Manas: Mining Software Repositories to Assist AutoML

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ABSTRACT

Today deep learning is widely used for building software. A software engineering problem with deep learning is that finding an appropriate convolutional neural network (CNN) model for the task can be a challenge for developers. Recent work on AutoML, more precisely neural architecture search (NAS), embodied by tools like *Auto-Keras* aims to solve this problem by essentially viewing it as a search problem where the starting point is a default CNN model, and mutation of this CNN model allows exploration of the space of CNN models to find a CNN model that will work best for the problem. These works have had significant success in producing high-accuracy CNN models. There are two problems, however. First, NAS can be very costly, often taking several hours to complete. Second, CNN models produced by NAS can be very complex that makes it harder to understand them and costlier to train them. We propose a novel approach for NAS, where instead of starting from a default CNN model, the initial model is selected from a repository of models extracted from *GitHub*. The intuition being that developers solving a similar problem may have developed a better starting point compared to the default model. We also analyze common layer patterns of CNN models in the wild to understand changes that the developers make to improve their models. Our approach uses commonly occurring changes as mutation operators in NAS. We have extended *Auto-Keras* to implement our approach. Our evaluation using 8 top voted problems from *Kaggle* for tasks including image classification and image regression shows that given the same search time, without loss of accuracy, *Manas* produces models with **42.9%** to **99.6%** fewer number of parameters than *Auto-Keras*' models. Benchmarked on GPU, *Manas*' models train **30.3%** to **641.6%** faster than *Auto-Keras*' models.

CCS CONCEPTS

• **Software and its engineering** → **Search-based software engineering**; • **Computing methodologies** → **Machine learning**.

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*This work was done when Md Johirul Islam was at Iowa State University.

KEYWORDS

Deep Learning, AutoML, Mining Software Repositories, MSR

1 INTRODUCTION

An increasingly larger number of software systems today are including deep learning. Deep learning uses a convolutional neural network (CNN) model, essentially a graph with weighted edges and nodes that are weighted functions, to convert inputs to the output. As more software systems incorporate deep learning, more software developers have to design and train CNN models. Designing a CNN model is very difficult, and developers often struggle leading to bugs. Model bugs are frequent bugs in CNN programs [29, 30].

Recent work on neural architecture search (NAS) aims to solve this problem [61]. NAS techniques start from a collection of default CNN models and search for a suitable model for the problem. The search space is defined by the collection of default models and a collection of mutation operators that are used to modify CNN models to create new candidates. NAS techniques have been implemented in industrial-strength tools such as *Auto-Keras* [31]. NAS techniques face two problems. First, NAS can be very costly, e.g., *Auto-Keras* takes 8-12 hours on high performance machines to search for models with reasonable accuracy (90+%). The accuracy drops rapidly if the search time is reduced. Second, CNN models produced by NAS can be very complex that makes it harder to understand them for maintenance, and costlier to train and retrain them.

We introduce *Manas* (Mining Assisted Neural Architecture Search) to alleviate the limitations of NAS. The fundamental intuition behind *Manas* is that mining and using the hand-developed models that are available in open-source repositories as default models or starting point of search can help NAS leverage human developer efforts. *Manas* applies this intuition in two ways. First, hand-developed models are mined to search for a better starting point for NAS. Second, the change patterns of the hand-developed models are mined to identify more suitable mutation operators for NAS.

We have realized *Manas* by extending *Auto-Keras*, the state-of-the-art NAS framework. *Auto-Keras* is open source and outperforms state-of-the-art methods like SEAS [17], NASBOT [32] making it a suitable baseline [31]. Some key technical contributions in *Manas*

Hi I was trying to train a cancer dataset, using the Resnet neural network by using the fine tuning approach

Here is how I used to fine-tune it.

```
image_input = Input(shape=(224, 224, 3))

model = ResNet50(input_tensor=image_input, include_top=True,weights='imagenet')
model.summary()
last_layer = model.get_layer('avg_pool').output
x= Flatten(name='flatten')(last_layer)
out = Dense(num_classes, activation='softmax', name='output_layer')(x)
custom_resnet_model = Model(inputs=image_input,outputs= out)
custom_resnet_model.summary()
```

(a) Problem

As far i can tell from your plots, you are overfitting. To avoid overfitting you should use Dropout. bellow i have added 2 new dense layers followed by 2 Dropout layers (you can reduce to 1 pair or increase to more). You can fine tune their parameters.

I would also like to propose a simpler representation of your network but your way would be just fine.

```
base_model = ResNet50(input_shape=(224, 224, 3), include_top=False,weights='imagenet'
x=base_model.output

x = Dense(512, activation='relu')(x) #add new layer
x = Dropout(0.5)(x) #add new layer
x = Dense(512, activation='relu')(x) #add new layer
x = Dropout(0.5)(x) #add new layer

out = Dense(62, activation='softmax', name='output_layer')(x)
```

(b) Solution

Figure 1: An example from *StackOverflow* showing the necessity to change model architecture

include *model matching*, a technique for matching the problem that the developer intends to solve with the hand-developed models mined from repositories, *model adaptation*, a technique for adapting the mined model to the problem context, *model transformation*, a technique for adapting the mined model for further improving metric values, and *training adaptation* that leverages mined parameter values from the repositories to change the optimizer.

To evaluate *Manas*, we use the top-8 problems from diverse domains obtained from Kaggle for machine learning tasks including image classification and image regression. Our evaluation shows that given the same amount of searching time, *Manas* generates simpler neural architectures than *Auto-Keras* without losing accuracy. In terms of the models' size, *Manas*' models have **42.9%** to **99.6%** fewer numbers of parameters than *Auto-Keras*' models. We observed up to **641.6%** faster training speed when training models produced by *Manas* as compared to those produced by *Auto-Keras*.

Our main contributions are the following:

- We have proposed a novel approach for NAS that leverages software repository mining.
- We have proposed methods to identify the suitable models by analyzing data characteristics and adapting models.
- We have utilized the common patterns to transform mined models to improve the performance of these models in terms of error rate, MSE, model complexity, and training time.
- We have implemented these ideas in a SOTA NAS framework, *Auto-Keras* [31]. **Our artifact is available here [21].**

The paper is organized as follows: §2 presents a motivating example, §3 presents preliminaries and problem statement, §4 describes the *Manas* approach for NAS, §6 describes the limitations and threats to validity of *Manas*, §7 describes related work, and §8 concludes.

2 MOTIVATION

Deep Learning has received much attention in both academia and industry. Therefore, many deep learning libraries and tools are created for supporting a large number of deep learning developers. Although these libraries and tools make deep learning more accessible, there are still many challenges. One of the challenging problems is constructing an appropriate CNN model architecture [53, 54], which also has been shown as a frequent bug in CNN programs by Islam *et al.* [29, 30]. For instance, Figure 1a shows a query [1] posted on *StackOverflow* where a developer is unable to find an appropriate

CNN model for their purpose. In particular, the question discusses the difficulty that the ResNet architecture does not give the result as the developer expected. In response, an expert suggests changing the CNN model. Figure 1b shows the solution of the expert for the question of the developer. The expert suggested that the developer should add dropout layers to minimize overfitting.

Neural architecture search (NAS) aims to solve this problem [61]. NAS takes the training data as an input to automatically define the neural network for that data. Moreover, NAS is able to tune the hyperparameter of the searched neural network. There are both commercial and open-source realizations of NAS. For example, a developer can pay about $20 per hour to use Google's AutoML. *Auto-Keras* is an AutoML system using NAS [31] created as an open-source alternative. *Auto-Keras* returns outstanding results compared with state-of-the-art handmade models on CIFAR10 [33], MNIST [34], and FASHION [57] datasets. *Auto-Keras* is shown to outperform state-of-the-art methods like SEAS [17], NASBOT [32].

