

# Neural Network Pruning based on Filter Importance Values Approximated with Monte Carlo Gradient Estimation

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**Abstract:** Neural network pruning is an effective way to reduce memory- and time requirements in most deep neural network architectures. Recently developed pruning techniques can remove individual neurons or entire filters from convolutional neural networks, making these “slim” architectures more robust and more resource-efficient. In this paper, we present a simple yet effective method that assigns probabilities to the network units – to filters in convolutional layers and to neurons in fully connected layers – and prunes them based on these values. The probabilities are learned by maximizing the expected value of a score function – calculated from the accuracy – that ranks the network when different units are tuned off. Gradients of the probabilities are estimated using Monte Carlo gradient estimation. We conduct experiments on the CIFAR-10 dataset with a small VGG-like architecture as well as on the lightweight version of the ResNet architecture. The results show that our pruning method has comparable results with different state-of-the-art algorithms in terms of parameter and floating point operation reduction. In case of the ResNet-110 architecture, our pruning method removes 72.53% of the floating point operations and 68.89% of the parameters, that marginally surpasses the result of existing pruning methods.

## 1 INTRODUCTION

Modern deep networks contain tens or hundreds of layers and within each layer there are a plethora of parameters (He et al., 2016). While these large networks can easily be used with sufficient memory and computing power – generally via GPUs or TPUs –, their use is complicated on resource-limited devices. The commonly used – embedded or IoT – devices have limited memory and computing power, they often run on batteries, meaning that energy consumption is also an important factor in the ergonomics of these devices. To reduce the memory, energy and power consumption of these networks, pruning can be applied on them. Studies showed that more than half of the network parameters can be removed such that their accuracy is not affected (Han et al., 2016b; He et al., 2019; Sandor et al., 2020).

Network pruning can be (1) unstructured pruning and (2) structured pruning. In unstructured pruning

individual parameters – weights – are removed from the network. This leads to highly compressed architectures with more than 90% of the parameters removed (Han et al., 2016b), but the left-over parameters need to be stored in sparse matrices that require special hardware (Han et al., 2016a) and special libraries (like sparse BLAS<sup>1</sup>) to be efficient. However, these resources may not be always available. Structured pruning on the other hand focuses on removing groups of parameters (Li et al., 2016; He et al., 2018, 2019): rows or columns from parameter matrices (e.g. kernels), entire neurons or filters – we call these *functional units*. Whilst there are less removed parameters, the resulting architecture does not require any special treatment.

This paper focuses on structured pruning of neural networks: we approximate the importance of the network functional units and remove the ones with small scores. Our main contributions:

- We introduce binary random variables associated with the functional units – we call them masks –,

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parameterize and infer these *hyper-parameters* by optimizing energy functions. We apply the log-derivative trick and Monte Carlo gradient estimation during the optimization.

- We show that the inferred values for the mask parameters can be used for pruning.
- We compare our method with different state-of-the-art pruning algorithms and show that it achieves comparable results with them.

## 2 PRUNING METHODS

Pruning is an active research field of neural network compression (Blalock et al., 2020). The first pruning techniques were presented in the beginning of 90s (Le Cun et al., 1990; Hassibi et al., 1993). These methods used the Hessian of the loss function to remove parameters. However, Hessian matrix calculation requires huge computational power and a lot of memory. Due to these obstacles, it is hard to apply on modern deep neural networks.

More recent work uses the magnitude of the weights as a criterion (Han et al., 2015). After pruning, the network is fine-tuned to regain the original accuracy. The intuition is that small weights have small impact, hence their absence will not affect overall performance. Han et al. (2016b) apply the unstructured magnitude pruning and adds quantization and Huffman coding to the pipeline. This way the authors managed to reduce the VGG architecture of Simonyan and Zisserman (2014) by a factor of 49.

