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Neural Architecture Search as Program Transformation Exploration

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ABSTRACT

Improving the performance of deep neural networks (DNNs) is important to both the compiler and neural architecture search (NAS) communities. Compilers apply program transformations in order to exploit hardware parallelism and memory hierarchy. However, legality concerns mean they fail to exploit the natural robustness of neural networks. In contrast, NAS techniques mutate networks by operations such as the grouping or bottlenecking of convolutions, exploiting the resilience of DNNs. In this work, we express such neural architecture operations as program transformations whose legality depends on a notion of representational capacity. This allows them to be combined with existing transformations into a unified optimization framework. This unification allows us to express existing NAS operations as combinations of simpler transformations. Crucially, it allows us to generate and explore new tensor convolutions. We prototyped the combined framework in TVM and were able to find optimizations across different DNNs, that significantly reduce inference time - over 3× in the majority of cases. Furthermore, our scheme dramatically reduces NAS search time. Code is available at this https url.

CCS CONCEPTS

• Computing methodologies \rightarrow Machine learning; • Software and its engineering \rightarrow Compilers.

KEYWORDS

program transformations, neural networks

ACM Reference Format:

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1 INTRODUCTION

Deep neural networks (DNNs) [39, 60] are everywhere, and there is a growing need to implement them efficiently [8, 12, 35]. This

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© 2021 Copyright held by the owner/author(s). Publication rights licensed to ACM. ACM ISBN 978-1-4503-8317-2/21/04...\$15.00 https://doi.org/10.1145/3445814.3446753 has led to an explosion in research from application [31] to hard-

Currently, there are two distinct communities optimizing DNNs for commodity devices. In one, neural architecture search (NAS) researchers explore different network models trading off size and accuracy. In the other, compiler developers take the resulting networks and explore optimizations to deliver hardware performance. We argue that this two-stage deployment process is artificial and can be unified. This paper shows that program and neural architecture transformations can be *interleaved* delivering significant performance improvement and greater expressivity. Before we describe our approach, let us briefly look at the two existing communities.

1.1 Program Transformations

Within compiler research, there has been much focus on restructuring underlying tensor computations [15, 55, 80]. This involves significant loop nest restructuring [1, 10] exploiting characteristics of the target device e.g. vectorization in SIMD units or memory coalescing in GPUs [18].

There are currently many methods [10, 57, 64, 68] in use. The polyhedral model [72, 83], in particular, is a natural fit for many standard neural network operations. A key aspect of a good program transformation representation is that sequences of transformations can be easily composed and checked for legality. This allows automatic exploration of a large space of options which can then be evaluated, and the best selected [64].

1.2 Neural Architecture Search

The neural architecture search (NAS) community is also concerned with accuracy, space, and performance [19, 76]. While there have been advances in NAS for language models [63], large scale studies of NAS tend to focus primarily on convolutional architectures [17, 81, 82]. At the neural architecture level, many techniques have been proposed for the automatic generation of networks under strict budgets [30, 65, 77, 79]. These methods focus on overall network structure, selectively replacing components such as convolutional layers with computationally cheaper methods to balance the tradeoff between accuracy and inference time.

One key strength of NAS is that it leverages the robustness of neural networks to deformation, reshaping and transforming them while incurring minimal damage to their ability to learn. This relies on networks maintaining their ability to extract feature representations for a given type of input, which we refer to as *representational capacity*. If an approximation or compression to a network does not inhibit its ability to learn from data, then it has not damaged the representational capacity of the network.

The ability of neural networks to weather compression without losing representational capacity is well-documented [16, 28, 40]. The legality of such neural architecture transformations, however, is not guaranteed and must be evaluated separately through either a training process or a small proxy task. Furthermore, while powerful, few NAS techniques explicitly take into account actual hardware behavior, and those that do face one of three problems. The first is that they only offer a black box solution with highly complex methods for predicating the search process on specific budgets (usually involving reinforcement learning [30, 65]). The second is that they do so while leaving the compilation pipeline fixed, often dismissing powerful candidate architectures because of an inappropriate, fixed choice of program transformations. The final problem is that they are limited to selecting from a pre-designed list of convolutional alternatives; they cannot synthesize their own.

1.3 Our Approach

We have two distinct communities who have the same goal but are siloed. NAS researchers assume the compiler is a black box bundled with the hardware, while compiler writers assume that the network architecture is set in stone. NAS designers can discover good networks but are limited to pre-defined options; compiler writers can efficiently exploit hardware structure but miss larger scale optimization opportunities.

What we want is the best of both worlds. We wish to combine neural architecture and compiler optimization in a unified framework. In this paper, we recast neural architecture search as program transformation exploration. By including transformations such as grouping and bottlenecking into the compiler optimization space, we leverage both the extensive history of program transformation research, and also discover new forms of neural architecture reduction that would not have been available to us otherwise.

Program transformations are necessarily restricted as they must be safe. Our solution is to unlock the space of *neural transformations* by introducing a new safety metric based on Fisher Potential [69]. It is a compile-time, cheap-to-compute metric that can reject damaging network changes, *eliminating the need to train while searching*. We, therefore, judge a transformation to be legal, not by data dependence preservation, but by the preservation of *representational capacity* (see section 5.2). This unification allows the exploration of a space that leads to more efficient implementations. It also shows that neural architecture options that previously required the engineering efforts of experts to develop, such as spatial bottlenecking, can be expressed as compositions of more fundamental transformations and discovered automatically (see section 5.3).