NAS has two limitations. First, it can be very costly. For example, *Auto-Keras* consumes 2,300% [47] more GPU computation time compared to using handmade model. Second, NAS often produces complex models that are hard to understand and time-consuming to train. To illustrate, we used *Auto-Keras* on another dataset, which is *Blood Cell* [39] collected from *Kaggle*. The model created by *Auto-Keras* for the *Blood Cell* problem has more than 2.3 million learnable parameters and more than 7 weight layers. The searched CNN models are constructed based on the architecture of the large default CNN models; thus, the models produced by *Auto-Keras* are often really large. Smaller CNN models train faster and save more energy [27, 44]. Han *et al.* have shown that reducing the number of parameters of deep learning models can reduce the training time by 3× to 4×, and energy comsumption by 3× to 7× [24]. The rest of this work describes our approach *Manas* that addresses these limitations. As an example, for *Blood Cell* dataset, *Manas* produces a model that decreases the error rate by 47.1%, the model depth by 14.3%, the model width by 87.0%, and increases the training speed by 56.9% compared with *Auto-Keras*'s model. The model produced by *Manas* has 6 layers and 0.3 million parameters (neurons), whereas the model produced by *Auto-Keras* has 7 layers, 2.3 million parameters.
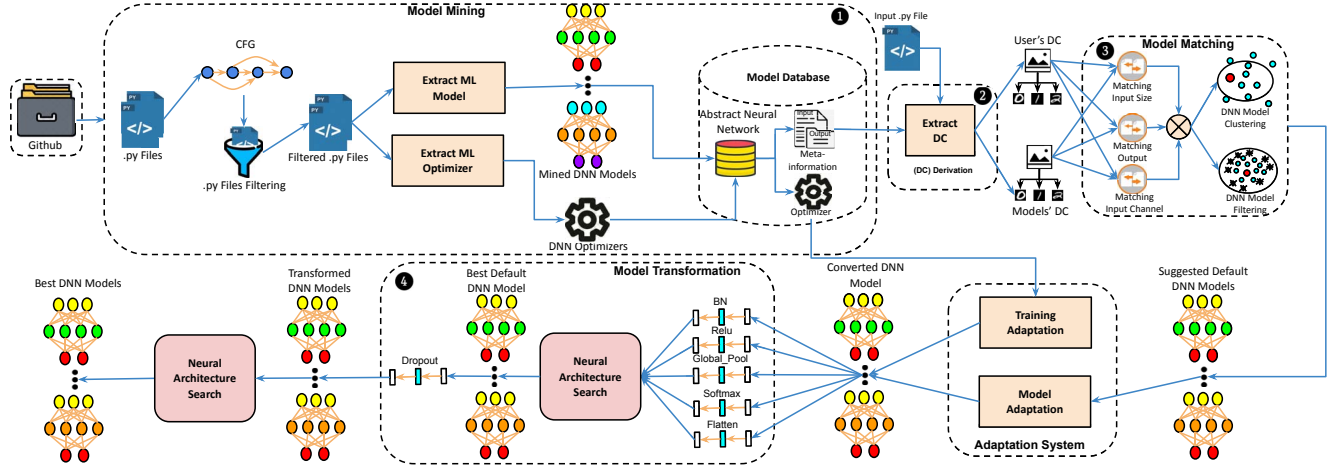
Figure 2: An Overview of *Manas*. Two inputs are mined models from repository (left-top), and user's initial file (middle-top).

3 PRELIMINARIES AND PROBLEM STATEMENT

3.1 Preliminaries (NAS)

We define NAS like it was defined in *Auto-Keras* [31]. Given a search space $\mathcal{S}$ and the input data $\mathcal{D}$ split into $\mathcal{D}_{train}$ and $\mathcal{D}_{val}$, the goal is to find an optimal neural architecture $\mathcal{N} \in \mathcal{S}$, which achieves the lowest value of the cost function in dataset $\mathcal{D}$. Search space $\mathcal{S}$ covers all the neural architectures created from default neural networks.

$$\mathcal{N} = \arg\min_{\mathcal{N} \in \mathcal{S}} Cost(\mathcal{N}'(\theta'), \mathcal{D}_{val}) \quad (1)$$

$$\theta' = \arg\min_{\theta} \mathcal{L}(\mathcal{N}'(\theta), \mathcal{D}_{train}) \quad (2)$$

Where *Cost* and $\mathcal{L}$ are the cost function and loss function, θ is the learned parameter of initial architectures $\mathcal{N}'$.

3.2 Problem statement

This work aims to utilize the neural networks from open source repositories to optimize the neural architecture search. We extract the data characteristics from both the input dataset and mined neural networks to determine better starting points (initial models) for NAS. Instead of using concrete default neural networks, the goal is to find optimal initial architecture $\mathcal{N}^*$ for NAS for each different input dataset. Similar to Equation 2, we obtain learned parameter θ^* of new initial architectures $\mathcal{N}^*$:

$$\theta^* = \arg\min_{\theta} \mathcal{L}(\mathcal{N}^*(\theta), \mathcal{D}_{train}) \quad (3)$$

The optimal initial architectures $\mathcal{N}^*$ support NAS to find out the optimal neural network faster. In other words, with the same amount of searching time and input dataset, new initial architectures help NAS to find out a neural network with lower error compared to the original NAS:

$$\arg\min Cost(\mathcal{N}^*(\theta^*), \mathcal{D}_{val}) < \arg\min Cost(\mathcal{N}'(\theta'), \mathcal{D}_{val}) \quad (4)$$

4 MANAS

Figure 2 shows the overview of *Manas*. *Manas* has five major components that we describe below.

❶ To initialize *Manas* for NAS, the model database must be populated by mining models from open source repositories. Currently, *Manas* collects high quality models from *GitHub* by extracting Python files from *Keras* projects. These projects are selected using certain filtering criteria to ensure code quality. Then, API usage is used to filter Python files to those that contain models. Finally, both the models and the values for optimizer are extracted to store in the model database. This database is constructed once and should be updated periodically as new models are added frequently.

❷ The data characteristics extracted from both the input data and mined models are used to select the suitable initial models for NAS from the database.

❸ Model matching matches the data characteristics extracted from an input dataset and the mined models to obtain a good starting point. It selects the models, which have the closest data characteristics with the input dataset by using the model clustering approach. In case there are too many initial models, the model filtering approach will be applied to reduce the number of models.

❹ The selected models are transformed by the model transformation approach based on related state-of-the-art papers and common layer API patterns of mined models. The transformation can enhance the performance of the models in terms of errors and training speed.

4.1 Model Mining

In order to extract CNN models and their optimizers from source code repositories, we build a source code analyzer based on the control flow graph (CFG). Figure 3 shows the overview of model mining process.
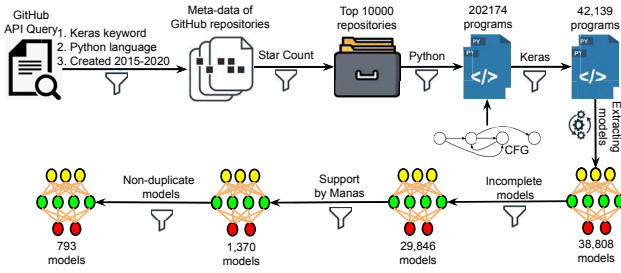
Figure 3: Model mining process.