Li et al. (2016) introduces a filter pruning approach with sensitivity analysis: filters in layers are sorted and pruned based on their  $\ell_1$  norms. This process is stopped if the accuracy of the network drops significantly. Yao et al. (2017) introduces a compressor-critic framework, where the filter importance values are learned by a recurrent neural network. This "compressor" network takes the parameters of the original network and outputs probabilities as importance values for the network units. To train the compressor network, the expected value of the original network's loss is minimized over the probabilities generated by the compressor. He et al. (2018) presents a soft filter pruning approach where filters with small  $\ell_2$  norm are iteratively set to zero but they are retrained afterwards – together with the other filters. This provides larger optimization space and the pruning and retraining process has less dependence on the pretrained model. While previous works utilize the *smaller norm less important* criterion, He et al. (2019) prunes deep neural networks based on filter

redundancy in layers: the geometric median of the filters are calculated and the ones close to this median are removed from the network. Discrimination-aware channel pruning (Liu et al., 2021) introduces additional discrimination-aware losses to remove the most discriminative channels. The paper introduces the channel pruning as a sparsity-inducing optimization problem and solve the convex objective with a greedy algorithm.

Our method is similar to the work of Yao et al. (2017) in that we use Monte Carlo method for the gradient estimation, however we use a simple factor model – using Bernoulli distributions – to approximate the probabilities – compared to their compressor RNN network. This means our model is simpler, more intuitive and requires less resources during pruning. The way the pruning is defined could be used as a dropout mechanism (Srivastava et al., 2014) as well: while dropout uses predefined probabilities to iteratively remove the units, our method learns these values based on the importance of the units.

## 3 OUR PROPOSED METHOD

Consider a dataset  $\mathcal{D}$  that contains  $N$  image and label pairs  $(\mathbf{x}_i, y_i)_{i=1}^N$ . Let  $\mu_{\mathcal{W}}(\mathbf{x})$  denote a neural network that predicts  $y$  for the given  $\mathbf{x}$  input, where  $\mathcal{W}$  is the set of network parameters.<sup>2</sup>

We define network pruning as finding a binary mask  $\mathbf{z} \in \{0, 1\}^{|\mathcal{W}|}$  that sets part of the functional units to 0, such that the accuracy remains sufficiently high. This binary mask basically defines a subnetwork in the original network. In pruning the question is always how to find an optimal  $\mathbf{z}$  mask?

To tackle this problem, we consider  $\mathbf{z}$  a vector of *binary* random variables where each  $z^i$  indicates whether the associated unit is active or not (i.e. is *dead*). Let  $P_{\boldsymbol{\theta}}(\mathbf{z})$  denote the joint probability distribution of the vector  $\mathbf{z}$ , where  $\boldsymbol{\theta}$  are hyper-parameters and let  $s(\mu_{\mathcal{W}}(\mathbf{x}|\mathbf{z}))$  be a score function of the mask  $\mathbf{z}$  and the input  $\mathbf{x}$ . As defined above,  $\mathbf{x}$  is an input image, but it could be an image batch as well.

Our goal is to learn the  $P_{\boldsymbol{\theta}}(\mathbf{z})$  probability distribution function such that the network score is as high as possible. More formally, we want to maximize the expected value of the score with respect to the probability distribution  $P_{\boldsymbol{\theta}}(\mathbf{z})$ :

<sup>2</sup>Here  $\mathcal{W}$  could "simply" be any vectorized form of the parameters, but in the experiments we used e.g. filter matrices as individual components and assigned a single mask bit to this subset.

$$\begin{aligned}\boldsymbol{\theta} &= \underset{\boldsymbol{\theta}}{\operatorname{argmax}} E_{\mathbf{z} \sim P_{\boldsymbol{\theta}}(\mathbf{z})} [s(\mu_{\mathcal{W}}(\mathbf{x}|\mathbf{z}))] \\ &= \underset{\boldsymbol{\theta}}{\operatorname{argmax}} S(P_{\boldsymbol{\theta}}),\end{aligned}\quad (1)$$

where we defined the expected score as  $S(P_{\boldsymbol{\theta}})$ .

