a fine-grained approximation technique for bounding the output difference of neurons for all possible inputs, which drastically improves over the coarse-grained concretizations used by ReluDiff. The second contribution is judiciously introducing symbolic variables to represent neurons in hidden layers whose difference bounds have accumulated significant approximation error. These two techniques are also complementary, i.e., when combined, the benefit is significantly greater than the sum of their individual benefits.

The overall flow of NeuroDiff is shown in Figure 1, where it takes as input two neural networks $f$ and $f'$, a set of inputs to the neural networks $X$ defined by box intervals, and a small constant $\epsilon$ that quantifies the tolerance for disagreement. We assume that $f$ and $f'$ have the same network topology and only differ in the numerical values of their weights. In practice, $f'$ could be the compressed version of $f$, or they could be networks constructed using the same network topology but slightly different training...
data. We also note that this assumption can support compression techniques such as weight pruning [14] (by setting edges’ weights to 0) and even neuron removal [10] (by setting all of a neuron’s incoming edge weights to 0). NeuroDiff then aims to prove \( \forall x \in X, |f'(x) - f(x)| < \epsilon \). It can return (1) verified if a proof can be found, or (2) undetermined if a specified timeout is reached.

Internally, NeuroDiff first performs a forward analysis using symbolic interval arithmetic to bound both the absolute value ranges of all neurons, as in single network verification, and the difference between the neurons of the two networks. NeuroDiff then checks if the difference between the output neurons satisfies \( \epsilon \), and if so returns verified. Otherwise, NeuroDiff uses a gradient-based refinement to partition \( X \) into two disjoint sub regions \( X_1 \) and \( X_2 \), and attempts the analysis again on the individual regions. Since \( X_1 \) and \( X_2 \) form independent sub-problems, we can do these analyses in parallel, hence gaining significant speedup.

The new convex approximations used in NeuroDiff are significantly more accurate than not only the coarse-grained concretizations in RELUVal [33] but also the standard convex approximations in single-network verification tools [39, 40, 47, 54]. While these (standard) convex approximations aim to bound the absolute value range of \( y = \text{ReLU}(x) \), where \( x \) is the input of the rectified linear unit (ReLU) activation function, our new convex approximations aim to bound the difference \( z = \text{ReLU}(x + \Delta) - \text{ReLU}(x) \), where \( x \) and \( x + \Delta \) are ReLU inputs of two corresponding neurons. This is significantly more challenging because it involves the search of bounding planes in a three-dimensional space (defined by \( x, \Delta \) and \( z \)) as opposed to a two-dimensional space as in the prior work.

The symbolic variables we judiciously add to represent values of neurons in hidden layers should not be confused with the symbolic inputs used by existing tools either. While the use of symbolic inputs is well understood, e.g., both in single-network verification [39, 40, 47, 54] and differential verification [33], this is the first time that symbolic variables are used to substitute values of hidden neurons during differential verification. While the impact of symbolic inputs often diminishes after the first few layers of neurons, the impact of these new symbolic variables, when judiciously added, can be maintained in any hidden layer.

We have implemented the proposed NeuroDiff in a tool and evaluated it on a large set of differential verification tasks. Our benchmarks consists of 49 networks, from applications such as aircraft collision avoidance, image classification, and human activity recognition. We have experimentally compared with RELUVal [33], the state-of-the-art tool which has also been shown to be superior to RELUVal [48] and DeepPoly [40] for differential verification. Our results show that NeuroDiff is up to 1,000X faster and 5X more accurate. In addition, NeuroDiff is able to prove many of the same properties as RELUVal while considering much larger input regions.

To summarize, this paper makes the following contributions:

- We propose new convex approximations to more accurately bound the difference between corresponding neurons of two structurally similar neural networks.
- We propose a method for judiciously introducing symbolic variables to neurons in hidden layers to mitigate the propagation of approximation error.
- We implement and evaluate the proposed technique on a large number of differential verification tasks and demonstrate its significant speed and accuracy gains.

The remainder of this paper is organized as follows. First, we provide a brief overview of our method in Section 2. Then, we provide the technical background in Section 3. Next, we present the detailed algorithms in Section 4 and the experimental results in Section 5. We review the related work in Section 6. Finally, we give our conclusions in Section 7.

## 2 OVERVIEW

In this section, we highlight our main contributions and illustrate the shortcomings of previous work on a motivating example.

### 2.1 Differential Verification

We use the neural network in Figure 2 as a running example. The network has two input nodes \( n_{0,1}, n_{0,2} \), two hidden layers with two neurons each \((n_{1,1}, n_{1,2})\) and \((n_{2,1}, n_{2,2})\), and one output node \( n_{3,1} \). Each neuron in the hidden layer performs a summation of their inputs, followed by a rectified linear unit (ReLU) activation function, defined as \( y = \max(0, x) \), where \( x \) is the input to the ReLU activation function, and \( y \) is the output.

Let this entire network be \( f \), and the value of the output node be \( n_{3,1} = f(x_1, x_2) \), where \( x_1 \) and \( x_2 \) are the values of input nodes \( n_{0,1} \) and \( n_{0,2} \), respectively. The network can be evaluated on a specific input by performing a series matrix multiplications (i.e., affine transformations) followed by element-wise ReLU transformations. For example, the output of the neurons of the first hidden layer is

\[
\begin{bmatrix}
    n_{1,1} \\
n_{1,2}
\end{bmatrix} = \text{ReLU}
\begin{bmatrix}
    1.9 & -1.9 \\
    1.0 & 1.1
\end{bmatrix}
\begin{bmatrix}
    x_1 \\
x_2
\end{bmatrix} = \begin{bmatrix}
    \text{ReLU}(1.9x_1 - 1.9x_2) \\
    \text{ReLU}(1.1x_1 + 1.0x_2)
\end{bmatrix}
\]
Differential verification aims to compare $f$ to another network $f'$ that is structurally similar. For our example, $f'$ is obtained by rounding the edge weights of $f$ to the nearest whole numbers, a network compression technique known as weight quantization. Thus, $f'$, $n_{k,j}'$, and $n_{k,j}' = f'(x_1, x_2)$ are counterparts of $f$, $n_{k,j}$ and $n_{k,j} = f(x_1, x_2)$ for $0 \leq k \leq 2$ and $1 \leq j \leq 2$. Our goal is to prove that $|f'(x_1, x_2) - f(x_1, x_2)|$ is less than some reasonably small $\epsilon$ for all inputs defined by the intervals $x_1 \in [-2, 2]$ and $x_2 \in [-2, 2]$. For ease of understanding, we show the edge weights of $f$ in black, and $f'$ in light blue in Figure 2.

2.2 Limitations of Existing Methods

Naively, one could adapt any state-of-the-art, single-network verification tool for our task, including DeepPoly [40] and Neurify [47]. Neurify, in particular, takes a neural network and an input region of the network, and uses interval arithmetic [27, 48] to produce sound symbolic lower and upper bounds for each output node. Typically, Neurify would then use the computed bounds to certify the absence of adversarial examples [43] for the network.