4.1.1 Meta-data collection. We collect the meta-data of *GitHub* repositories by using a *GitHub* API query [1]. Meta-data contains basic information of a repository such as authors, repository's name, etc. The query allows us to obtain the meta-data of the *GitHub* repositories with three filtering criteria including Python programming language, containing *Keras* keyword, and created date between 2015-01-01 and 2020-12-31. From the meta-data, we obtain *GitHub* URLs of the top 10,000 repositories with the most star count to ensure the quality of the models [8]. The URLs help us to access the repositories to collect the CNN models.

4.1.2 Keras programs collection. We obtain 202,174 Python programs from collected repositories; however, only 42,139 programs use *Keras* API. In particular, we use CFG to analyze their import statements of Python programs to only collect which one imports *Keras* API.

4.1.3 Models extraction. In this work, since *Manas* only works with *Keras* CNN, we will explain how we extract CNN from *Keras* programs. We use CFG to extract a model from a deep learning program. We manually create a list of function calls used to build neural networks of *Keras* based on *Keras*' documentation [2]. Then, CFG examines every API call to collect the functions contained in the list and their connections. The collected functions represent the layers in the models. The connections between functions represent the layer connections in the models. The reason for using CFG is to collect complete models from programs even if they contain branches. For example, the CFG contains a convolutional block followed by a dense block in the "if" branch and a skip connection in the "else" branch. If a convolutional block combines with a dense block or a convolutional block combines with a skip connection to be a complete model, we will extract those parts in the branches to collect the complete model. CFG is used to handle the situation that there is a loop containing a part of a model in the CNN programs. If the number of iterations is available to extract that part completely, we will collect it to complete the model. If there is a method call in the program containing a part of the model, we will collect it to complete the model. For the other cases, when the parts in the branches cannot complete the model or cannot be extracted, we will ignore them. For instance, a part of the model in a loop whose numbers of iterations are unavailable cannot be extracted. After this step, we collected 38,808 models from 42,139 *Keras* programs. The number of models is less than the number of *Keras* programs

because many programs do not contain models. We assume that a program contains a model if it has at least an API used to build a model.

Rather than mining only neural architectures, we also mine their optimizers that deep learning developers carefully select after spending manual efforts on retraining their models. Then, whenever a model is selected as an initial model, *Manas* trains the model with its optimizer instead of the default one. Optimizers are algorithms deciding how the parameters of the models change. Every optimizer has strengths and weaknesses; thus, it is necessary to choose a suitable optimizer. While models are the decisive factor to the performance of *Manas*, we note that it is wasteful if we cannot fully use these models. In other words, good models with wrong optimizers cannot produce a good performance. To obtain the optimizer, we use the same process of extracting CNN models by creating a list of functions related to the optimizer. After that, CFG analysis is used to obtain the API call, which contains the optimizer.

4.1.4 Incomplete models detection. A complete model includes an input layer, hidden layers, and an output layer [60]. The input layer is the first convolutional layer of the neural network, which is distinguished from the other convolutional layers based on its parameters. In particular, only the API call representing the input layer contains *input_shape* parameter. The output layer is the last linear layer of the CNN. Hidden layers are the layers between the input layer and the output layer, including convolutional layers, activation layers, and fully connected layers (linear layers). Therefore, if an extracted model does not have convolutional layers, activation layers, or linear layers, we will consider that the model is incomplete. By removing incomplete models, there are 29,846 models left.

4.1.5 Supported by Manas. As *Manas* currently supports a few kinds of layers that are the convolutional layer, the linear layer, the batch normalization layer, the concatenate layer, the add layer, the max pooling layer, the dropout layer, the Softmax layer, the ReLU layer, the flatten layer, and the global pooling layer, we filter out the models containing unsupported layers. After this step, we have 1,370 models left.

4.1.6 Model duplication detection. We obtain 793 models after removing the duplicate ones. We consider if two models have the same abstract neural network (ANN), there will be a duplicate model. We store the extracted CNN models in a database as an abstract neural network, which is an abstract representation of the neural network. This representation has the structure as a network where the nodes are API calls, and the connections are the order between API calls. We use this representation to adapt models and their optimizers into *Manas*. From each node, we obtain the name of layer and its parameters, which can be converted into an API call. Figure 4 presents an example of ANN built from an mined model. Notice that if an activation parameter is implemented as an argument inside a layer, we consider that the activation is a separate layer.

	Conv2D(64, kernel_size = (3, 3),activation='relu', input_shape=(3, 120, 180))	1	{'func': 'Conv2D', 'input_shape': [3, 120, 180], 'arg2 ': 64, 'kernel_size': [3, 3]}
2		2	{'func': 'relu'}
3	MaxPooling2D(pool_size=(2,2))	3	{'func': 'MaxPool2d', 'pool_size': [2, 2]}
4	Conv2D(32, activation='relu')	4	{'func': 'Conv2D', 'arg1': 64, 'arg2': 32}
5		5	{'func': 'relu'}
6	MaxPooling2D(pool_size=(2,2))	6	{'func': 'MaxPool2d', 'pool_size': [2, 2]}
7	Dropout(0.25)	7	{'func': 'Dropout', 'arg1': 0.25}
8	Flatten()	8	{'func': 'Flatten'}
9	Dense(20, activation='softmax')	9	{'func': 'linear', 'arg1': 128, 'arg2': 20}
		10	{'func': 'softmax'}
		11	
12	SGD(lr=0.01, decay=1e−6)	12	{'func': 'SGD', 'lr': 0.01, 'decay': 1e−06}

(a) Extracted model **(b) Abstract Neural Network**

Figure 4: Building an abstraction of neural network from extracted model

4.2 Data Characteristics Derivation

We analyzes the input data and input/output layers of the mined models to extract their data characteristics. Taking an image classification as an example, we obtain the input data characteristics including the input size and the output channel from the image data. The data characteristics extracted from a mined model also are the input size and the output channel are obtained by analyzing the first convolutional layer and the last linear layer of a CNN model, which are the input layer and the output layer the model, respectively.

EXAMPLE 1. *Line 1 is an API call representing for the input layer of a CNN. We extract the value of the argument* input_shape *from the input layer to obtain the input size and the input channel, which are (120, 180) and 3, respectively. The output channel is obtained from the output layer 3. By extracting the first argument's value of the output layer, we obtain the value of the output channel.*

Conv2D('input_shape': [3, 120, 180], activation='relu')

...

Dense(10, activation='relu')

Since we currently focus on image problems like image classification and image regression, we use the input size, the input channel, and the output channel of image as data characteristics. First, the input size includes the height and the width extracted from the input data and mined models. Second, the input channel represents the number of primary colors in the image. Third, the output channel is the number of output categories of the data and models.

4.3 Model Matching

Model matching is a ranking system used to find good models for a certain problem by using the data characteristics. Instead of using constant default models, model matching finds the suitable models to uses them as new default models for NAS.