Our contributions are as follows:

- (1) We reformulate popular Neural Architecture Search techniques as *program transformations*.
- (2) We use Fisher Potential to provide transformation safety guarantees without the need to train.
- (3) We unify the transformation and architecture search spaces, discovering new types of convolution.
- (4) We evaluate 3 networks using these operations, ResNet, ResNext and DenseNet, on 4 platforms and demonstrate, in most cases, more than 3× inference speedup over a TVM baseline.

2 OVERVIEW

Here we develop a simple example to illustrate the direct connection between models and code, and hence, neural architecture search (NAS) and program transformation. This is followed by an outline of our approach.

2.1 Models and Code

The fundamental building block of a DNN is the tensor convolution, a generalization of a basic convolution.

Basic Convolution. Consider the first row in Figure 1. This shows a basic convolution where each element in the 2D output matrix is the weighted sum of the corresponding and neighboring elements of the 2D input matrix. The weights are stored in a small 2D filter or weight matrix W whose size corresponds to the size of the neighborhood. This is shown both diagrammatically on the left and as code on the right.

Tensor Convolution. A tensor convolution is a generalization of the basic convolution as shown in row 2 of Figure 1. The input is now a 3D tensor with a new dimension, C_i , referred to as the input channels. Similarly, the output is expanded by the number of output channels (C_o). The weight matrix is now a 4D tensor of $C_o \times C_i$ channels with two spatial dimensions (height, width). Two new loops scan the channels, the outermost, C_i shown by an arrow, while each of the tensors arrays have appropriate extra dimensions.

2.2 Models and Code Transformations

In the compiler setting, transformations change structure while maintaining meaning.

Code Transformation. Loop interchange changes the order of computation and memory access without affecting the values computed. In row 2 of Figure 1, interchanging the outer loop iterators gives the code in row 3 with the C_o and C_i loops interchanged. We can represent this using polyhedral notation $[C_i, C_o] \mapsto [C_o, C_i]$ or TVM-like syntax: .interchange(CI,CO). This is shown diagrammatically with the arrow denoting the new outermost, C_o loop order. From a neural architecture perspective nothing has changed; the number of computations and memory accesses remains the same, as do the values of the output matrix. From a code perspective, the memory access behavior is significantly altered, affecting execution time.

Model Transformation. A popular choice in NAS for reducing convolutional complexity is bottlenecking [65]. For a convolution with C_o filters, we choose a bottlenecking factor B. This reduces the size of the weight and output tensors and reduces the range of the *outermost loop iterator* by a factor B. Again this can be represented as $[C_o] \mapsto [C_o' < C_o/B]$ or .bottleneck(B). This is shown in row 4 of Figure 1. From a NAS point of view, this is a standard operation reducing complexity with a minimal effect on representational capacity. From a program transformation point of view, this is illegal as the computed values are changed.

2.3 Combined Space

If we consider swapping a full convolution for a reduced convolution as a program transformation, we can combine them. For

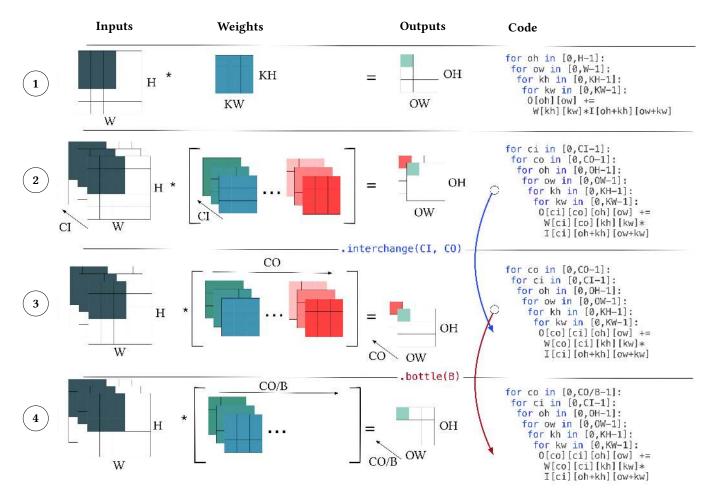


Figure 1: An illustration of the relation between neural architecture components, code and transformations. The first row shows a basic convolution. The second shows a tensor convolution. In the third row, we see a program transformation: loop interchange. The fourth row shows a neural architecture transformation: bottlenecking. Here, the number of weights is divided by B. The accuracy of the network after training does not change, but the number of operations being performed is drastically reduced.

example, we could apply loop interchange to the code in Figure 1 row 4, ($[C_o' < C_o/B, C_i] \mapsto [C_i, C_o' < C_o/B]$). With C_i now the outermost iterator, we can reapply bottlenecking, giving input channel bottlenecking. Such an optimization is both semantically invalid, and also unavailable in existing neural architecture search spaces. However, given the robustness of neural networks to noise, in some specific cases it may be just as representationally preserving as output channel bottlenecking. It is only through the lens of program transformations over loop nests that we are able to automatically access such operators.

3 NEURAL ARCHITECTURE SEARCH (NAS)

In this section we briefly describe NAS, which seeks to automate the design of neural networks for given tasks and budgets. For more details, please see [76].

Starting from an overall network skeleton, NAS attempts to design one or more cells that slot into different locations within the skeleton. Cells are described as a DAG with nodes as intermediate feature maps (or tensors) and edges representing possible operations (for example, convolution or tensor product). The task is then to find the best DAG (or cell) that can be slotted into the skeleton and trained on a given dataset. Recent work takes cells as predefined options and attempts to select where best to place each of the cells in the skeleton [69, 77]. to match specific constraints.

3.1 Neural Architectures Components

The vast majority of neural architectures consist of sequences of inter-connected components. The convolution operation dominates computational complexity and is the primary component of interest in NAS [17, 81]. We now introduce the convolution operation, and the variants that are considered as possible substitutions in our NAS baseline.