To maximize the expected value, we need to optimize the parameterized probability distribution by gradient ascent:

$$\boldsymbol{\theta}_{k+1} = \boldsymbol{\theta}_k + \alpha \nabla_{\boldsymbol{\theta}} S(P_{\boldsymbol{\theta}})|_{\boldsymbol{\theta}_k} \quad (2)$$

However, calculating the expectation is not possible due to the large number of mask combinations (in total  $2^{|\mathcal{W}|}$  possibilities) where the network score should be evaluated. Instead, we approximate the gradient by developing Monte Carlo estimators (Robert and Casella, 2010).

$$\begin{aligned}\nabla_{\boldsymbol{\theta}} S(P_{\boldsymbol{\theta}}) &= \nabla_{\boldsymbol{\theta}} E_{\mathbf{z} \sim P_{\boldsymbol{\theta}}(\mathbf{z})} [s(\mu_{\mathcal{W}}(\mathbf{x}|\mathbf{z}))] \\ &= \int_{\mathbf{z}} \nabla_{\boldsymbol{\theta}} P_{\boldsymbol{\theta}}(\mathbf{z}) s(\mu_{\mathcal{W}}(\mathbf{x}|\mathbf{z})) d\mathbf{z}\end{aligned}\quad (3)$$

In Eq. (3) the gradient of the probability distribution appears ( $\nabla_{\boldsymbol{\theta}} P_{\boldsymbol{\theta}}(\mathbf{z})$ ). This can be expressed using the log-derivative trick:

$$\nabla_{\boldsymbol{\theta}} P_{\boldsymbol{\theta}}(\mathbf{z}) = P_{\boldsymbol{\theta}}(\mathbf{z}) \nabla_{\boldsymbol{\theta}} \log P_{\boldsymbol{\theta}}(\mathbf{z}), \quad (4)$$

and replacing Eq. (4) into Eq. (3), we get:

$$\begin{aligned}\nabla_{\boldsymbol{\theta}} S(P_{\boldsymbol{\theta}}) &= \int_{\mathbf{z}} P_{\boldsymbol{\theta}}(\mathbf{z}) \nabla_{\boldsymbol{\theta}} \log P_{\boldsymbol{\theta}}(\mathbf{z}) s(\mu_{\mathcal{W}}(\mathbf{x}|\mathbf{z})) d\mathbf{z} \\ &= E_{\mathbf{z} \sim P_{\boldsymbol{\theta}}(\mathbf{z})} [\nabla \log P_{\boldsymbol{\theta}}(\mathbf{z}) s(\mu_{\mathcal{W}}(\mathbf{x}|\mathbf{z}))],\end{aligned}\quad (5)$$

where we have the product of a probability distribution and a function that we can evaluate. Since in Eq. (5) we have an expected value, we can rewrite the expression and obtain that  $\nabla_{\boldsymbol{\theta}} S(P_{\boldsymbol{\theta}})$  is the expected value of the score function times the gradient of the log probability distribution.

Using the Monte Carlo method, we can approximate the gradient by deriving a general-purpose estimator using  $N$  samples from the  $P_{\boldsymbol{\theta}}(\mathbf{z})$  distribution:

$$\nabla_{\boldsymbol{\theta}} S(P_{\boldsymbol{\theta}}) \approx \frac{1}{N} \sum_{i=1}^N \nabla_{\boldsymbol{\theta}} \log P_{\boldsymbol{\theta}}(\mathbf{z}^i) s(\mu_{\mathcal{W}}(\mathbf{x}|\mathbf{z}^i)) \quad (6)$$

Since the estimated gradient can have higher variance, the convergence of the optimization can be slower. To tackle this, we apply simple variance reduction techniques following the work of Yao et al. (2017): we subtract the moving average of the score from the actual score value and divide it by  $\sqrt{1-v}$ , where  $v$  denotes the variance of the score. We used the above formulation in our algorithm (see Section 3.3).

### 3.1 Score of the Network

Eq. (1) contains a function that scores the network when a  $\mathbf{z}^i$  mask is applied on it. This score has to be high when the network performs "well" and low otherwise. To calculate the score for a given  $\mathbf{z}^i$  mask, the network is evaluated on a random image batch from the validation set (meaning that we train the probability distribution on the validation set). We experiment with 3 different score functions and measure how quickly the probabilities converge to 0 or 1 and how the pruning affects the network accuracy.