4.3.1 Model clustering. Model clustering uses the data characteristics of both the input dataset and mined neural networks to select appropriate initial architectures $\mathcal{N}^*$ for NAS. First of all, *Manas* clusters the mined models based on the data characteristics of the models and the input dataset. Secondly, in a meta-feature (data characteristics) space, our approach identifies closest clusters to the input dataset. Lastly, *Manas* uses all the models in the closest cluster to the dataset as the initial architectures for NAS. Formally,

we determine the initial architectures $\mathcal{N}^*$ for dataset $\mathcal{D}$ in Equation 3 as follows:

$$\mathcal{N}^* = \{C_k \mid n \in C_k, C_k \in C\} \tag{5}$$

$$C = \{\mathcal{G}(\Delta o_k, \Delta s_k) \mid \Delta i_k = 0, k \in [1, |\mathcal{S}|]\} \tag{6}$$

$$dist(n, \mathcal{D}) = \begin{cases} \min \Delta o \text{ if } \min \Delta i = 0 \\ \min \Delta s \text{ if } \min \Delta i = 0, \min \Delta o = 0 \end{cases} \tag{7}$$

In Equation 6, C is a set of clusters of neural network detected by clustering algorithm G-means [23], which uses a statistical test to automatically decide the number of clusters. Δo_k, Δi_k, and Δs_k are measured as follows:

$$\Delta i_k = |i - i_k| \tag{8}$$

$$\Delta o_k = |o - o_k| \tag{9}$$

$$\Delta s_k = \sqrt{(w - w_k)^2 + (h - h_k)^2} \tag{10}$$

Where i, o, w, and h are the input channel, the number of output classes, the input width, and the input height of the dataset, respectively. Similarly, i_k, o_k, w_k, and h_k are the input channel, the number of output classes, the input width, and the input height of a model k, respectively. The idea behind the clustering equations is that there are two types of the input channel, including 1 and 3. Therefore, we classify the neural networks that have the same input channel first. Then, we use (Δo, Δs) as the input of G-means to split the mined models into clusters. After that we identify the closest model n to the input dataset like Equation 7. The closest model is identified based on the priority of Δo and Δs that Δo takes precedence over Δs. We have tried to run our tool in different orders of priority Δo and Δs; however, this order of priority gives us the best results. The closest cluster to the input dataset is the cluster which contains the closest model. Then, we select all the models in the closest cluster to the input dataset as the initial models $\mathcal{N}^*$ for NAS shown in Equation 5.

4.3.2 Model filtering. If the number of initial models found by the model clustering approach is too large, we use model filtering to filter some models to increase the performance of *Manas*. In the searching process, *Manas* trains all the default neural networks to select the best one. After that NAS is applied to tune the selected model. Thus, with a specific time budget, the more time *Manas* spends on trying the default models, the less time it spends on NAS for model searching. To filtering neural networks, we detect the equivalent architectures. We treat each neural architecture as a graph, whose trainable layers like convolutional layers or dense layers represent vertices, connections between two trainable layers represent edges, and channels of the outgoing trainable layers represent the weights of edges between two vertices. We use Cosine similarity to measure the similarity between two vertices. M_{ij} is an element of the similarity matrix of vertex v_i of a graph G_A and vertex v_j of a graph G_B, which is measured as follows:

$$M_{ij} = \frac{\sum_{k=1}^n w_{ik} w_{jk}}{\sqrt{\sum_{l=1}^m w_{il}^2} \sqrt{\sum_{n=1}^k w_{jl}^2}} \tag{11}$$

Suppose that v_k is the common neighbor of vertex v_i and vertex v_j, we have w_{ik} and w_{jk} are the weights of edges $v_k v_i$ and $v_k v_j$,

respectively. We also have w_{il} or w_{jl} are the weights of edges that incident with each of the vertices v_i and v_j, respectively.

$$w_{pq} = \begin{cases} \text{weight of the edge between } v_p \text{ and } v_q \text{ in a graph } G \\ 0 \text{ otherwise} \end{cases}$$

$$(12)$$

We create a similarity matrix M for each pair of graphs and classify equal matrices into the same classes. In this work, we filter the models by only using the neural network whose graphs belong to the class having the highest number of matrices. In other words, we will only choose the most used architecture in the determined initial architecture for NAS. We consider the most frequent architecture to be more suitable than others because many deep learning developers repeatedly use these neural architectures. Non-trainable layers like dropout or activation are not used as the factors to detect the similarities of the neural architectures because we want to utilize the various usages of these layers to increase the performance of *Manas*. Particularly, different models can have the same graph structure because of the differences in non-trainable layers; therefore, a graph can refer to many different neural architectures.

4.4 Model Transformation

Even though the mined models are from the top star *GitHub* repositories, they are not perfect. Therefore, we transform the initial architectures by modifying or adding the batch normalization (BN) layer, the flatten layer, the activation layers, the global average pooling (GAP) layer, and the dropout layer to optimize NAS in terms of speed, errors, and the number of parameters. We choose these layers to modify the default models because of two reasons. Firstly, we have found many common patterns related to these layers. Secondly, many recent studies have shown the effectiveness of these layers on increasing the performance of deep learning models. We have created a set of pre-defined rules to transform networks based on related state-of-the-art papers and common function calls' patterns from mined models. *Manas* analyzes these architectures to determine whether the network satisfies the pre-defined rules. If these rules are satisfied, the pre-defined model transformations will be applied. The pre-defined model transformations support NAS to ignore transformations offered by our approach and focus on the other transformations. The model transformations do not work for *Auto-Keras'* initial neural architectures because those default models have already included these transformations.

4.4.1 Batch normalization layer constraint. We add a new batch normalization layer between the convolutional layer and the activation layer to increase training speed [28]. Using BN means that we modify the activations to normalize the input layer to decrease the training time. Many well-known neural architectures like ResNet [25], DenseNet [26], EfficientNet [50] use BN to increase the training speed. BN is also popular in optimizing NAS [13, 18, 61].

EXAMPLE 2. *According to Figure 5a, the original model has a convolutional layer connects to a ReLU layer, which is an activation function layer. Thus, in Figure 5b, following the batch normalization layer constraint, Manas adds a new BN layer between the convolutional layer and the ReLU layer like the following example.*

	(a) Original model		(b) Transformed model
1	(0): Conv2d(3, 32, ...)	1	(0): Conv2d(3, 32, ...)
2		2	(1): BatchNorm2d(32, ...)
3	(1): Tanh()	3	(2): ReLU()
4	(2): MaxPool2d(kernel=2, stride=2, ...)	4	(3): MaxPool2d(kernel=2, stride=2, ...)
5		5	(4): Dropout2d(p=0.5)
6	(3): Conv2d(32, 32, ...)	6	(5): Conv2d(32, 32, ...)
7		7	(6): BatchNorm2d(32, ...)
8	(4): Tanh()	8	(7): ReLU()
9	(5): MaxPool2d(kernel=2, stride=2, ...)	9	(8): MaxPool2d(kernel=2, stride=2, ...)
10		10	(9): Dropout2d(p=0.5)
11	(6): Flatten()	11	(10): GlobalAvgPool2d()
12	(7): Linear(in=32, out=32, ...)	12	(11): Linear(in=32, out=32, ...)
13	(8): ReLU()	13	(12): ReLU()
14		14	(13): Dropout2d(p=0.25)
15	(9): Linear(in=32, out=2, ...)	15	(14): Linear(in=32, out=2, ...)
16	(10): Softmax()	16	(15): Softmax()

Figure 5: Original CNN model vs transformed CNN model

	(a) Original model		(b) Transformed model
1	(0): Conv2d(3, 32, ...)	1	(0): Conv2d(3, 32, ...)
2		2	(1): BatchNorm2d(32, ...)
3	(1): ReLU()	3	(2): ReLU()

Figure 6: Example of batch normalization layer constraint

4.4.2 Global average pooling layer constraint. We use GAP to reshape the data into the correct format for fully connected layers to prevent overfitting [35]. Moreover, we use the mined models as default models for NAS; thus, the original input size of the initial models and the input size of the dataset can be different, which can cause a shape mismatch bug. However, using GAP can solve this problem since it does not care about the input shape.