However, for our task, the bounds must be computed for both networks $f$ and $f'$. Then, we subtract them, and concretize to compute lower and upper bounds on $f'(x_1, x_2) - f(x_1, x_2)$. In our example, the individual bounds would be (approximately, due to rounding) $[LB(f), UB(f)] = [-0.94x_1 - 0.62x_2 - 6.51, 0.71x_1 - 2.35x_2 + 7.98]$ and $[LB(f'), UB(f')] = [-0.94x_1 - 0.44x_2 - 6.75, 0.75x_1 - 2.25x_2 + 8.00]$ for nodes $n_{3,1}$ and $n_{3,1}'$, respectively. After the subtraction, we would obtain the bounds $[LB(f') - UB(f), UB(f') - LB(f)] = [-1.65x_1 + 1.9x_2 - 14.73, 1.68x_1 - 1.63x_2 + 14.5]$. After concretization, we would obtain the bounds $[-21.83, 21.12]$. Unfortunately, the bounds are far from being accurate.

The ReluDiff method of Paulsen et al. [33] showed that, by directly computing a difference interval layer-by-layer, the accuracy can be greatly improved. For the running example, ReluDiff would first compute bounds on the difference between the neurons $n_{1,1}$ and $n_{1,1}'$, which is $[0, 1]$, and then similarly compute bounds on the difference between outputs of $n_{1,2}$ and $n_{1,2}'$. Then, the results would be used to compute difference bounds of the subsequent layer. The reason it is more accurate is because it begins computing part of the difference bound before errors have accumulated, whereas the naive approach first accumulates significant errors at each neuron, and then computes the difference bound. In our running example, ReluDiff [33] would compute the tighter bounds $[-3.1101, 2.5600].$

While ReluDiff improves over the naive approach, in many cases, it uses concrete values for the upper and lower bounds. In practice, this approach can suffer from severe error-explosion. Specifically, whenever a neuron of either network is in an unstable state – i.e., when a ReLU’s input interval contains the value 0 – it has to concretize the symbolic expressions.

2.3 Our Method

The key contribution in NeuroDiff, our new method, is a symbolic and fine-grained approximation technique that both reduces the approximation error introduced when a neuron is in an unstable state, and mitigates the explosion of such approximation error after it is introduced.

2.3.1 Convex Approximation for the Difference Interval.

Our first contribution is developing convex approximations to directly bound the difference between two neurons after these ReLU activations. Specifically, for a neuron $n$ in $f$ and corresponding neuron $n'$ in $f'$, we want to bound the value of $ReLU(n') - ReLU(n)$. We illustrate the various choices using Figures 3, 4, and 5.

The naive way to bound this difference is to first compute approximations of $y = ReLU(n)$ and $y' = ReLU(n')$ separately, and then subtract them. Since each of these functions has a single variable, convex approximation is simple and is already used by single-network verification tools [40, 47, 49]. Figure 6 shows the function $y = ReLU(n)$ and its bounding planes (shown as dashed-lines) in a two-dimensional space (details in Section 3). However, as we have already mentioned, approximation errors would be accumulated in the bounds of $ReLU(n)$ and $ReLU(n')$ and then amplified by the interval subtraction. This is precisely why the naive approach performs poorly.

The ReluDiff method of Paulsen et al. [33] improves upon the new approximation by computing an interval bound on $n' - n$, denoted $\Delta$, then rewriting $z = ReLU(n') - ReLU(n)$ as $z = ReLU(n + \Delta) - ReLU(n)$, and finally bounding this new function instead. Figure 3 shows the shape of $z = ReLU(n + \Delta) - ReLU(n)$ in a three-dimensional space. Note that it has four piece-wise linear subregions, defined by values of the input variables $n$ and $\Delta$. While the bounds computed by ReluDiff [33], shown as the (horizontal) yellow planes in Figure 4, are sound, in practice they tend to be loose because the upper and lower bounds are both concrete values. Such eager concretization eliminates symbolic information that $\Delta$ contained before applying the ReLU activation.

In contrast, our method computes a convex approximation of $z$, shown by the (tiled) yellow planes in Figure 5. Since these tiled bounding planes are in a three-dimensional space, they are significantly more challenging to compute than the standard two-dimensional convex approximations (shown in Figure 6) used by single network verification tools. Our approximations have the advantage of introducing significantly less error than the horizontal planes used in ReluDiff [33], while maintaining some of the symbolic information for $\Delta$ before applying the ReLU activation.

We will show through experimental evaluation (Section 5) that our convex approximation can drastically improve the accuracy of the difference bounds, and are particularly effective when the input region being considered is large. Furthermore, the tiled planes shown in Figure 5 are for the general case. For certain special cases, we obtain even tighter bounding planes (details in Section 4). In the running example, using our new convex approximations would improve the final bounds to $[-1.97, 1.42].$

2.3.2 Symbolic Variables for Hidden Neurons.

Our second contribution is introducing symbolic variables to represent the output values of some unstable neurons, with the goal of limiting the propagation of approximation errors after they are introduced. In the running example, since both $n_{1,1}$ and $n_{1,1}'$ are in unstable states, i.e., the input intervals of the ReLUs contain the value 0, we may introduce a new symbol $x_3 = ReLU(n_{1,1}') - ReLU(n_{1,1})$. In all subsequent layers, whenever the value of $ReLU(n_{1,1}') - ReLU(n_{1,1})$ is needed, we use the bounds $[x_3, x_3]$ instead of the actual bounds.
The reason why using $x_3$ can lead to more accurate results is because, even though our convex approximations reduce the error introduced, there is inevitably some error that accumulates. Introducing $x_3$ allows this error to partially cancel in the subsequent layers. In our running example, introducing the new symbolic variable $x_3$ would be able to improve the final bounds to $[-1.65, 1.18]$. While creating $x_3$ improved the result in this case, carelessly introducing new variables for all the unstable neurons can actually reduce the overall benefit (see Section 4). In addition, the computational cost of introducing new variables is not negligible. Therefore, in practice, we must introduce these symbolic variables judiciously, to maximize the benefit. Part of our contribution in NeuroDiff is in developing heuristics to automatically determine when to create new symbolic variables (details in Section 4).

3 BACKGROUND

In this section, we review the technical background and then introduce notations that we use throughout the paper.

3.1 Neural Networks

We focus on feed-forward neural networks, which we define as a function $f$ that takes an $n$-dimensional vector of real values $x \in \mathbb{X}$, where $\mathbb{X} \subseteq \mathbb{R}^n$, and maps it to an $m$-dimensional vector $y \in \mathbb{Y}$, where $\mathbb{Y} \subseteq \mathbb{R}^m$. We denote this function as $f : \mathbb{X} \rightarrow \mathbb{Y}$. Typically, each dimension of $y$ represents a score, such as a probability, that the input $x$ belongs to class $i$, where $1 \leq i \leq m$.