The **loss-score** function is used from the work of Sandor et al. (2020):

$$s_i = \frac{\mathcal{L}_{\max} - \mathcal{L}_i}{\mathcal{L}_{\max} - \mathcal{L}_{\min}}, \quad (7)$$

where  $\mathcal{L}_i$  is the network loss with the  $\mathbf{z}^i$  mask,  $\mathcal{L}_{\min}$  and  $\mathcal{L}_{\max}$  are the minimum and maximum values among the  $\mathcal{L}_i$  losses. This way the score is 1 when the network loss is the smallest (network performs well) and 0 when the loss is the highest.

The **acc-score** uses the accuracy as a score function. Let  $\mathcal{X}$  denote a random batch of images from the validation set and  $\operatorname{Acc}(\mu_{\mathcal{W}}(\mathcal{X}|\mathbf{z}^i))$  denote the network accuracy on the image batch when  $\mathbf{z}^i$  mask is applied on it. Then the score of the  $\mathbf{z}^i$  mask is simply:

$$s_i = \operatorname{Acc}(\mu_{\mathcal{W}}(\mathcal{X}|\mathbf{z}^i)) \quad (8)$$

The **exp-acc-score** applies a scaling and an exponential function on the accuracy:

$$s_i = \exp\left(\frac{\operatorname{Acc}(\mu_{\mathcal{W}}(\mathcal{X}|\mathbf{z}^i))}{\beta}\right) \quad (9)$$

While the accuracy can be used as a score function, it cannot capture fine-grained details: if  $\mathbf{z}^i$  and  $\mathbf{z}^j$  differs only in a single value, the scores could be very close to each other. This results small difference between the gradients as well, resulting slow convergence during the factor model optimization. The exp-acc-score function increases the distance between the scores when the accuracy values are similar.

The experiments with the different score functions are presented in section 4.1.

### 3.2 Probability Distribution

An important question is how to represent the probability distribution over the set of random variables  $\mathbf{z}$ ? For simplicity, we assume independence between the elements and define the probability distribution as a product of Bernoulli distributions:

$$P_{\boldsymbol{\theta}}(\mathbf{z}) = \prod_i p_i^{z_i} (1-p_i)^{1-z_i}, \quad (10)$$

where the probability that  $z^i = 1$  depends on the  $\theta_i$  parameter:  $P_{\theta_i}(z^i = 1) = p_i = \sigma(\theta_i)$ .

Using the factor model from Eq. (10), the log probability in Eq. (6) can be written as:

$$\begin{aligned} \log P_{\theta}(\mathbf{z}) &= \log \prod_i p_i^{z^i} (1 - p_i)^{1 - z^i} \\ &= \sum_i z^i \log p_i + (1 - z^i) \log(1 - p_i) \end{aligned} \quad (11)$$

### 3.3 Network Pruning Algorithm

To use the factor model as a probability distribution, we assume independence between the network units (neurons and filters). Since this assumption is clearly not true in case of a multilayer network, we assume independence only between units from the same layer. This way, we prune the network layer by layer, learning separate probability distributions for each layer. A formalized version is presented in Algorithm 1.

Algorithm 1: Network pruning.

**Require:**  $\mu_{\mathcal{W}}$  pre-trained network

- 1:  $l \leftarrow$  index of first or last layer
- 2: **while** stopping condition not met **do**
- 3:   initialize  $P_{\theta}(\mathbf{z})$  for layer  $l$
- 4:   **for** some predefined steps **do**
- 5:     sample  $\{\mathbf{z}^1, \dots, \mathbf{z}^N\}$  masks from  $P_{\theta}(\mathbf{z})$
- 6:     calculate  $\nabla_{\theta} S(P_{\theta})$  ▷ from Eq. (6)
- 7:     update  $\theta$  ▷ from Eq. (2)
- 8:   prune layer  $l$  based on  $P_{\theta}(\mathbf{z})$
- 9:   fine-tune  $\mu_{\mathcal{W}}$
- 10:  $l \leftarrow$  next layer index

## 4 EXPERIMENTS

We analyze our pruning algorithm on different architectures trained on the CIFAR-10 (Krizhevsky et al., 2009) dataset.