EXAMPLE 3. *According to Figure 5a, the original model used the flatten layer to pass the feature map through the CNN. Therefore, in Figure 5b, following the constraint about GAP layer, Manas transforms flatten into GAP like the following example.*

	(a) Original model		(b) Transformed model
1	Flatten()	1	GlobalAvgPool2d()
2	Linear(in=32, out=32, ...)	2	Linear(in=32, out=32, ...)

Figure 7: Example of global average pooling layer constraint

4.4.3 Activation layer constraint. We investigate the patterns of usages of activation functions used in the mined model. We have found 3218 hidden layers used in 793 models, where ReLU is used 2946 times, accounting for 94.55%. Therefore, we replace the current activation layers of convolutional layers with ReLU.

EXAMPLE 4. *According to Figure 5a, the original model uses Tanh for the convolutional layer. Thus, following the constraint of the activation layer, Manas transforms Tanh to ReLU like following example like the following example.*

1	*(3): Conv2d(32, 32, ...)*		1	*(5): Conv2d(32, 32, ...)*
2			2	*(6): BatchNorm2d(32, ...)*
3	*(4): Tanh()*		3	*(7): ReLU()*

(a) Original model   **(b) Transformed model**

Figure 8: Example of activation layer constraint

4.4.4 Dropout layer constraint. We investigate the frequency of dropout [48] layers and their drop rates used in the mined model. Out of 806 times that dropout is used in hidden layers, the drop rate of 0.25 is used 385 times, which accounts for 47.8%. Out of 753 times that dropout is used in fully connected layers, the drop rate of 0.5 is used 529 times, which accounts for 70.3%. Thus, we add dropout layers with a drop rate of 0.25 and 0.5 to the hidden layers and fully connected layers, respectively.

EXAMPLE 5. *According to Figure 5a, the original model does not use the dropout layer. Therefore, in Figure 5b, following dropout layer constraint, Manas adds dropout layers into a convolutional layer with 0.25 drop rate and a dropout layer into the fully connected layer with 0.5 drop rate like the following example.*

1	*Conv2d(32, 32, ...)*		1	*Conv2d(32, 32, ...)*
2	*ReLU()*		2	*ReLU()*
3	*MaxPool2d(kernel=2, stride=2, ...)*		3	*MaxPool2d(kernel=2, stride=2, ...)*
4			4	*Dropout2d(p=0.5)*

(a) Original model   **(b) Transformed model**

Figure 9: Example of dropout layer constraint

All the transformations except dropout transformation are applied simultaneously to the mined models before searching. After the best model is selected from candidate models, dropout transformation is applied to the best model. Since adding dropout does not always improve the model's errors, we utilize NAS to identify whether using dropout is good or bad.

5 EVALUATION

5.1 Experimental Setup

We implement *Manas* by extending *Auto-Keras* [31]. All experiments use Python 3.6, with 16GB GPU Tesla V100. In these experiments, we use 8 datasets for image classification and image regression, which are obtained from *Kaggle* based on the vote count. The efficiency and effectiveness of *Manas* are evaluated in three aspects. Firstly, we evaluate the metric values including error rate and MSE that is lower the better, model complexity, and training speed of *Manas* by comparing it with *Auto-Keras*. Secondly, we evaluate the model matching approach's efficiency and effectiveness. Lastly, we evaluate the efficiency and effectiveness of model transformation and training adaptation approaches. In these comparisons, *Manas* include model matching, model transformation, and NAS. The algorithm NAS used by *Manas* explores the search space via morphing the neural architectures guided by the Bayesian optimization algorithm. Mining the models is the most time-consuming task, which took about 24 hours to complete mining all the models. However, we only do this one time, so we do not count the time to mine the model in the comparison of *Manas* and *Auto-Keras*.

5.2 Mined models

Since many models for different *Kaggle* datasets have already been published on *GitHub*, it is possible that some *Kaggle* models of the testing dataset have already been included in our mined models. To avoid this problem, for each dataset, we have examined all *GitHub* repositories of selected initial models of each dataset. There is no information showing that those initial models are specifically created for the input dataset. Moreover, for each input dataset, we have compared the initial model of *Manas* with all models from *Kaggle*. *Manas*' initial model is not one of *Kaggle*'s models. Moreover, since most of the models from *Kaggle* are published as Jupyter Notebooks, we only mine the model written as Python files.

We only collect models from the top 10,000 repositories with the most star count to ensure high-quality models. We also do sanity checks by removing incomplete models and duplicate models. Moreover, we evaluate the mined models by training them on MNIST with 50 epochs [44]. The average accuracy of the mined models on MNIST is 90.98%.

5.3 Datasets

To evaluate the performance of our method, we use **8** different image datasets collected from *Kaggle*: *Blood Cell* [39], *Breast Cancer* [40], *Flower* [37], *Intel Image Classification* (IIC) [5], *Malaria* [3], *MNIST: Ham* [36], *Sign Language Digits* (SD) [38], and *Sign Language* (SL) [51]. Our goal was to utilize more complex datasets compared to well-known datasets such as MNIST, CIFAR10, and FASHION. Most of our datasets have large sizes of images. For example, in the intel image classification [5] dataset, the image size is 150x150 while the image size of MNIST is only 28x28. Secondly, the number of images in our datasets is much larger than MNIST, CIFAR10, and FASHION. For instance, the breast cancer dataset [40] has 277,524 images. Lastly, the number of classification of the evaluated datasets is up to 10 classes. These dataset are often used for image classification task; however, we treat prediction targets as numerical values for image regression task. For example, we will treat the prediction targets of the MNIST dataset as integers ranging from 0 to 9 as numerical values to be directly used as the regression targets. We only use image datasets because *Auto-Keras* only apply NAS for image data. we divide it into two subsets, 80% of randomly selected images are used for training and the remaining images for validation.

5.4 Results

5.4.1 RQ1: How efficient is Manas? To evaluate the efficiency of *Manas*, we run both *Manas* and *Auto-Keras* on 8 datasets for image classification and image regression. We vary the search time from 2 hours to 20 hours, which are described in Figure 10. Table 1 shows the error rate, MSE, the depth, the number of parameters, and the training speed of the best models of *Manas* and *Auto-Keras*. By the best model, we mean one that has the lowest error rate or MSE after search timeout. We run both Manas and Auto-Keras 5 times for each dataset with random training and validation sets. In the table, we report the information of the lowest errors of Manas and Auto-Keras [15]. By comparing *Manas* and *Auto-Keras*, two conclusions can be drawn.