A network with $l$ layers has $l$ weight matrices, each of which is denoted $W_k$, for $1 \leq k \leq l$. For each weight matrix, we have $W_k \in \mathbb{R}^{l_{k-1} \times l_k}$ where $l_{k-1}$ is the number of neurons in layer $(k-1)$ and likewise for $l_k$, and $l_0 = n$. Each element in $W_k$ represents the weight of an edge from a neuron in layer $(k-1)$ to one in layer $k$. Let $n_{k,j}$ denote the $j^{th}$ neuron of layer $k$, and $n_{k-1,i}$ denote the $i^{th}$ neuron of layer $(k-1)$. We use $W_k[i,j]$ to denote the edge weight from $n_{k-1,i}$ to $n_{k,j}$. In our motivating example, we have $W_1[1,1] = 1.9$ and $W_1[1,2] = 1.1$.

Mathematically, the entire neural network can be represented by $f(x) = f_l(W_l \cdot f_{l-1}(W_{l-1} \cdot \ldots f_1(W_1 \cdot x) \ldots))$, where $f_k$ is the activation function of the $k^{th}$ layer and $1 \leq k \leq l$. We focus on neural networks with ReLU activations because they are the most widely implemented in practice, but our method can be extended to other activation functions, such as sigmoid and tanh, and other layer types, such as convolutional and max-pooling. We leave this as future work.

3.2 Symbolic Intervals

To compute approximations of the output nodes that are sound for all input values, we leverage interval arithmetic [27], which can be viewed as an instance of the abstract interpretation framework [5]. It is well-suited to the verification task because interval arithmetic...
is soundly defined for basic operations of the network such as addition, subtraction, and scaling.

Let $I = [LB(I), UB(I)]$ be an interval with lower bound $LB(I)$ and upper bound $UB(I)$. Then, for intervals $I_1, I_2$, we have addition and subtraction defined as $I_1 + I_2 = [LB(I_1) + LB(I_2), UB(I_1) + UB(I_2)]$ and $I_1 - I_2 = [LB(I_1) - UB(I_2), UB(I_1) - LB(I_2)]$, respectively. For a constant $c$, scaling is defined as $c \times I = [c \times LB(I), c \times UB(I)]$ when $c > 0$, and $c \times I = [c \times UB(I), c \times LB(I)]$ otherwise.

While interval arithmetic is a sound over-approximation, it is not always accurate. To illustrate, let $f(x) = 3x - x$, and say we are interested in bounding $f(x)$ when $x \in [-1, 1]$. One way to bound $f$ is by evaluating $f(I)$ where $I = [-1, 1]$. Doing so yields $3 \times [-1, 1] - [-1, 1] = [-4, 4]$. Unfortunately, the most accurate bounds are $[-2, 2]$.

There are (at least) two ways we can improve the accuracy. First, we can soundly refine the result by dividing the input intervals into disjoint partitions, performing the analysis independently on each partition, and then unioning the resulting output intervals together. Previous work has shown the result will be at least as precise [48], and often better. For example, if we partition $x \in [-1, 1]$ into $x \in [-1, 0]$ and $x \in [0, 1]$, and perform the analysis for each partition, the resulting bounds improve to $[-3, 3]$.

Second, the dependence between the two intervals are not leveraged when we subtract them, i.e., that they were both $x$ terms and hence could partially cancel out. To capture the dependence, we can use symbolic lower and upper bounds [48], which are expressions in terms of the input variable, i.e., $I = [x, x]$. Evaluating $f(I)$ then yields the interval $f_I = [2x, 2x]$, for $x \in [-1, 1]$. When using symbolic bounds, eventually, we must concretize the lower and upper bound equations. We denote concretization of $LB(I_f) = 2x$ and $UB(I_f) = 2x$ as $LB(I_f) = -2$ and $UB(I_f) = 2$, respectively. Compared to the naive solution, $[-4, 4]$, this is a significant improvement.

When approximating the output of a given function $f : X \rightarrow Y$ over an input interval $X \subseteq X$, one may prove soundness by showing that the evaluation of the lower and upper bounds on any input $x \in X$ are always greater than and less than, respectively, to the true value of $f(x)$. Formally, for an interval $I$, let $LB(I)(x)$ be the evaluation of the lower bound equation on input $x$, and similarly for $UB(I)(x)$. Then, the approximation is considered sound if for all $x \in X$, we have $LB(I)(x) \leq f(x) \leq UB(I)(x)$.

### 3.3 Convex Approximations

While symbolic intervals are exact for linear operations (i.e. they do not introduce error), this is not the case for non-linear operations, such as the ReLU activation. This is because, for efficiency reasons, the symbolic lower and upper bounds must be kept linear. Thus, developing linear approximations for non-linear activation functions has become a significant area of research for single neural network verification [40, 47, 49, 54]. We review the basics below, but caution that they are different from our new convex approximations in NeurD iff.

We denote the input to the ReLU of a neuron $n_{k,j}$ as $S^{in}(n_{k,j})$ and the output as $S(n_{k,j})$. The approach used by existing single-network verification tools is to apply an affine transformation to the upper bound of $S^{in}(n_{k,j})$ such that $UB(S^{in}(n_{k,j}))(x) \geq 0$, where $x \in X$, and $X$ is the input region for the entire network. For the lower bound, there exist several possible transformations, including the one used by Neurify [47], shown in Figure 6, where $n = S^{in}(n_{k,j})$ and the dashed lines are the upper and lower bounds.

We illustrate the upper bound transformation for $n_{1,1}$ of our motivating example. After computing the upper bound of the ReLU input $UB(S^{in}(n_{1,1})) = 1.9x_1 - 1.9x_2$, where $x_1 \in [-2, 2]$ and $x_2 \in [-2, 2]$, it computes the concrete lower and upper bounds. We denote these as $LB(S^{in}(n_{1,1})) = -7.6$ and $UB(S^{in}(n_{1,1})) = 7.6$. We refer to them as $I$ and $u$, respectively, for short hand. Then, it computes the line that passes through $(u, u)$ and $(0, l)$. Letting $y = UB(S^{in}(n_{1,1}))$ be the upper bound equation of the ReLU input, it computes the upper bound of the ReLU output as $UB(S(n_{1,1})) = \frac{u}{l}(y - l) = 0.95x_1 - 0.95x_2 + 3.81$.

When considering a single ReLU of a single network, convex approximation is simple because there are only three states that the neuron can be in, namely active, inactive, and unstable. Furthermore, in only one of these states, convex approximation is needed. In contrast, differential verification has to consider a pair of neurons, which has up to nine states to consider between the two ReLUs. Furthermore, different states may result in different linear approximations, and some states can even have multiple linear approximations depending on the difference bound of $\Delta = n' - n$.

As we will show in Section 4, there are significantly more considerations in our problem domain.