Standard Convolution. Figure 1 row 2 shows the standard convolution operation, in which an input volume of size $H_i \times W_i \times C_i$ is convolved with a set of C_0 filters of size $K_h \times K_w \times C_i$, each producing an individual feature map in the output. The convolution operation is

$$\forall c_{o}, w_{o}, h_{o} \quad O(c_{o}, w_{o}, h_{o}) = \sum_{c_{i}}^{C_{i}} \sum_{k_{h}}^{K_{h}} \sum_{k_{w}}^{K_{w}} I(c_{i}, w_{o} + k_{w}, h_{o} + k_{h}) * K(c_{o}, c_{i}, k_{w}, k_{h}).$$

$$(1)$$

Bottlenecked Convolution. A popular choice for reducing convolutional complexity is bottlenecking [29] as shown in Figure 1 row 4. A bottlenecking factor B is selected, reducing the number of weights/filters to C_o/B . ResNets [29] frequently feature bottleneck blocks that consist of trios of convolutions, one to bring the number of channels down, another for processing, and a final one to bring the number of channels back up. A similar pattern is relied on in several NAS techniques [65, 66]. The bottlenecked convolution operation is

$$\forall c'_{o} < \frac{C_{o}}{B}, w_{o}, h_{o} \quad O(c'_{o}, w_{o}, h_{o}) =$$

$$\sum_{c_{i}}^{C_{i}} \sum_{k_{h}}^{K_{h}} \sum_{k_{v}}^{K_{w}} I(c_{i}, w_{o} + k_{w}, h_{o} + k_{h}) * K(c'_{o}, c_{i}, k_{w}, k_{h}).$$
(2)

Grouped Convolution. Here, the C_i channel input is split along the channel dimension into G groups, each of which has C_i/G channels. Each group is independently convolved with its input split, producing C_o/G channels which are concatenated along the channel dimension. Let $O = [O'_1; \ldots, O'_G]$ i.e. each slice $O'_g, g \in 1, \ldots G$ is concatanated to form O. I'_g is similarly defined. Group convolution is then as follows:

$$\forall c'_{o}, w_{o}, h_{o} \quad O(c'_{o}, w_{o}, h_{o}) = \sum_{c'_{i}}^{C'_{i}} \sum_{k_{h}}^{K_{h}} \sum_{k_{w}}^{K_{w}} I(c'_{i}, w_{o} + k_{w}, h_{o} + k_{h}) * K(c'_{o}, c'_{i}, k_{w}, k_{h}).$$
(3)

This reduces the number of basic convolutions from $C_o \times C_i$ to $G \times C_o/G \times C_i/G = (C_o \times C_i)/G$ reducing the number of operations used by a factor of G. Note that we can think of standard convolution as grouped convolution with G = 1.

Depthwise Convolution. Notice that if the number of groups is equal to the number of input channels and the number of output channels, $G = C_i = C_o$, then there is a single, 2D convolutional filter for each input channel C_i . This is known as depthwise convolution and is a special case of group convolution.

3.2 NAS Search Space Example

Typically, the NAS heuristic finds the best cell to fill in the skeleton by choosing appropriate operations. Figure 2 gives an example cell from [17]. The skeleton is ResNet-like with 5 cells in series. Each cell contains 4 inter-connected nodes (A,B,C,D). Between nodes, downsampling takes place where the spatial dimensions are halved in size while doubling the channel depth. This restricts the types of operations available on each edge. This gives a total of 15625 possible cells, which captures most of the available options

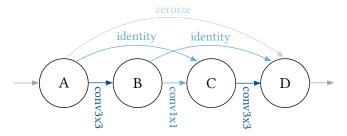


Figure 2: An illustration of a Neural Architecture Search design space (as used in [17]). Every cell has exactly four nodes (A,B,C,D) representing intermediate feature map states. Edges represent operations that transform intermediate states from source to target node, taken from a list of options specified by the designer.

within cell-based NAS techniques, each of which the authors train exhaustively on various datasets.

Rather than designing cells by choosing which of a predetermined list of options is best, we instead wish to choose a sequence of transformations from an original model which will allow us to step through the cell design space. We describe the program transformations we use in the next section.

4 PROGRAM TRANSFORMATIONS

Due to the restricted, static, convex and affine nature of tensor convolutions, it is natural to describe program transformations of them in the well-studied polyhedral model. In this section we give a brief introduction to the model; for more a detailed description, we refer the reader to [72, 75]. The polyhedral model of a program consists of three main components:

The **domain** is a collection of the possible statement instances that occur within the iteration space of a set of loop bounds. We can represent each statement instance with a multidimensional co-ordinate corresponding to the iterator values of the loops that surround it. As the constraints on the loops are affine, the bounded domain forms a convex polyhedron.

A **set of accesses** are affine mappings of the iteration space to memory. Two statement instances have a dependence ordering between them if they have accesses to the same memory location and at least one of them is a write.

A **schedule** assigns a timestamp to each statement instance, dictating the order in which they are executed. Different schedules for the same program represent possible program transformations. The transformed schedules are determined to be semantically preserving if dependence ordering is preserved.

To illustrate this, consider the implementation of the 1×1 convolution at the start of a residual block listed in Algorithm 1. We can describe the domain as follows:

$$\begin{array}{ll} \mathtt{S1}(c_o, h, w) & \mid 0 \leq c_o < C_o \land 0 \leq h < H \land \\ 0 \leq w < W \\ \\ \mathtt{S2}(c_o, h, w, c_i) \mid 0 \leq c_o < C_o \land 0 \leq h < H \land \\ 0 \leq w < W \land 0 \leq c_i < C_i \end{array}$$

Algorithm 1 Naive implementation of 1×1 tensor convolution.