Table 1: *Manas* Classification & Regression Results

Data	Image Classification								Image Regression							
	Error rate (%)		Depth (layers)		Param # (million)		Speed (epoch/min)		MSE		Depth (layers)		Param # (million)		Speed (epoch/min)	
	AK	MN	AK	MN	AK	MN	AK	MN	AK	MN	AK	MN	AK	MN	AK	MN
Blood Cell	18.9	10.0 (↓ 47.1%)	7	6 (↓ 14.3%)	2.3	0.3 (↓ 87.0%)	5.1	8.0 (↑ 56.9%)	0.16	0.11 (↓ **31.3 %**)	21	9 (↓ 57.1%)	11.2	0.1 (↓ 99.1%)	2.7	8.4 (↑ 211.1%)
Breast Cancer	6.9	6.9 (↓ 0.0%)	21	8 (↓ 61.9%)	11.2	0.5 (↓ 95.5%)	0.2	0.5 (↑ 150%)	0.05	0.06 (↑ 20.0%)	26	7 (↓ 73.1%)	11.4	0.04 (↓**99.6%**)	0.2	0.6 (↑ 200.0%)
Flower	14.6	12.8 (↓ 12.3%)	121	11 (↓ **90.9%**)	7.0	1.5 (↑ 78.7%)	2.7	3.8 (↓ 40.7%)	0.64	0.65 (↓ 1.5%)	21	9 (↓ 57.1%)	11.2	0.4 (↓ 96.4%)	3.3	4.3 (↑ 30.3%)
IIC	8.9	8.6 (↓ 3.4%)	27	7 (↓ 74.1%)	15.5	0.5 (↓ 96.8%)	0.7	2.6 (↑ 73.1%)	0.58	0.59 (↑ 1.7%)	122	5 (↓ **95.9%**)	7.0	0.3 (↓ 95.7%)	0.7	2.4 (↑ 242.9%)
Malaria	2.3	1.6 (↓ 30.4%)	30	7 (↓ 76.7%)	20.3	0.3 (↓ **98.5%**)	1.7	4.3 (↑ 152.9%)	0.02	0.02 (↓ 0%)	21	8 (↓ 61.9%)	11.2	0.4 (↓ 96.4%)	2.4	4.7 (↑ 95.8%)
MNIST: Ham	20.6	20.3 (↓ 1.5%)	7	6 (↓ 14.3%)	0.9	1.6 (↑ 77.8%)	2.2	1.4 (↓ 36.4%)	0.81	0.73 (↓ 9.9%)	26	13 (↓ 50.0%)	11.9	0.4 (↓ 96.6%)	1.1	5.2 (↑ **372.7%**)
SD	0.5	0.0 (↓ **100.0%**)	22	11 (↓ 50.0%)	11.3	0.7 (↓ 93.8%)	8.9	66.0 (↑**641.6%**)	0.05	0.1 (↑ 100.0%)	21	12 (↓ 42.9%)	11.2	0.4 (↓ 96.4%)	48.4	78.2 (↑ 61.6%)
SL	0.0	0.0 (↓ 0.0%)	22	6 (↓ 72.7%)	11.3	0.3 (↓ 97.3%)	5.5	8.0 (↑ 45.5%)	0	0 (↓ 0.0%)	27	10 (↓ 63.0%)	12.6	7.2 (↓ 42.90%)	5.0	2.6 (↓ 48.0%)
Avg	9.1	7.5 (↓**17.6%**)	32.1	7.8 (↓**75.7%**)	10.0	0.7 (↓**93.0%**)	3.4	11.8 (↑**247.1%**)	0.3	0.3 (↓**0.0%**)	35.6	9.1 (↓**74.4%**)	11.0	1.2 (↓**89.1%**)	8.0	13.3 (↑ **66.3%**)

In the table, Avg, AK and MN represent average, *Auto-Keras*, and *Manas*, respectively. In each cell, ↓ and ↑ represent a decrease percentage and an increase percentage, respectively.
In each evaluation metric, the bold value show the best percentage change of *Manas* compared to *Auto-Keras*.

First of all, Table 1 shows that *Manas* produces models, which has lower error rate or MSE compared with *Auto-Keras*' models. The lower errors of *Manas* compared with *Auto-Keras* indicates that using mined models as the starting points for NAS can produce better models. Notably, *Manas* outperforms *Auto-Keras* by achieving **17.6%** lower error rate on average. For some problems like *IIC* and *MNIST: HAM* for image classification task, the decrease of the error rate of *Manas* compared with *Auto-Keras* is small because the models created by *Manas* and *Auto-Keras* have reached the limit of error rate. Therefore, a small decrease in error rate is a major improvement. We use the error rate as the main evaluation metric to clearly point out this improvement of *Manas* to *Auto-Keras*.

Secondly, *Manas* achieves lower errors with less complicated models compared to *Auto-Keras*. Using mined models on NAS significantly decreases the complexity of the produced models. On average, models generated by *Manas* are **75.7%** less deep and **93.0%** less wide compared to the models generated by *Auto-Keras* for image classification task. Similarly, *Manas* creates models with **74.4%** less deep and **89.1%** less wide compare to *Auto-Keras*'s models for image regression task. Simpler DNN models train faster and save more energy [27] than the complex ones. Han *et al.* have shown that reducing the number of parameters of deep learning models can reduce the training time by 3× to 4×, and energy comsumption by 3× to 7× [24]. Table 1 also shows that on average, *Manas*' models run faster than *Auto-Keras*' model **247.1%** and **66.3%** in image classification and image regression, respectively.

5.4.2 RQ2: How efficient are model transformation and optimizers modification? We create an ablation study to observe the efficiency of the model transformation and modifying optimizers via mining separately.

- *Original Manas* (OM) represents mined models + no transformation + no optimizer + NAS.
- *Transformed Manas* (TM) represents mined models + transformations + no optimizer + NAS.
- *Manas* (MN) represents for mined models + transformations + optimizers + NAS.
- *Auto-Keras* (AK) represents NAS.

We observe the error values of *Original Manas*, *Transformed Manas*, *Manas*, and *Auto-Keras* by executing them on 8 datasets for 20 hours for both image classification and image regression. To evaluate the efficiency of the model transformation, we compare *Original Manas* and *Transformed Manas*. We compare *Transformed Manas*

and *Manas* to evaluate the efficiency of optimizers modification. To evaluate the combination of all methods, we compare *Manas* and *Auto-Keras*.

Notice that we have applied the model transformation on *Auto-Keras*' models; however, this did not lead to any improvement because those default models use BN layers or Dropout layers appropriately. Figure 10b does not have the *Original Manas* series because the best models of *Original Manas* and *Transformed Manas* are the same. In other words, the model already contains all the transformation constraints, so there is no transformation applied to the model. Similarly, Figure 10e does not have the *Transformed Manas* series because the optimizers of the best model of *Transformed Manas* and *Manas* are the same. The selected model's optimizer of this problem is not available; therefore, we use the default optimizer of *Auto-Keras* for this model. From the results are shown in Figure 10, we draw four observations.

First of all, model transformation increases the performance of *Manas* in terms of errors and converge time. Figures 10a, 10c, 10f, 10g show that *Transformed Manas* achieve lower errors than *Original Manas* most of the time. The activation layers, the GAP layer, and the dropout layers contribute to the better errors of *Transformed Manas* compared to *Original Manas*. For example, the dropout layers can prevent overfitting, which decreases the errors. The model transformation also helps *Manas* to converge faster. As we can observe Figures 10a, 10c, 10f, 10g, *Original Manas* has higher errors compared to *Transformed Manas* in the first 2 hours of training, which indicate the *Transformed Manas* converge faster than *Original Manas*. *Transformed Manas* has a fast converge speed thanks to BN.

Secondly, mined optimizers help *Manas* to reduce errors and converge faster. In most of problems, the best models of *Manas* have lower errors than the best models of *Transformed Manas*. Moreover, *Manas* can converge easier with the mined optimizers. For instance, the Figures 10d, 10f, 10g shows that *Transformed Manas* has trouble converging. After 8 hours of searching, *Transformed Manas* cannot find out better models while *Manas* succeeds. *Transformed Manas* has trouble in converging since it does not have an suitable optimizer for those problems like *Manas*.