### 4 OUR APPROACH

We first present our baseline procedure for differential verification of feed-forward neural networks (Section 4.1), and then present our algorithms for computing convex approximations (Section 4.3) and introducing symbolic variables (Section 4.4).

#### 4.1 Differential Verification – Baseline

We build off the work of Paulsen et al. [33], so in this section we review the relevant pieces. We assume that the input to NeurD iff consists of two networks $f$ and $f'$, each with $l$ layers of the same size. Let $n'_{k,j}$ in $f'$ be the neuron paired with $n_{k,j}$ in $f$. This implicitly creates a pairing of the edge weights between the two networks. We first introduce additional notation.

- We denote the difference between a pair of neurons as $\Delta_{k,j} = n'_{k,j} - n_{k,j}$. For example, $\Delta_{1,1} = 0.1$ under the input $x_1 = 2, x_2 = 1$ in our motivating example shown in Figure 2.
- We denote the difference in a pair of edge weights as $W^{in}_{k,j} = W^{in}_{k,j}$. For example, $W^{in}_{k,j} = 2.0 - 1.9 = 0.1$.
- We extend the symbolic interval notation to these terms. That is, $S^{in}(\Delta_{k,j})$ denotes the interval that bounds $n'_{k,j} - n_{k,j}$ before applying ReLU, and $S(\Delta_{k,j})$ denotes the interval after applying ReLU.

Given that we have computed $S(n_{k-1,j})$, $S(n'_{k-1,j})$, $S(\Delta_{k-1,j})$ for every neuron in the layer $k - 1$, now, we compute a single $S(\Delta_{k,j})$ in the subsequent layer $k$ in two steps (and then repeat for each $1 \leq j \leq l_k$).

First, we compute $S^{in}(\Delta_{k,j})$ by propagating the output intervals from the previous layer through the edges connecting to the target
4.3 New Convex Approximations for $S(\Delta_{k,j})$

Now, we are ready to present our new approximations, which are linear symbolic expressions derived from Equations 1 and 2.

We first assume that $n_{k,j}$ and $n'_{k,j}$ could both be unstable, i.e., they could take values both greater than and less than 0. This yields bounds for the general case in that they are sound in all states of $n_{k,j}$ and $n'_{k,j}$ (Sections 4.3.1 and 4.3.2). Then, we consider special cases of $n_{k,j}$ and $n'_{k,j}$, in which even tighter upper and lower bounds are derived (Section 4.3.3).

To simplify notation, we let $n, n'$, and $\Delta$ stand in for $n_{k,j}, n'_{k,j}$, and $\Delta_{k,j}$ in the remainder of this section.

4.3.1 Upper Bound for the General Case. Let $l = \overline{UB}(S^{in}(\Lambda))$ and $u = \overline{UB}(S^{in}(\Lambda))$. The upper bound approximation is:

$$\overline{UB}(S(\Lambda)) = \begin{cases} \overline{UB}(S^{in}(\Lambda)) & \text{if } \overline{UB}(S^{in}(\Lambda)) \geq 0 \\ \overline{UB}(S^{in}(\Lambda)) & \text{if } \overline{UB}(S^{in}(\Lambda)) < 0 \\ \overline{UB}(S^{in}(\Lambda)) & \text{otherwise} \end{cases}$$

That is, when the input’s (delta) upper bound is greater than 0 for all $x \in X$, we can use the input’s upper bound unchanged. When the upper bound is always less than 0, the new output’s upper bound is then 0. Otherwise, we apply a linear transformation to the upper bound, which results in the upper plane illustrated in Figure 5. We prove all three cases sound.

Proof. We consider each case above separately. In the following, we use Equation 1 to derive the bounds, but we note a symmetric proof using Equation 2 exists and produces the same bounds.

Case 1: $\overline{UB}(S^{in}(\Lambda)) \geq 0$. We first show that, according to Equation 1, when $0 \leq \Delta$ we have $z_{1} \leq \Delta$. This then implies that, if $\overline{UB}(S^{in}(\Lambda)) \geq 0$, then $z_{1} \leq \overline{UB}(S^{in}(\Lambda))(x)$ for all $x \in X$, and hence it is a valid upper bound for the output interval.

Assume $0 \leq \Delta$. We consider two cases of $n$. First, consider $0 \leq n$. Observe $0 \leq n \wedge 0 \leq \Delta \implies 0 \leq n \wedge \Delta$. Thus, the ReLU’s of Equation 1 simplify to $z_{1} = n + \Delta - n = \Delta \implies z_{1} \leq \Delta$. When $n < 0$, Equation 1 simplifies to $z_{1} = ReLU(n + \Delta)$. Since $n < 0$, we have $n + \Delta \leq \Delta \leq 0 \implies ReLU(n + \Delta) \leq \Delta$. Thus, $z_{1} = ReLU(n + \Delta) \leq \Delta$, so the approximation is sound.

Case 2: $\overline{UB}(S^{in}(\Lambda)) < 0$. This case was previously proven [33], but we restate it here. $\overline{UB}(S^{in}(\Lambda)) \leq 0 \implies n' \leq n \implies ReLU(n') \leq ReLU(n) \implies ReLU(n') \leq \overline{UB}(n) \leq 0$.

Case 3. By case 1, any $\overline{UB}(S(\Lambda))$ that satisfies $\overline{UB}(S(\Lambda))(x) \geq 0$ and $\overline{UB}(S(\Lambda))(x) \geq \overline{UB}(S^{in}(\Lambda))(x)$ for all $x \in X$ is sound. Both inequalities hold by Lemma 4.1, with $x = UB(S^{in}(\Lambda)), l = UB(S^{in}(\Lambda)), u = UB(S^{in}(\Lambda))$, and $l' = 0$.

We illustrate the upper bound computation on node $n_{1,1}$ of our motivating example. Recall that $UB(S^{in}(n_{1,1})) = 0.1x_{1} - 0.1x_{2}$. Since $UB(S^{in}(n_{1,1})) = -0.4$ and $UB(S^{in}(n_{1,1})) = 0.4$, we are in the third case of our linear approximation above. Thus, we have $UB(S^{in}(n_{1,1})) = (0.1x_{1} - 0.1x_{2} - (-0.4)) \frac{0.4}{(-0.4)} = 0.5x_{1} - 0.5x_{2} + 0.2$. This is the upper bounding plane illustrated in Figure 5. The volume under this plane is 50% less than the upper bounding plane of ReLU_DIFF shown in Figure 4.
4.3.2 Lower Bound for the General Case. Let \( l = \overline{B}(S_{in}(\Delta)) \) and \( u = \overline{B}(S_{in}(\Delta)) \), the lower bound approximation is:

\[
\text{LB}(S(\Delta)) = \begin{cases} 
\overline{B}(S_{in}(\Delta)) & \overline{B}(S_{in}(\Delta)) \leq 0 \\
0 & \overline{B}(S_{in}(\Delta)) > 0 \\
(\overline{B}(S_{in}(\Delta)) - u) \cdot \frac{-1}{w_{1}} & \text{otherwise}
\end{cases}
\]

That is, when the input lower bound is always less than 0, we can leave it unchanged. When it is always greater than 0, the new lower bound is then 0. Otherwise, we apply a transformation to the lower bound, which results in the lower plane illustrated in Figure 5. We prove all three cases sound.