We can also describe the schedule as follows:

$$T_{S1}(c_o, h, w) = (c_o, h, w)$$

 $T_{S2}(c_o, h, w, c_i) = (c_o, h, w, c_i)$

We now briefly discuss the transformations we consider.

Loop Interchange. Interchanging two nested loops involves applying a permutation to the schedule of the enclosed statements. For example, in the algorithm shown in 1 and a statement S1 we can express this as simply as:

$$T_{S1}(c_o, h, w) = (c_o, w, h)$$

Strip-Mining. This is performed by mapping an iterator into two new iterators whose combined range is that of the original. A constant strip-mining factor is selected which forms the range of the new inner loop. The outer loop range is that of the original divided by the strip-mine factor. For example, to strip mine the inner c_i loop of Algorithm 1 we have

$$T_{S2}(c_o, h, w, c_i) = (c_o, w, h, c_i/32, c_i \pmod{32})$$

Tiling. Tiling is a combined transformation consisting of stripmining followed by interchange. For example, to tile c_i loop of 1 we have

$$T_{S2}(c_o, h, w, c_i) = (c_i/32, c_o, w, h, c_i \pmod{32})$$

There are many more transformations that can be expressed in the polyhedral model. For further detail, we point the reader to the excellent polyhedral literature [27, 71].

4.1 Legality

Legality in the polyhedral framework is determined by preservation of data dependences. If there is a data dependence between two dynamic statement instances under the original program schedule, the relative ordering between these statements must be preserved under the new transformed schedule.

In the case of Algorithm 1, we can see that S1 is the definition, or source, of a dependence that must occur before its usage (in S2). For an instance i of statement S1 and j of statement S2, assuming constant data dependences, we can represent dependences in matrix form D. Elements of D are non-negative integers such that an element $d_{S1,S2}$ indicates that i must execute before j (formally, i < j, where < denotes lexicographical ordering).

Given a linear transformed schedule of the iteration space I in matrix form T and a dependence matrix D, then a transformation is legal iff:

$$\forall i, j, S1, S2, D \quad i \rightarrow j \in d_{S1,S2} \rightarrow T(i) \leq T(j)$$

Algorithm 2 Grouping transformation of tensor convolution.

which is to say that the lexicographical ordering is preserved.

5 UNIFIED SPACE

Our approach to joining NAS and compiler optimizations is to describe convolutional alternatives as polyhedral program transformations. Due to the affine nature of convolutions, this is relatively straightforward, except for checking transformation legality.

We adapt a newly developed legality check based on Fisher information [69], called Fisher Potential, that allows us to check the legality of transformations without having to retrain the network each time. This leads to a dramatic reduction of transformation search time as shown in Section 7.

5.1 Extending the Polyhedral Model

Bottlenecking. The bottleneck transformation is a reduction in the outer iterator in the domain node by parameterized constant factor B. Let the vector \underline{J} denote the iterators spanning the domain of the tensor loop nest, $\underline{J} = [c_o, c_i, h, w, k_h, k_w]$ where c_o is the outermost iterator and $\underline{J'} = [c_i, h, w, k_h, k_w]$ be the enclosed inner iterators, then bottlenecking can be expressed as

$$T_{S}(c_{o}, J') = (c'_{o}, J') \mid c'_{o} < C_{o}/B$$

The value *B* is constrained such that $C_0 \pmod{B} \equiv 0$. This gives a factor *B* reduction in computation. Figure 1 row 4 shows an example of bottlenecking.

Grouping. Grouping can be thought of as tiling the two outer iterators by a common factor and then discarding one of the iterators. The original iterator domains must both be divisible by the grouping factor G. Let $J' = [c_i, J'']$, such that $J = [c_o, c_i, J'']$ then we can define grouping as

$$T_{S}(c_{o}, c_{i}, J'') = (g, c_{o}/G, c_{i}/G, J')$$

to give the algorithm in Figure 2. Note as C_o, C_i and G are compiletime known constants each of the loop bounds is affine. Examining the code we see that each slice g of the output array refers only to the corresponding slice of the weight and input array.

Depthwise. Depthwise convolutions can be considered as a special case of group convolutions where the group size equals the number of output channels, C_o , $G = C_o$. For this transformation to be possible, the number of input and output channels must be equal, $C_o = C_i$. This means the two inner loops will have strip

Algorithm 3 Depthwise transformation of tensor convolution.

```
for (g=0; g<Co; g++)
for (oh=0; oh<OH; oh++)
for (ow=0; w<OW; ow++)
for (kh=0; kh<KH; kh++)
for (kw=0; kw<KW; kw++)
O[g][oh][ow] +=
W[g][kh][kw] *
I[g][oh+kh][ow+kw];</pre>
```

counts of 1 as $C_o/G=C_i/G=1$ and $T_{\mathbb{S}}(c_o,c_i,J'')=(g,1,1,J')$ which can be trivially simplified to

$$T_{\mathsf{S}}(c_0, c_i, J^{\prime\prime}) = (q, J^{\prime})$$

5.2 Fisher Potential as a Legality Check

The transformations described above fail to preserve traditional program semantics. We instead state that a schedule transformation for a neural network with respect to a task is legal if the final classification accuracy on the held out data is the same, or similar to within a small δ . Training all possible transformed networks to determine accuracy, however, would be prohibitively expensive.

To address this, we employ Fisher Potential [69] as a pre-training legality check. It is a cheap-to-compute measure that is effective at rejecting any architectures whose final test accuracy is significantly below the original starting point, without needing to perform training. Informally, Fisher Potential is, locally, the total information that each loop nest (layer) contains about class labels under a simplifying assumption of conditional independence. We note that the specific measure could easily be swapped out for another, such as [46], or any of the measures developed in [2], and that this is an area of active development.