Thirdly, *Auto-Keras* may have lower errors than *Manas* in the first few hours; however, in the last hours, *Manas* often finds out better models than *Auto-Keras*. For example, Figures 10b, 10c, 10d, and 10m show that *Auto-Keras* has the errors at the beginning; however,
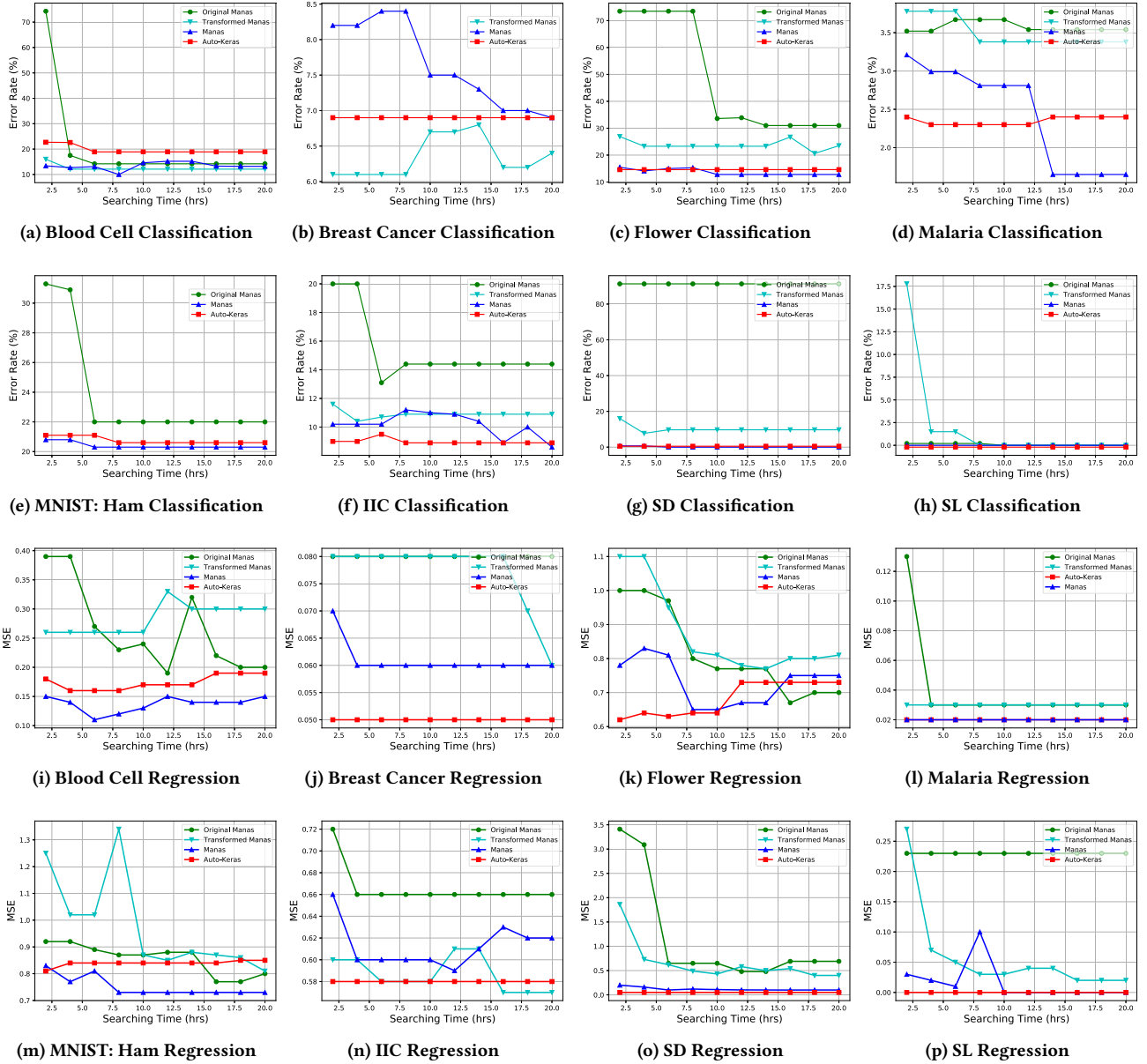
Figure 10: Error Rate and MSE of *Auto-Keras*, *Original Manas*, *Transformed Manas*, and *Manas* over time

as time gone we have not seen any improvement in *Auto-Keras* results. The reason for this problem is that *Auto-Keras* starts with very complicated models that take a lot of time to train; therefore, the number of models is searched by *Auto-Keras* are small, which decreases the chance to find out good models of *Auto-Keras*. *Manas* starts with simple models, which trains faster. Therefore, *Manas* may not be off to a good start, but in the end, it still outperforms *Auto-Keras*. The observation shows the benefit of using simple mined models in NAS.

Lastly, Figure 10 indicates that *Manas* obtains lower error rates than *Auto-Keras* in almost different time periods. As we can observe in Figures 10a and 10i, *Manas* always outperforms *Auto-Keras* in terms of error rate in the 20-hours of searching. Figure 10 shows that the longer searching time sometimes gives worse models. During the searching process, the NAS estimates the errors of searched models and selects the best one. However, the estimation may not be accurate, leading to an incorrect choice. This problem of NAS indicates the importance of using a simple model as a starting point. Simple models can train faster that increases the chance for NAS

Table 2: Efficiency of Model Matching

Data	Blood Cell		Breast Cancer		Flower		IIC		Malaria		Ham		SD		SL	
Task	IC	IM	IC	IM	IC	IM	IC	IM	IC	IM	IC	IM	IC	IM	IC	IM
Total	793															
MC	5	5	5	7	18	6	11	46	5	7	6	11	69	4	5	5
MF	-	-	-	-	-	-	-	12	-	-	-	-	12	-	-	-

In each cell, IC, IM, MM, and MF represent image classification, image regression, model clustering and model filtering, respectively. The unit of all the data in the table is the number of models.

Table 3: *Manas* vs *Auto-Keras* for Well-known Problems

Data	Error Rate (%)		Depth (layers)		Param # (million)		Speed (epoch/min)	
	AK	MN	AK	MN	AK	MN	AK	MN
CIFAR10	7	7.7 (↑ 10.0%)	21	10 (↓ 52.4%)	19.4	0.8 (↓ 95.9%)	1	2.9 (↑ 190.0%)
Fashion	5.2	11 (↑ 111.5%)	22	9 (↓ 59.1%)	14.7	1.8 (↓ 87.8%)	2.1	1.8 (↓ 14.3%)
MNIST	0.5	0.7 (↑ 40.0%)	23	6 (↓ 73.9%)	19.3	2.9 (↓ 85.0%)	1.3	3.1 (↑ 138.5%)
Avg	4.2	6.5 (↑ 54.8%)	22	8.3 (↓ 62.3%)	17.8	1.8 (↓ 89.9%)	1.5	2.6 (↑ 73.3%)

In each cell, Avg, AK and MN represent average, *Auto-Keras*, and *Manas*, respectively.

to search for more models. Since the estimation of NAS can be incorrect; thus, if we increase the number of searched model, we can increase the chance to obtain better models. For example, after 8-hours of searching, NAS goes wrong with *IIC* problem when it produces worse models than before. However, when we keep searching for new models, NAS gradually fixes the problem to obtains a good model.

5.4.3 RQ3: How efficient is model matching? The goal of model matching is not only selecting good default models for *Manas* but also reducing the number of default models. From Table 1, we observe the efficiency of model matching, when *Manas* can outperform *Auto-Keras* in many different perspectives. Table 2 shows that model matching, including model clustering and model filtering, can significantly reduce the number of default models for *Manas*.To balance the time that *Manas* spends between initial architectures and NAS, we applied the models filtering when the number of initial neural networks is larger than 40 since *Manas* often takes more than 10 hours, half of the amount of time used in our evaluations to complete training these neural networks. We use all 793 mined models as input for each problem, which may take few days to complete training. By using model matching, we decrease the number of default models by 99% on average. Taking *SD* in image classification as an example, after using DNN clustering, there are remaining 69 default models. Therefore, we use DNN filtering on this dataset to reduce the number of models of *SD* from 69 models to 12 models, which eliminates 82.6% number of the DNN model.