Similar to (Frankle and Carbin, 2019), we use a small VGG-like (Simonyan and Zisserman, 2014) architecture with two convolutional layers, a max pool layer and three fully connected layers. Both convolutional layers contain 64,  $3 \times 3$  filters while the fully connected layers have 256, 256 and 10 neurons respectively. In the hidden layers ReLU activation function is used. The network is trained on the CIFAR-10 dataset for 10 epochs (using Adam optimizer (Kingma and Ba, 2015) with  $2 \times 10^{-4}$  learning rate and early stopping condition) and reaches 69.76% accuracy on the validation dataset and 68.95% on the test dataset.

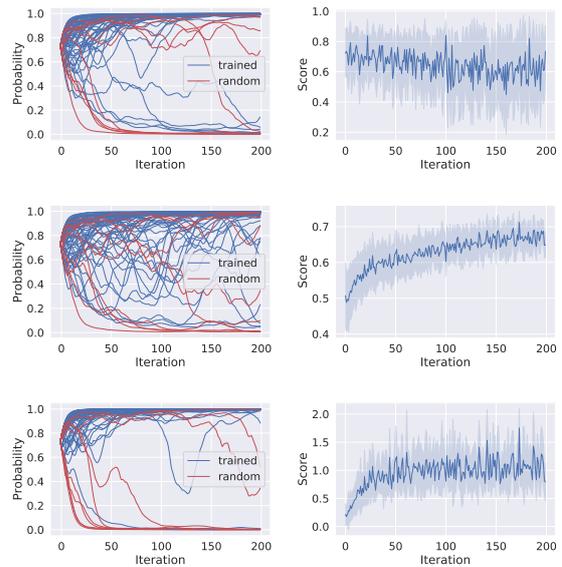


Figure 1: Left: Inferred probabilities for the 74 filters in the first convolutional layer (10 of them are filters with random weights). Right: standard deviation of the scores during the training process. Top: loss-score function, middle: acc-score function, bottom: exp-acc-score function.

### 4.1 Score Function

To analyze the different score functions presented in section 3.1, we prune the trained VGG-like network: First, we insert 10 extra filters (containing random weights) in the first convolutional layer. Then we prune the network and measure the true positive rate: how many extra filters are pruned from the network. A score function is better if the pruning algorithm can identify more filters with random weights.

In case of each function, we train our factor model from Eq. (10) for 200 iterations and remove the filters with small probabilities (we use 0.2 as a threshold). We repeat the process until at least 10 filters are removed from the network and measure the validation and test accuracy of the pruned networks. For each of the 3 score functions we repeat the experiment 5 times and report the average validation and test accuracy, the number of removed filters and the true positives (number of removed filters that contain random weights). Results are reported in Table 1. As the table shows, the pruning method detects part of the inserted random filters. However, the true positive rate and the pruned network accuracy varies. When the network loss and accuracy is used to calculate the score (first 2 rows in the table), on average more than 11 filters are removed from the network. However, around 6-7 of them are true positives, that means 30–40% of the removed units contains trained weights. In these cases the network accuracy decreases quite heavily as well:

Table 1: Pruning results using the three different score functions. The validation and test accuracy, true and false positives (Tp and Fp) are averages from 5 different experiments. We insert 10 random filters in each experiment, thus the ideal case would be that the algorithm prunes only those 10 filters.

Score function	Val. acc. (%)	Test acc. (%)	Tp	Tp+Fp
loss-score	68.84 (↓ 0.91%)	67.76 (↓ 1.18%)	6.8	11.2
acc-score	67.99 (↓ 1.76%)	66.72 (↓ 2.22%)	6.0	11.0
exp-acc-score	70.11 (↑ 0.35%)	68.56 (↓ 0.38%)	8.4	10.2