Fisher Potential refers to the use of aggregated Fisher Information at initialization to estimate the effectiveness of network architectures before training. Where [40] used the diagonal of the Hessian in order to compress networks, [50, 67] showed that this computation could be approximated via the Fisher Information Matrix. This approximation was empirically shown to be effective at estimating the importance of individual neurons in a neural network [49], and to correlate to final accuracy [26]. In SNIP [41], the authors used a measure heavily related to Fisher Information *at initialization* in order to prune connections in the network prior to the training process. This was adapted in [69] to perform architecture search, by summing the Fisher Information over neurons to get layer-wise importance estimates. It is this form of the measure that we use in this paper.

Formal Definition. For a single channel, c, of a convolutional filter in a network, consider that for some input minibatch of N examples, its outputs are an $N \times W \times H$ tensor where W and H are the channel's spatial width and height. We refer to this tensor as the activation A of this channel and denote the entry corresponding to example n in the mini-batch at location (i, j) as A_{nij} . As the network has a loss function \mathcal{L} , then we can get the gradient of the loss with respect to this activation channel $\frac{\partial \mathcal{L}}{\partial A}$. Let us denote this

gradient as g and index it as g_{nij} . The channel c error, Δ_c can then be computed by

$$\Delta_c = \frac{1}{2N} \sum_{n}^{N} \left(-\sum_{i}^{W} \sum_{j}^{H} A_{nij} g_{nij} \right)^2. \tag{4}$$

This gives us a filter-wise score for a particular channel. In order to gauge the sensitivity of the full convolution, we sum Δ_c over each channel (as in [69]):

$$\Delta l = \sum_{c_o}^{C_o} \Delta_{c_o} \tag{5}$$

This score is summed for each of the convolutional blocks in the network. For an original network and a proposed alternative architecture, we reject the proposal if its score is below that of the original.

To summarise, the Fisher Potential of a proposed network architecture is the sum of the Fisher Information for each layer when given a single random minibatch of training data, as performed in [69].

Example. Let us consider, candidate networks designed from NAS-Bench-201 [17] as shown in Figure 3. Each point represents a different neural architecture of which there are 15625. The *y*-axis shows final CIFAR-10 top-1 error (lower is better), and the *x*-axis shows the Fisher Potential assigned to the model *at initialization* (higher is better). We can see that without requiring any training, Fisher Potential is able to filter out poorly-performing architectures, visible in the cluster of low scoring networks on the left with poor final errors. Many good networks are also discarded — this is unfortunate but acceptable for our scenario, since the space is large and densely populated with networks. We note again that this measure in particular could be swapped out for an improved one in future.

5.3 Expressive Power

Neural Architecture Search techniques rely on hand-designed exploration spaces. themselves are engineered by hand. For example, a recent paper found that bottlenecking could be applied in the spatial domain [53]. This can now be added to a list of candidate operations. However, spatial bottlenecking is automatically captured within our framework as the combination of existing transformations. As an example, consider the convolution in row 2 of Figure 1. We use the shorthand notation $[i] \xrightarrow{B(b)} [i(b)]$ to denote bottlenecking domain i by factor b and $[i,j] \xrightarrow{\mathrm{int}} [j,i]$ to denote interchange. Then the spatial bottleneck operation can be constructed as the following sequence of transformations:

$$T_{S}: [C_{o}, C_{i}, H, W, K_{h}, K_{w}] \xrightarrow{\text{int.}}$$

$$[H, W, C_{o}, C_{i}, K_{h}, K_{w}] \xrightarrow{\text{B(b).}}$$

$$[H(b), W, C_{o}, C_{i}, K_{h}, K_{w}] \xrightarrow{\text{int.}}$$

$$[W, H(b), C_{o}, C_{i}, K_{h}, K_{w}] \xrightarrow{\text{B(b).}}$$

$$[W(b), H(b), C_{o}, C_{i}, K_{h}, K_{w}] \xrightarrow{\text{int.}}$$

$$[C_{o}, C_{i}, H(b), W(b), K_{h}, K_{w}] \xrightarrow{\text{int.}}$$

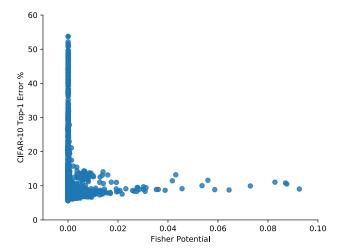


Figure 3: Fisher Potential as a rejection filter for invalid architectures. Each point is a different neural architecture from NAS-Bench-201 [17]. The y-axis shows final CIFAR-10 top-1 error (lower is better), and the x-axis shows the Fisher Potential assigned to the model at initialization (higher is better).

The expressiveness of the polyhedral model, equipped with these new transformations, means that there are novel transformations that can be derived from this unified framework.

6 IMPLEMENTATION AND SETUP

We expressed our new transformations as polyhedral transformations and implemented the resulting operators in TVM [10]. We selected TVM as its API allows composition of transformation sequences, and is a well-accepted state-of-the-art optimizing compiler for convolutional neural networks.

Transformation Space. An overview of the transformations used in this paper is given in Table 1. The first four are standard program transformations available within TVM. To these, we add our two neural architecture transformations. Finally, for GPU platforms, we enable GPU mapping transformations.

Baseline TVM. TVM allows users to implement schedules for each new operator by hand. Due to the large number of operators involved, for each device we use TVM's default schedules; automatic schedule design [84] has yet to be incorporated into TVM. We then enable auto-tuning of parameter values within the schedule to find best performance.