Proof. We consider each case above separately. In the following, we use Equation 1 to derive the bounds, but we note a symmetric proof using Equation 2 exists and produces the same bounds.

Case 1: \( \overline{B}(S_{in}(\Delta)) \leq 0 \). We first show that according to Equation 1, when \( \Delta \leq 0 \) we have \( \Delta \leq z_1 \). This then implies that, if \( \overline{B}(S_{in}(\Delta)) \leq 0 \), we have \( \overline{B}(S_{in}(\Delta))(x) \leq z_1 \) for all \( x \in X \), and hence it is a valid lower bound for the output interval.

Assume \( \Delta \leq 0 \). We consider two cases of \( n + \Delta \). First, let \( 0 \leq n + \Delta \). Observe that \( 0 \leq n + \Delta \) \( \land \Delta \leq 0 \) \( \iff \) \( 0 \leq n \), so we can simplify Equation 1 to \( z_1 = n + \Delta - n = \Delta \leq z_1 \). Second, let \( n + \Delta < 0 \) \( \iff \) \( \Delta < -n \). Then, Equation 1 simplifies to \( z_1 = -\text{ReLU}(n) = -\max(0, n) = \min(0, -n) \). Now observe \( \Delta < -n \) \( \land \Delta < 0 \) \( \iff \) \( \Delta < \min(0, -n) = z_1 \).

Case 2: \( \overline{B}(S_{in}(\Delta)) > 0 \). This case was previously proven sound [33], but we restate it here. \( \overline{B}(S_{in}(\Delta)) \geq 0 \iff n' \geq n \iff \text{ReLU}(n') \geq \text{ReLU}(n) \iff \text{ReLU}(n') - \text{ReLU}(n) \geq 0 \).

Case 3. By case 1, any \( \overline{B}(S(\Delta)) \) that satisfies \( \overline{B}(S(\Delta))(x) \leq 0 \) and \( \overline{B}(S(\Delta))(x) \leq \overline{B}(S_{in}(\Delta))(x) \) for all \( x \in X \) will be valid. Both inequalities hold by Lemma 4.2, with \( x = \overline{B}(S_{in}(\Delta)), u' = 0, l = \overline{B}(S_{in}(\Delta)) \), and \( u = \overline{B}(S_{in}(\Delta)) \).

We illustrate the lower bound computation on node \( n_{1,1} \) of our motivating example. Recall that \( \overline{B}(S_{in}(n_{1,1})) = 0.1x_1 - 0.1x_2 \). Since \( \overline{B}(S_{in}(n_{1,1})) = 0.4 \) and \( \overline{B}(S_{in}(n_{1,1})) = 0.4 \), we are in the third case of our linear approximation. Thus, we have \( \overline{B}(S_{in}(n_{1,1})) = 0.1x_1 - 0.1x_2 - (0.4) \cdot \frac{-0.4}{0.4 + 0.4} = 0.05x_1 - 0.05x_2 - 0.2 \). This is the lower bounding plane illustrated in Figure 5. The volume above this plane is 50% less than the lower bounding plane of \( \text{ReLU}_{\Delta} \) shown in Figure 4.

4.3.3 Tighter Bounds for Special Cases. While the bounds presented so far apply in all states of \( n \) and \( n' \), under certain conditions, we are able to tighten these bounds even further. Toward this end, we restate the following two lemmas proved by Paulsen et al. [33], which will come in handy. They are related to properties of Equations 1 and 2, respectively.

Lemma 4.3. \( \text{ReLU}(n + \Delta) - n = \max(-n, \Delta) \)

Lemma 4.4. \( n' - \text{ReLU}(n' - \Delta) = \min(n', \Delta) \)

These lemmas provide bounds when \( n \) and \( n' \) are proved to be linear based on the absolute bounds that we compute.

Figure 8: Tighter upper bounding plane.

Figure 9: Tighter lower bounding plane.

Tighter Upper Bound When \( n' \) Is Linear. In this case, we have \( U_B(S(\Delta)) = \overline{B}(S_{in}(\Delta)) \), which is an improvement for the second or third case of our general upper bound.

Proof. By our case assumption, Equation 2 simplifies to the one in Lemma 4.4. Thus, \( z_2 = \min(n', \Delta) \iff z_2 \leq \Delta \).

Tighter Upper Bound When \( n \) Is Linear. \( \overline{B}(S_{in}(\Delta)) \leq \overline{B}(S_{in}(\Delta)) \leq \overline{B}(S_{in}(\Delta)) \leq \overline{B}(S_{in}(\Delta)) \). We illustrate the \( z_1 \) plane under these constraints in Figure 8. Let \( l = \overline{B}(S_{in}(\Delta)) \), and let \( u = \overline{B}(S_{in}(\Delta)) \), and \( l' = -\overline{B}(S_{in}(\Delta)) \), we use Lemma 4.1 to derive \( U_B(S(\Delta)) = (\overline{B}(S_{in}(\Delta)) - l) \ast \frac{u - l'}{u_{1'}} + l' \). This results in the upper plane of Figure 8. This improves over the third case in our general upper bound because it allows the lower bound of \( U_B(S(\Delta)) \) to be less than 0.

Proof. By our case assumption, Equation 1 simplifies to the one in Lemma 4.3. By Lemma 4.1, we have for all \( x \in X \), \( U_B(S(\Delta))(x) \geq -\overline{B}(S_{in}(\Delta)) \) and \( U_B(S(\Delta))(x) \geq \overline{B}(S_{in}(\Delta))(x) \). These two inequalities imply \( U_B(S(\Delta)) \geq \max(-n, \Delta) \).

Tighter Lower Bound When \( n \) Is Linear. Here, we can use the approximation \( \text{LB}(S(\Delta)) = \overline{B}(S_{in}(\Delta)) \). This improves over the second and third cases of our general lower bound.

Proof. By our case assumption, Equation 1 simplifies to the one in Lemma 4.3. Thus, \( z_1 = \max(-n, \Delta) \iff z_1 \geq \Delta \).