5.4.4 RQ4: How efficient is Manas for well-known problems? We also evaluate *Manas* with the well-known datasets like FASHION, MNIST, or CIFAR10 for the image classification task. Table 3 shows the error rate, depth, number of parameters, and speed of the best models of *Manas* and *Auto-Keras*. As can be observed, *Auto-Keras* achieve better error rates on these datasets than *Manas*; however, *Auto-Keras* produces larger models to achieve these error rates while *Manas* uses much smaller models to get close to the error rates of *Auto-Keras* in CIFAR10 and MNIST problems. Particularly,

Manas' models are 62.3% shorter and 89.9% than *Auto-Keras'* models on average, which increases the training speed of *Manas'* models by 73.3% compared to *Auto-Keras'* models. *Auto-Keras* achieves better error rates than *Manas* since *Auto-Keras* uses ResNet and DenseNet as initial models. These neural networks are well-tuned to achieve outstanding results on these datasets, which once again shows that using good initial architecture optimizes NAS.

6 LIMITATIONS AND THREATS TO VALIDITY

6.1 Limitations

In this work, *Manas* directly derives the data characteristics from the image dataset, which is limited to the image classification and image regression problems. We believe that *Manas* is not directly applied to other problems such as natural language processing (NLP) or video classification; however, our approach of mining models to identify a good starting point candidate should be applicable to any AutoML problems. For example, data characteristics of NLP problems also include the input shape, which are input time steps and the number of features, and the total number of output classes. Time steps represent the maximum length of the input sequence, which could either be the number of words or the number of characters depending on what we want. The number of features is the number of dimensions we feed at each time step. We can analyze the input layer and output layer to extract data characteristics from models. We can analyze the input dataset to obtain its data characteristics. The extracted data characteristics can be used to find better starting points for NLP problems in AutoML systems.

Manas can only work with few kinds of layers since it only uses the layers that *Auto-Keras* supports. This limitation can decrease the performance of *Manas* because if *Auto-Keras* supported more layers, we can mine more kinds of models from the software repositories.

6.2 Internal Validity

We have tried our best to obtain the results of *Manas* and *Auto-Keras* on as many datasets as possible. Because of the time limit, *Manas* is currently evaluated on 8 datasets. All the source code, trained models, datasets, and evaluation data are public for reproduction to mitigate these threats.

6.3 External Validity

First of all, *Manas* only focuses on image classification/regression problems. We rely on meta-features to find good starting points for NAS; therefore, one possible threat is that meta-features (data characteristics) do not work for other types of problems. However, Auto-Sklearn [20] and Auto-Sklearn 2.0 [19] mitigate this threat by showing that using meta-features can increase the performance of AutoML systems in terms of training speed and accuracy on structured datasets. Secondly, *Manas* only focuses on CNN. Thus, another threat is that the model transformation approach does not work for other types of models. Nevertheless, Cambronero *et al.* [10] propose AMS showing that using unspecified complementary and functionally related API components can improve the performance of AutoML systems for classical models such as Linear Regression or Random Forest. The difference between *Manas* and AMS is that AMS applies these transformations to search space while *Manas* applies these transformations to default models.

7 RELATED WORK

7.1 Neural architecture search

NAS is a technique for automatically finding appropriate neural architectures which can outperform most of the hand-designed neural networks. Specifically, NAS needs a the training dataset as the input to create a powerful neural architecture. There are many different approaches for a NAS system; however, most of them have three main components which are search space, search strategy, and optimization strategy. The search space represents the search boundary of a NAS system limiting what kinds of neural network can be searched and optimized. For instance, Baker *et al.* [4] use the convolutional architecture with pooling, linear transformations as a search space. Around the same time, Zoph *et al.* [61] use a similar search space; however, the authors use more skip connection for the search space. The search strategy is used to search appropriate models in a defined search space. There are many approaches to search models such as reinforcement learning [4, 9, 59, 61, 62] or evolutionary algorithms [46, 49, 58]. This optimization strategy supports NAS to guide the network search process. The optimization strategy evaluates a searched model with training data without training these models. Recently, many methods are proposed to optimize NAS [32, 41, 45]. In our work, *Manas* mines the neural networks from repositories to enhance the power of NAS by supporting it to have a better starting point.

7.2 AutoML

AutoML is a process for constructing an appropriate model architecture for a specific problem automatically. Many features that AutoML can provide to deep learning users, such as automated data augmentation, automated hyperparameter tuning, or automated model selection. A lot of AutoML systems have been created like Auto-WEKA [52] on top of WEKA [22, 56], Auto-Sklearn [20] on top of Scikit-learn, which support deep learning user to automate tuning hyperparameter and model selection. Some other AutoML systems can support deep learning users to automate optimizing the full ML pipeline. For instance, TPOT [42, 43] uses evolutionary programming to optimize ML software. However, a main disadvantage of these systems are very slow because of the high GPU computation requirement. Recently, *Auto-Keras* is created to handle this problem, which has implemented network morphism [12, 55] to reduce the searching time of NAS. Network morphism is a technique to morph a neural architecture without changing its functionality. Nevertheless, even though *Auto-Keras* apply network morphism technique, it still takes a lot of GPU computation. Our approach uses DNN model mining and common layer patterns to enhance the performance of AutoML system.

7.3 Mining software repositories

Cambronero *et al.* [11] proposed AL, a system that leverages existing machine learning code from repositories to synthesize final pipelines. AL can generate ML pipelines for a wide range of problems without any manual selection. Cambronero *et al.* [10] also proposed AMS, which automated generates new search space for AutoML systems by utilizing source code repositories. The new

search space is created based on an input ML pipeline, which increases the performance of AutoML systems. However, these only operate classical machine learning models, whereas *Manas* works with neural networks.

7.4 Meta-features

Auto-Sklearn [20] uses 38 meta-features of structured datasets to find a better starting point. Feurer *et al.* [19] proposes Auto-Sklearn 2.0, which reduces the number of meta-features to three, including the number of data points, the number of features, and the number of classes. The reasons for the reduction are that good meta-features are time-consuming and memory-consuming to generate. We also do not know which meta-features work best for which problem. Unlike Auto-Sklearn and Auto-Sklearn 2.0, *Manas* uses meta-features to find a better starting point for NAS, which works for neural networks. Moreover, *Manas* also proposes the meta-features, which helps NAS find a better starting point for image datasets.

8 CONCLUSION

We present *Manas*, a technique for NAS, which uses the mining technique to assist NAS. The key idea of *Manas* is to mine models from repositories to enhance NAS. In particular, we use CNN models mined from software repositories as the default model of NAS. From a large number of models, we use the model matching approach to find good models for a problem. We also apply some transformations for those models to enhance their performances. With better default models, *Manas* can increase NAS's performance, which leads to better CNN models as search results. Our experiment shows that *Manas* can produce better CNN models in terms of the error rate and MSE, the model complexity, and the training speed than *Auto-Keras*. Future work will involve extending *Manas* to problems beyond those tackled in this paper, such as video classification. We can utilize the code change patterns [16] in ML programs to improve the results. Moreover, the proposed approach of *Manas* can be applied to other automated tools, e.g., AutoAugment [14] of other components in ML pipeline [7]. The technique can also be utilized to use AutoML to address the other problems in ML, such as fairness bug [6].

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