Search. Our current search process is relatively naive. We enumerate random sequences of transformations through TVM, and generate 1000 configurations of the resulting operations for each network. We then check which candidates pass our implementation of the Fisher Potential legality test and select the best performing one.

Table 1: A description of the autotuning primitives available to TVM for optimizing operations, including GPU mapping, standard primitives, and our two new optimizations.

Optimization	Description
Program Transformations	
reorder	Interchange nested loops
tile	Cache and register blocking
unroll	Loop unrolling
prefetch	Memory coalescing between threads
split	Divide iteration into multiple axes
fuse	Combine two axes into one
Neural Architecture Transformations	
bottleneck	Reduce domain by factor B
group	Slice and offset two loops by factor <i>G</i>
Mapping to GPU	
blockIdx	Block-wise parallelism
threadIdx	Threads within blocks
vthread	Striding thread access

6.1 Experimental Setup

<code>Platforms.</code> We evaluate on an ARM A57 mobile CPU (mCPU), and an Nvidia 128-Core Maxwell mobile GPU (mGPU) available on the Jetson Nano board. We also evaluate an Intel Core i7 (CPU) and an Nvidia 1080Ti (GPU). This is representative of a wide range of deployment targets, from mobile to server class. We use TVM v1 . 7 compiled with CUDA 10.1 and LLVM 8.0. For training and accuracy evaluation we use <code>PyTorch v1</code> . 4.

Neural Models. We evaluate our methodology on three popular networks: ResNet-34 [29], ResNeXt-29 [78], and DenseNet-161 [34]. These networks were chosen to represent a range of convolutional architectures, from standard 3×3 convolutions in ResNet-34 to grouped convolutions in ResNeXt and a heavy reliance on 1×1 convolutions in DenseNet. CIFAR-10 models are trained for 200 epochs with Stochastic Gradient Descent (SGD) [6] and a learning rate of 0.1, decayed by a factor of 10 at epochs 60, 120, and 160. ImageNet models were trained using the default PyTorch script 1, which trains for 90 epochs with SGD, starting from a learning rate of 0.1 and decaying by a factor of 10 every 30 epochs.

Comparison. The models are implemented with each operation written as a TVM Tensor Expression [10], which is an einsum-style syntax for expressing tensor computations. This is lowered to TVM IR, where our transformations can be employed. This allows for a fair comparison of each approach.

For each network, we consider three approaches. First we compile the model using TVM using its default schedule (labeled TVM). We report the best performance achieved after auto-tuning. Next, we use BlockSwap [69] as NAS to compress the modifiable convolutions in the network, followed by compilation with TVM (labeled NAS). Finally, we apply our unified approach as described in the previous section (labeled Ours).

¹https://github.com/pytorch/examples/tree/master/imagenet

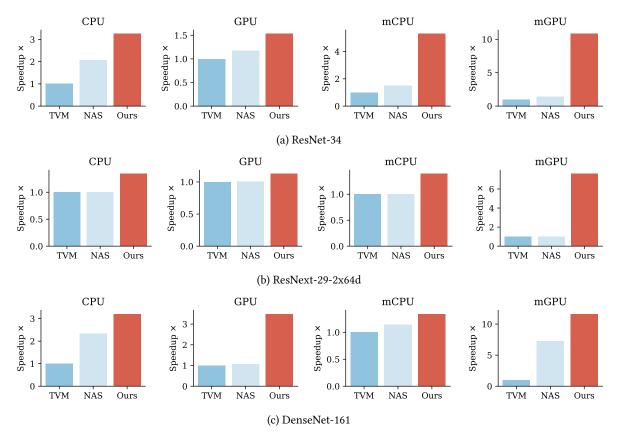


Figure 4: End-to-end performance for several networks on different hardware devices on CIFAR-10. TVM represents the original model compiled with TVM default schedules. The NAS columns represent the BlockSwap-compressed copies of the models which are then compiled with TVM default schedules. Ours represents our unified NAS-compilation strategy, with new operators stacked into the network blocks.

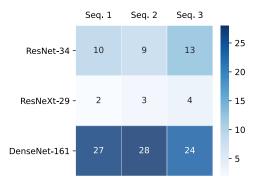


Figure 5: Frequency of operation application.

7 RESULTS

In this section we first present the performance of the three approaches on different networks and platforms for the CIFAR-10 dataset. This is followed by an analysis of accuracy, size, search time and the highest performing new convolutions found. We then

drill down into one network and examine the impact of transformations layer-by-layer. In order to compare against alternative NAS techniques, we evaluate the performance of FBnet across the three networks. Next, we evaluate the performance of our approach on the ImageNet dataset additionally examining the tradeoff in accuracy and performance across variants of the ResNet and DenseNet family of networks. Finally we show that our approach allows for fine-grained exploration of an accuracy/size tradeoff, producing a new Pareto optimal network design.

7.1 CIFAR-10 Network Performance

The results for each network are shown in Figure 4. The combination of TVM and NAS is a strong baseline, however, for each of the models our unified method is able to generate models with improved hardware performance. In general there is more performance to gain on the mGPU than other platforms, as relaxed memory pressure from smaller designs is of increasing importance.

ResNet-34. NAS is able to find a modest improvement of $1.12 \times$ speedup over TVM on the GPU platform, increasing to $2 \times$ on i7 CPU, showing the power of compressed convolutions. Our unified approach is able to find further improvement giving $2 \times$ and $3 \times$

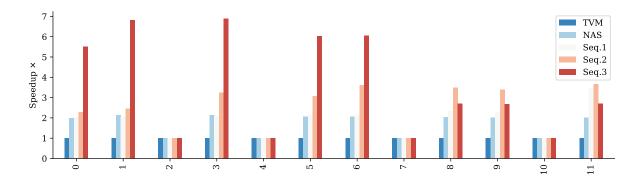


Figure 6: Exploring different sequences of transformations for an individual layer of ResNet-34 on the Intel Core i7 CPU. NAS is the result of applying grouping with factor 2 first, then compiling with TVM. The other three sequences are interleaved transformations produced by our method.