Tighter Lower Bound when \( n' \) Is Linear. \( \text{LB}(S_{in}(\Delta)) \leq \text{LB}(S_{in}(\Delta')) \leq \text{LB}(S_{in}(\Delta)) \). We illustrate the \( z_2 \) plane under these constraints in Figure 9. Here, letting \( l = \text{LB}(S_{in}(\Delta)), u = \text{LB}(S_{in}(\Delta)), \) and \( u' = \text{LB}(S_{in}(\Delta')) \), we can use Lemma 4.2 to derive the approximation \( \text{LB}(S(\Delta)) = (\text{LB}(S_{in}(\Delta)) - u) \ast \frac{u - u'}{u_{1'}} + u' \). This results in the lower plane of Figure 9. This improves over the third case, since it allows the upper bound to be greater than 0.
Proof. By our case assumption, Equation 2 simplifies to the one shown in Lemma 4.4. By Lemma 4.2, we have for all \( x \in X \), \( \text{LB}(S(\Delta))(x) \leq \text{LB}(S^{n}(\Delta))(x) \) and \( \text{LB}(S(\Delta))(x) \leq \text{LB}(S^{n}(n')) \). These two inequalities imply \( \text{LB}(S(\Delta))(x) \leq \min(n', \Delta). \) \( \Box \)

4.4 Intermediate Symbolic Variables for \( S(\Delta) \)

While convex approximations reduce the error introduced by ReLU, even small errors tend to be amplified significantly after a few layers. To combat the error explosion, we introduce new symbolic terms to represent the output values of unstable neurons, which allow their accumulated errors to cancel out.

We illustrate the impact of symbolic variables on \( n_{3,1} \) of our motivating example. Recall we have \( S(\Delta_{1,1}) = [0.05x_{1} - 0.05x_{2} - 0.2, 0.05x_{1} - 0.05x_{2} + 0.2] \). After applying the convex approximation, we introduce a new variable \( x_{3} \) such that \( x_{3} = [0.05x_{1} - 0.05x_{2} - 0.2, 0.05x_{1} - 0.05x_{2} + 0.2] \). Then we set \( S(\Delta_{2,1}) = [x_{3}, x_{3}] \), and propagate this interval as before. After propagating through \( n_{2,1} \) and \( n_{2,2} \) and combining them at \( n_{3,1} \), the \( x_{3} \) terms partially cancel out, resulting in the tighter final output interval \([-1.65, 1.18]\).

In principle, symbolic variables may be introduced at any unstable neurons that introduce approximation errors, however there are efficiency vs. accuracy tradeoffs when introducing these symbolic variables. One consideration is how to deal with intermediate variables referencing other intermediate variables. For example, if we decide to introduce a variable \( x_{4} \) for \( n_{2,1} \), then \( x_{4} \) will have an \( x_{3} \) term in its equation. Then, when we are evaluating a symbolic bound that contains an \( x_{4} \) term, which will be the case for \( n_{3,1} \), we will have to recursively substitute the bounds of the previous intermediate variables, such as \( x_{3} \). This becomes expensive, especially when it is used together with our bisection-based refinement [33, 48]. Thus, in practice, we first remove any back-references to intermediate variables by substituting in their lower bounds and upper bounds into the new intermediate variable’s lower and upper bounds, respectively.

Given that we do not allow back-references, there are two additional considerations. First, we must consider that introducing a new intermediate variable wipes out all the other intermediate variables. For example, introducing a new variable at \( n_{2,1} \) wipes out references to \( x_{3} \), thus preventing any \( x_{3} \) terms from canceling at \( n_{3,1} \). Second, the runtime cost of introducing symbolic variables is not negligible. The bulk of computation time in NeuroDiff is spent multiplying the neuron’s weight matrices by the neuron’s symbolic bound equations, which is implemented using matrix multiplication. Since adding variables increases the matrix size, this increases the matrix multiplication cost.

Based on these considerations, we have developed heuristics for adding new variables judiciously. First, since the errors introduced by unstable neurons in the earliest layers are the most prone to explode, and hence benefit the most when we create variables for them, we rank them higher when choosing where to add symbolic variables. Second, we bound the total number of symbolic variables that may be added, since our experience shows that introducing symbolic variables for the earliest \( N \) unstable neurons gives drastic improvements in both run time and accuracy. In practice, \( N \) is set to a number proportional to the weighted sum of unstable neurons in all layers. Formally, \( N = \sum_{k=1}^{K} y^{k} \times N_{k} \), where \( N_{k} \) is the number of unstable neurons in layer \( k \) and \( y^{k} = \frac{1}{k} \) is the discount factor.

5 EXPERIMENTS

We have implemented NeuroDiff and compared it with ReluDiff [33], the state-of-the-art tool for differential verification of neural networks. NeuroDiff builds upon the codebase of ReluDiff [32], which was also used by single-network verification tools such as ReluVal [48] and Neurify [47]. All use OpenBLAS [55] to optimize the symbolic interval arithmetic (namely in applying the weight matrices to the symbolic intervals). We note that NeuroDiff uses the algorithm from Neurify to compute \( S(n_{k,j}) \) and \( S(n'_{k,j}) \), whereas ReluDiff uses the algorithm of ReluVal. Since Neurify is known to compute tighter bounds than ReluVal [47], we compare to both ReluDiff, and an upgraded version of ReluDiff which uses the bounds from Neurify to ensure that any performance gain is due to our optimizations and not due to using Neurify’s bounds. We use the name ReluDiff+ to refer to ReluDiff upgraded with Neurify’s bounds.

5.1 Benchmarks

Our benchmarks consist of the 49 feed-forward neural networks used by Paulsen et al. [33], taken from three applications: aircraft collision avoidance, image classification, and human activity recognition. We briefly describe them here. As in Paulsen et al. [33], the second network \( f' \) is generated by truncating the edge weights of \( f \) from 32 bit to 16 bit floats.

ACAS Xu [16]. ACAS (aircraft collision avoidance system) Xu is a set of forty-five neural networks, each with five inputs, six hidden layers of 50 units each, and five outputs, designed to advise a pilot (the ownship) how to steer an aircraft in the presence of an intruder aircraft. The inputs describe the position and speed of the intruder relative to the ownship, and the outputs represent scores for different actions that the ownship should take. The scores range from \([-0.5, 0.5]\). We use the input regions defined by the properties of previous work [17, 48].

MNIST [21]. MNIST is a standard image classification task, where the goal is to correctly classify 28 x 28 pixel greyscale images of handwritten digits. Neural networks trained for this task take 784 inputs (one for each pixel) each in the range [0, 255], and compute ten outputs — one score for each of the ten possible digits. We use three networks of size 3x100 (three hidden layers of 100 neurons each), 2x512, and 4x1024 taken from Weng et al. [49] and Wang et al. [47]. All achieve at least 95% accuracy on holdout test data.

Human Activity Recognition (HAR) [1]. The goal for this task is to classify the current activity of human (e.g. walking, sitting, laying down) based on statistics from a smartphone’s gyroscopeic sensors. Networks trained on this task take 561 statistics computed from the sensors and output six scores for six different activities. We use a 1x500 network.