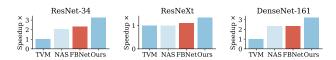


Figure 7: Intel Core i7 performance of FBNet on the three networks. FBNet is able to improve over NAS with large training cost. Our approach outperforms FBNet with no training required

speedup respectively. The improvement is more dramatic on the smaller platforms with $5\times$ and $10\times$ speedups on the mCPU and mGPU.

ResNext-29-2x64d. NAS is unable to find any improvement here due to the already highly compact structure of the network. There are simply no NAS options that improve performance over the TVM baseline across all platforms. Despite this our approach is able to find small improvements of $1.3\times$ and $1.1\times$ on the CPU and GPU platforms by combining NAS and program transformations. This increases to $1.4\times$ on the embedded CPU and $7\times$ on the mGPU.

DenseNet-161. The impact of NAS is much more varied here. While it can find 2.2× improvement on the CPU, it has a negligible improvement over TVM on the GPU. It also struggles on the mCPU but finds $6\times$ on the mGPU. Our approach is able to improve by over $3\times$ for both server platforms, but only finds $1.2\times$ on the mCPU while acheiving $10\times$ on the mGPU.

7.2 Analysis

Accuracy For all CIFAR-10 networks pictured in Figure 4, changes in accuracy were less than 1% in absolute difference. The ResNet-34 in Figure 6 had an original ImageNet Top-1 and Top-5 accuracy of 73.2% and 91.4% respectively. The final network compiled with our transformations had a Top-1 and Top-5 accuracy of 73.4% and 91.4%; it was slightly more accurate than the original but considerably faster. To give a broader overview, we present the accuracy of several ImageNet models in Figure 8, which shows that for each model accuracy degradation is small or non-existent.

Size: One benefit of these neural transformations is their effect on model size, both in terms of weights and runtime memory usage. We found that CIFAR-10 networks could be compressed in size by $2-3\times$. Likewise, the ImageNet ResNet-34 in Figure 6 was compressed from 22M parameters to 9M without a loss in accuracy.

Search time: Our search process involves suggesting configurations and rejecting or accepting them based on Fisher Potential. Since Fisher Potential is extremely cheap to compute, the search time overhead introduced by our method is small, less than 5 minutes on a CPU. During this time we were able to explore 1000 different configurations, discarding approximately 90% of the candidate transformation sequences through the Fisher Potential legality check.

7.3 Transformation Sequence Case Studies

There were 3 particular transformation sequences that dominated the list of best performing transformations. In neural architecture terms, the resulting operations of each of these sequences of transformations is a new convolution-like operator, previously unavailable to NAS.

Sequence 1 applies grouping to the kernels over the spatial domain of the input. These are then concatenated to form one output. The exact sequence is: [split \rightarrow interchange \rightarrow group \rightarrow interchange \rightarrow fuse].

Sequence 2 is an operator in which the output channels have been unrolled by factor 16 and then the remaining iteration domain has been grouped by factor G=2. The exact sequence is: [unroll \rightarrow group \rightarrow interchange] Sequence 3 is an operator that was devised by splitting up the iteration domain of the output channels, and applying different levels of channel grouping to each new domain (e.g. G=2 on first half, G=4 on the second half). The exact sequence is: [split \rightarrow group \rightarrow interchange \rightarrow group].

Figure 5 shows how often each of these sequences appear in our best performing networks. ResNext-29 has the fewest instances as it contains the fewest layers, while DenseNet-161 has the most layers and hence most instances. Each of them is applicable across all networks and may be more widely applicable as standard tensor convolutions for other networks.

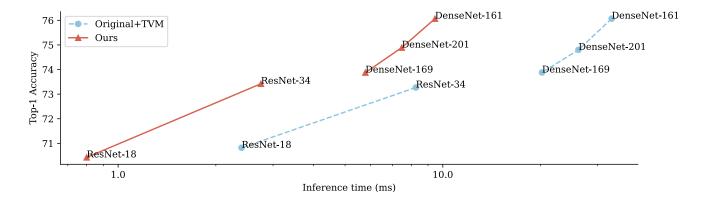


Figure 8: Accuracy vs inference time (log scale) of our approach (Ours) compared to TVM (Original + TVM) when applied to different variants of ResNet and DenseNet on the ImageNet dataset. Our approach gives a significant reduction in inference time.

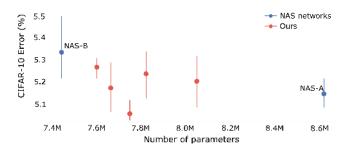


Figure 9: Two NAS models (the blue points labeled NAS-A and NAS-B) composed of grouped blocks (with g=2 and g=4 respectively) can be chained together by a series of parametrized transformations in our framework, yielding the points in red. Each point is the mean of three training runs, with error bars.

7.4 Exploring Layer-Wise Optimizations

In Figure 6, we examine the impact of our sequences layer-by-layer on one network on one platform: ResNet-34 on the i7. The exact configurations mirror the experiment in the original TVM paper [10]. While we achieve a 3x improvement overall as shown in Figure 4, the speedup achieved varies considerably across layers. In 4 of the 11 layers no performance improvement is found, as Fisher Potential marks these individual layers to be extremely sensitive to compression. Simple grouping with a factor G=2 is able to give around $2\times$ speedup across 7 of the 11 layers. Sequence 1 is able to give a small improvement in most cases — particularly layer 11 — where spatial reduction yields new parallelization opportunities. Sequence 2 provides a slight improvement in many cases, and through the later layers it is able to offer larger speedups through improved data reuse. Sequence 3 is the best option for most early layers but suffers compared to other sequences in the later layers.