5.2 Experimental Setup

Our experiments aim to answer the following research questions:

1. Is NeuroDiff significantly faster than state-of-the-art?
Thus, to save space, we will only show the results that compare variables, the differences are on the order of seconds. Since they are slower than scale the speedups of points above the diagonal line are wins for \( \text{iff}^+ \) in seconds. Thus, points on the diagonal line are ties, while in seconds, and the y-axis the execution time of NeuroDiff a differential verification task: the x-axis is the execution time of \( \text{iff}^+ \) using a scatter plot in Figure 10, where each point represents (our method) and \( \text{ReluVal} \) instead of \( \text{ReluDiff} \). The results show that, on all benchmarks, the improved improvement?

To answer these questions, we run both \( \text{NeuroDiff} \) and \( \text{ReluD}-+ \) on all benchmarks and compare their results. Both \( \text{NeuroDiff} \) and \( \text{ReluD}-/\text{ReluD}+ \) can be parallelized to use multithreading, so we configure a maximum of 12 threads for all experiments. Our experiments are run on a computer with an AMD Ryzen Threadripper 2950X 16-core processor, with a 30-minute timeout per differential verification task.

While we could try and adapt a single-network verification tool to our task as done previously [33], we note that \( \text{ReluD}+ \) has been shown to significantly outperform (by several orders of magnitude) this naive approach.

### 5.3 Results

In the remainder of this section, we present our experimental results in two steps. First, we present the overall verification results on all benchmarks. Then, we focus on the detailed verification results on the more difficult verification tasks.

#### 5.3.1 Summary of Results on All Benchmarks

Our experimental results show that, on all benchmarks, the improved \( \text{ReluD}+ \) slightly but consistently outperforms the original \( \text{ReluD} \) due to its use of the more accurate component from \( \text{Neurify} \) instead of \( \text{ReluVal} \) for bounding the absolute values of individual neurons. Thus, to save space, we will only show the results that compare \( \text{NeuroDiff} \) (our method) and \( \text{ReluD}+ \).

We summarize the comparison between \( \text{NeuroDiff} \) and \( \text{ReluD}+ \) using a scatter plot in Figure 10, where each point represents a differential verification task: the x-axis is the execution time of \( \text{NeuroDiff} \) in seconds, and the y-axis the execution time of \( \text{ReluD}+ \) in seconds. Thus, points on the diagonal line are ties, while points above the diagonal line are wins for \( \text{NeuroDiff} \).

The results show that \( \text{NeuroDiff} \) outperformed \( \text{ReluD}+ \) for most verification tasks. Since the execution time is in logarithmic scale the speedups of \( \text{NeuroDiff} \) are more than 1000X for many of these verification tasks. While there are cases where \( \text{NeuroDiff} \) is slower than \( \text{ReluD}+ \), due to the overhead of adding symbolic variables, the differences are on the order of seconds. Since they are all on the small MNIST networks and the HAR network that are very easy for both tools, we omit an in-depth analysis of them.

In the remainder of this section, we present an in-depth analysis of the more difficult verification tasks.

#### 5.3.2 Results on ACAS Networks

For ACAS networks, we consider two different sets of properties, namely the original properties from Paulsen et al. [33] where \( \epsilon = 0.05 \), and the same properties but with \( \epsilon = 0.01 \). We emphasize that, while verifying \( \epsilon = 0.05 \) is useful, this means that the output value can vary by up to 10%. Considering \( \epsilon = 0.01 \) means that the output value can vary by up to 2%, which is much more useful.

Our results are shown in Tables 1 and 2, where the first column shows the property, which defines the input space considered. The next three columns show the results for \( \text{NeuroDiff} \), specifically the number of verified networks (out of the 45 networks), the number of unverified networks, and the total run time across all networks. The next three show the same results, but for \( \text{ReluD}+ \). The final column shows the average speedup of \( \text{NeuroDiff} \).

The results show that \( \text{NeuroDiff} \) makes significant gains in both speed and accuracy. Specifically, the speedups are up to two and three orders of magnitude for \( \epsilon = 0.05 \) and 0.01, respectively. In addition, at the more accurate \( \epsilon = 0.01 \) level, \( \text{NeuroDiff} \) is able to complete 53 more verification tasks, out of the total 142 verification tasks.

**Table 1: Results for ACAS networks with \( \epsilon = 0.05 \).**

| Property | NeuroDiff (new) | ReluD-/+ | Speedup |
|----------|----------------|----------|---------|
|          | proved | undet. | time (s) | proved | undet. | time (s) |         |
| \( p_1 \) | 45     | 0      | 322.6    | 44     | 0      | 4800.6  | 9.2     |
| \( p_2 \) | 42     | 0      | 2.3      | 42     | 0      | 4.1     | 1.8     |
| \( p_3 \) | 42     | 0      | 1.7      | 42     | 0      | 2.8     | 1.7     |
| \( p_4 \) | 1      | 0      | 0.2      | 1      | 0      | 0.2     | 1.4     |
| \( p_5 \) | 2      | 0      | 0.6      | 2      | 0      | 2.4     | 0.7     |
| \( p_6 \) | 0      | 1      | 1400.4   | 0      | 1      | 1800.0  | 1.3     |
| \( p_7 \) | 1      | 0      | 132.2    | 1      | 0      | 361.8   | 2.7     |
| \( p_8 \) | 1      | 0      | 0.6      | 1      | 0      | 2.3     | 3.7     |
| \( p_9 \) | 1      | 0      | 0.9      | 1      | 0      | 7.1     | 8.0     |
| \( p_{10} \) | 1      | 0      | 0.2      | 1      | 0      | 0.3     | 1.6     |
| \( p_{11} \) | 1      | 0      | 2.8      | 1      | 0      | 360.9   | 129.4   |
| \( p_{12} \) | 1      | 0      | 5.8      | 1      | 0      | 5.1     | 0.9     |
| \( p_{13} \) | 2      | 0      | 0.3      | 2      | 0      | 93.9    | 196.2   |
| \( p_{14} \) | 2      | 0      | 0.6      | 2      | 0      | 65.0    | 113.2   |