7.5 Alternative NAS comparison

To provide an alternative evaluation against an existing NAS technique, we re-implement FBNet [77] using the convolutional blocks available in our NAS space, and our three baseline networks as the skeletons into which the selected cells are inserted. Figure 7, shows the performance of resulting networks on CIFAR-10 relative to TVM, NAS and our approach on the Intel Core i7 for each network. In each case, FBNet does provide a modest improvement over NAS. However, it is worth noting that this involves an expensive training step at each stage of evaluation (requiring ~3 GPU days per network). Our approach is able to consistently improve over FBNet, with no training required.

7.6 ImageNet Network Performance

We also applied our technique to the ImageNet classification dataset, to show that the method transfers beyond CIFAR-10. In Figure 8, we first show the resultant networks for running our method on ResNet-18, ResNet-34, DenseNet-161, DenseNet-201, and DenseNet-169, with their inference times recorded on the Intel i7 CPU. For each network, accuracy is within 2%, and there are significant gains in inference time.

7.7 Interpolating Between Models

The additional expressive power of our method allows us to explore the search space of networks in a more fine-grained manner than traditional NAS techniques. This is because NAS techniques simply choose operations from a list, whereas we generate new ones. To illustrate this consider Figure 9 where we plot the accuracy and inference time of a ResNet-34 with two BlockSwap-based models (the blue points labeled NAS-A and NAS-B). We can explore alternatives by a series of parametrized transformations in our framework, yielding the points in red. Each of these points is a new possible blocktype that would not be accessible to a traditional NAS technique unless explicitly written by the human designer. During this interpolation we are also able to find a Pareto optimal point.

8 RELATED WORK

Compiler optimization. There has been much interest in autotuning DNN code generators [10, 23, 37, 48, 64, 72]. Polyhedral compilers are particularly well-suited [72, 83] as they have in-built abstractions for exploiting parallelism and memory layout in a principled form. Polyhedral compilation itself benefits from a wealth of literature [5, 13, 24, 38, 71, 75] that describes a unified compilation scheme and associated set of legality checks delivering state-of-the-art performance across a wide range of standard compiler benchmarks [27, 73, 74], as well as image and neural network specific ones [51, 72, 83].

Other popular approaches include loop synthesis [10] and rewrite rules [64] which optimize through parametrized schedules. In particular, TVM [10] is an adaptation of Halide [57] with specific extensions for deep neural network abstractions. Triton [68] is a tile-based IR for parallelizing tensor-based computation. Most of these frameworks focus on operator-level optimization, though higher level graph transformation has also shown promising results [37].

Though the toolchains are exceptionally feature-rich, there is evidence that some implementations have been highly-engineered for specific workloads, at the cost of general support for newer optimizations [4]. All current approaches, however, are limited by their inability to exploit NAS transformations.

Reducing the amount of computation at the cost of accuracy has been examined in a compiler context using loop perforation [20, 59, 62] in approximate computing. Within approximate computing, there is extensive work on probabilistic program transformations [47, 58, 85] and language support for approximation [25, 52]. Where some approximate computing methods accept small absolute differences in the outputs of programs, our transformations may render the numerical outputs of our programs completely different while maintaining the legality of the program. The benefit of Fisher Potential, while it does not give any guarantees on behavior, is that its domain specificity allows us to capture these transformations.

Model Optimization. From a machine-learning perspective, deep neural networks should be highly compressible as there is significant evidence that they are vastly overparametrized [16, 21]. There are several popular strategies that can be employed. One can prune a network by removing unimportant connections between weights [22, 28, 50]. However, the sparsity introduced can lead to poor memory behavior [14]. An alternative is to retain dense weight tensors by pruning channels [42] although the resulting irregular structures cause a significant slowdown [56].

Network distillation [3, 31, 70], takes a trained large network and uses its outputs to assist in the training of a smaller network. This, however, leads to a large design space as there are many ways to make a network smaller [11, 32, 33, 36, 78].

Neural architecture search instead automates the process of finding efficient networks. In [86] the authors use an RNN to generate network descriptions and filter the options using reinforcement learning. Several networks are generated and trained in parallel, which requires a large quantity of possible networks to be stored at once. To reduce this overhead, an alternative method is to design one *supernet* that contains all of the possible subnetworks [54]. This allows all models to have access to a shared set of weights,

reducing the computational requirement. Subsequent works have made extensive use of this technique [7, 44, 45, 77].

However, there is some evidence to suggest that weight sharing schemes aggressively hamper the ability of the NAS agent to identify good architectures, and under such constrained spaces, random architecture search provides a competitive baseline [43, 61]. Although NAS is a rapidly developing area, no-one to the best of our knowledge has considered formalizing NAS as program transformation.

9 CONCLUSIONS AND FUTURE WORK

This paper presents a new unified program transformation approach to optimizing convolution neural networks by expressing neural model operations as formal program transformations. We develop a new legality check based on Fisher potential, and show that different types of convolution can be described as transformations of more fundamental building blocks. We implemented this approach in TVM and show that our combined approach significantly outperforms both TVM and state-of-the-art NAS. We eliminate the need to train while searching, dramatically reducing search time. Future work could consider further transformations in the neural architecture space in order for a more extensive analysis, and to expand the possibilities beyond that of just convolutional networks.

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