**Table 2: Results for ACAS networks with \( \epsilon = 0.01 \).**

| Property | NeuroDiff (new) | ReluD-/+ | Speedup |
|----------|----------------|----------|---------|
|          | proved | undet. | time (s) | proved | undet. | time (s) |         |
| \( p_1 \) | 41     | 4      | 11400.1  | 15     | 30     | 557.86  | 4.9     |
| \( p_2 \) | 42     | 0      | 14.3     | 35     | 7      | 13642.2 | 957.2   |
| \( p_3 \) | 42     | 0      | 3.8      | 37     | 5      | 9115.0  | 2398.1  |
| \( p_4 \) | 0      | 0      | 0.3      | 0      | 1      | 1800.0  | 5528.5  |
| \( p_{15} \) | 2      | 0      | 1.0      | 2      | 0      | 0.8     | 0.8     |
| \( p_{16} \) | 0      | 1      | 1800.0   | 0      | 1      | 1800.0  | 1.0     |
| \( p_{17} \) | 1      | 0      | 1115.9   | 0      | 1      | 1800.0  | 1.6     |
| \( p_{18} \) | 1      | 0      | 2.8      | 0      | 1      | 1800.0  | 738.2   |
| \( p_{19} \) | 1      | 0      | 1.6      | 1      | 0      | 1.1     | 0.7     |
| \( p_{20} \) | 1      | 0      | 0.3      | 0      | 1      | 1800.0  | 5673.8  |
| \( p_{21} \) | 1      | 0      | 132.2    | 0      | 1      | 1800.0  | 13.6    |
| \( p_{22} \) | 1      | 0      | 15.9     | 1      | 0      | 14.8    | 0.9     |
| \( p_{23} \) | 2      | 0      | 1289.3   | 0      | 2      | 3600.0  | 2.3     |
| \( p_{24} \) | 2      | 0      | 579.4    | 2      | 0      | 3600.0  | 6.2     |
5.3.3 Results on MNIST Networks. For MNIST, we focus on the 4x1024 network, which is the largest network considered by Paulsen et al. [33]. In contrast, since the smaller networks, namely 3x100 and 2x512 networks, were handled easily by both tools, we omit their results. In the MNIST-related verification tasks, the goal is to verify $\epsilon = 1$ for the given input region. We consider the two types of input regions from the previous work, namely global perturbations and targeted pixel perturbations, however we use input regions that are hundreds of orders of magnitude larger.

First, we look at the global perturbation. For these, the input space is created by taking an input image and then allowing a perturbation of +/- $p$ greyscale units to all of its pixels. In the previous work, the largest perturbation was $p = 3$. Figure 11 compares NeuroDiff and ReluDiff+ on $p = 3$ all the way up to 8, where the x-axis is the perturbation applied, and the y-axis is the percentage of verification tasks (out of 100) that each can handle.

The results show that NeuroDiff can handle perturbations up to +/- 6 units, whereas ReluDiff+ begins to struggle at 4. While the difference between 4 and 6, may seem small, the volume of input space for a perturbation of 6 is $784^6 / 784^4 \approx 1.1 \times 10^{138}$ times larger than 4, or in other words, 138 orders of magnitude larger.

Next, we show a comparison of the epsilon verified by a single forward pass for a perturbation of 8 on the MNIST 4x1024 network in Figure 12. Points above the blue line indicate NeuroDiff performed better. Overall, NeuroDiff is between two and three times more accurate than ReluDiff+.

Finally, we look at the targeted pixel perturbation properties. For these, the input space is created by taking an image, randomly choosing $n$ pixels, and setting there bounds to [0, 255], i.e., allowing arbitrary changes to the chosen pixels. We again use the 4x1024 MNIST network. The results are summarized in Table 3. The first column shows the number of randomly perturbed pixels. We can again see very large speedups, and a significant increase in the size of the input region that NeuroDiff can handle.

5.3.4 Contribution of Each Technique. Here, we analyze the contribution of individual techniques, namely convex approximations and symbolic variables, to the overall performance improvement.

In Table 4, we present the average $\epsilon$ that was able to be verified after a single forward pass on the 4x1024 MNIST network for each of the four techniques: ReluDiff+ (baseline), NeuroDiff with only convex approximations, NeuroDiff with only intermediate variables, and the full NeuroDiff.

Overall, the individual benefits of the two proposed approximation techniques are obvious. While convex approximation (alone) consistently provides benefit as perturbation increases, the benefit of symbolic variables (alone) tends to decrease. In addition, combining the two provides much greater benefit than the sum of their individual contributions. With perturbation of 8, for example, convex approximations alone are 1.59 times more accurate than ReluDiff+, and intermediate variables alone are 1.01 times more accurate. However, together they are 2.93 times more accurate.

The results suggest two things. First, intermediate symbolic variables perform well when a significant portion of the network is already in the stable state. We confirm, by manually inspecting the experimental results, that it is indeed the case when we use a perturbation of 3 and 8 in the MNIST experiments. Second, the convex approximations provide the most benefit when the pre-ReLU delta intervals are (1) significantly wide, and (2) still contain a significant
amount of symbolic information. This is also confirmed by manually inspecting our MNIST results: increasing the perturbation increases the overall width of the delta intervals.

6 RELATED WORK

Aside from ReluDiff [33], the most closely related to our work are those that focus on verifying properties of single networks as opposed to two or more networks. These verification approaches can be broadly categorized into those that use exact, constraint solving-based techniques and those that use approximations.

On the constraint solving side, several works have adapted off-the-shelf solvers [2–4, 7, 45], or even implemented solvers specifically for neural networks [17, 18]. On the approximation side, many use techniques that fit into the framework of abstract interpretation [5]. For example, many works have leveraged abstract domains such as intervals [16, 48, 49, 54], polyhedra [39, 40], and zonotopes [9, 41].

In addition, these verification techniques have also been combined [15, 41, 47], or entirely different approaches [6, 12, 38], such as bounding a network’s lipschitz constant, have been studied. These verification techniques can also be integrated into the training process to produce more robust and easier to verify networks [8, 25, 26, 50]. These works are orthogonal, though we believe their techniques can be adapted to our domain.

A related but tangential line of work focuses on discovering interesting behaviors of neural networks, though without any guarantees. Most closely related to our work are differential testing techniques [23, 34, 52], which focus on finding disagreements between a set of networks. However, these techniques do not attempt to prove the equivalence or similarity of multiple networks.

Other works are more geared towards single network testing, and use white-box testing techniques [22, 31, 42, 44, 51], such as neuron coverage statistics, to assess how well a network has been tested, and also report interesting behaviors. Both of these can be thought of as adapting software engineering techniques to machine learning.

In addition, many works use machine learning techniques, such as gradient optimization, to find interesting behaviors, such as adversarial examples [19, 24, 28, 29, 53]. These interesting behaviors can then be used to retrain the network to improve robustness [11, 36]. Again, these techniques do not provide guarantees, though we believe they could be integrated into NeuroDiff to quickly find counterexamples.

Finally, our work draws inspiration from classic software engineering techniques, such as regression testing [37], differential assertion checking [20], differential fuzzing [30], and incremental symbolic execution [13, 35], where one version of a program is used as an “oracle”, to more efficiently test or verify a new version of the same program. In our case, \( f \) can be thought of as the oracle, while \( f' \) is the new version.

7 CONCLUSIONS

We have presented NeuroDiff, a scalable differential verification technique for soundly bounding the difference between two feedforward neural networks. NeuroDiff leverages novel convex approximations, which reduce the overall approximation error, and intermediate symbolic variables, which control the error explosion, to significantly improve efficiency and accuracy of the analysis. Our experimental evaluation shows that NeuroDiff can achieve up to 1000X speedup and is up to five times as accurate.

ACKNOWLEDGMENTS

This work was partially funded by the U.S. Office of Naval Research (ONR) under the grant N00014-17-1-2896.
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