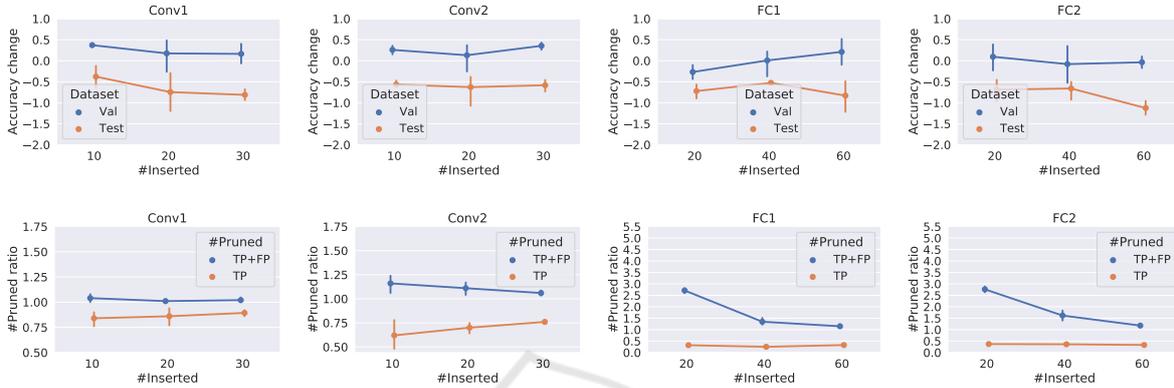


Figure 2: Up: Accuracy change of the pruned networks when different number of filters (10, 20, 30) or neurons (20, 40, 60) are inserted into the layers. Down: true and false positive rates of the pruning.

0.91% and 1.76% on the validation set and 1.18% and 2.22% on the test set. Results are more promising in case of the exponential score function: on average 8.4 from 10 random filters are found by the algorithm and the pruned network accuracy increases by 0.35% on the validation set and decreases only 0.38% on the test dataset.

In Figure 1, we present the dynamics of the inferred probabilities and scores during the factor model training. The left figures show how the mask probabilities converge to 0 and 1 during this process. It is easy to see that the convergence is much faster in case of the loss-score and exp-acc-score functions (top and bottom). Using these functions, the values quickly moves close to 0 or 1 depending on the filter’s importance. Slow convergence with the acc-score function is related to the value of the scores (right figure): during the factor model training the standard deviation of the scores decreases. This means that the difference between the gradients decreases as well, leading to slow convergence. The standard deviation of the scores is much higher in case of the loss-score and exp-acc-score functions. This is important, since as the probabilities start to converge (the network accuracy approximate the original accuracy), a small change on the  $\mathbf{z}^i$  mask leads to higher change on the score – meaning the difference between the gradients are higher.

While the dynamics of the probabilities with loss-score and exp-acc-score functions are similar (top and

bottom figures), the values converges a bit quicker with the latter. Moreover, as Table 1 shows, the result are also better with this function. This is because exp-acc-score function provides consistent values during the training process while the values of loss-score depends on  $\mathcal{L}_{\min}$  and  $\mathcal{L}_{\max}$ : the same  $\mathbf{z}_i$  mask can have different scores in different training iterations as  $\mathcal{L}_{\min}$  and  $\mathcal{L}_{\max}$  changes - which is very likely, since the masks are randomly sampled. This varying score affects the gradients that leads to slower convergence.

## 4.2 Pruning Randomly Inserted Filters from Trained Networks

Based on the presented experiments in section 4.1, we select the exp-acc-score function and examine the pruning algorithm on different layers of the VGG-like network. Similar to section 4.1, we insert randomly initialized filters and neurons into the trained VGG-like network, however, we vary the number of random filters and the target layers as well. In case of the convolutional layers, we insert 10, 20 and 30 random filters while in the fully connected layers we insert 20, 40 and 60 random neurons one after another. We repeat each experiment 5 times and report the average change of the validation and test accuracy (between the pruned and original networks), the average number of removed filters (TP + FP) and the average number of the removed random filters (TP). Figure 2

presents the results of this experiment.

The first 4 figures show the accuracy change of the pruned networks compared to the original network accuracy (trained network, no random filters inserted). Since the probability distribution is optimized to maximize the expected value of the scores calculated on the validation set, the validation accuracy of the pruned network outperforms the validation accuracy of the original network in almost all 4 cases. This validates that Eq. (6) correctly estimates the gradient and the gradient method can increase the expected value of the score. In case of the test accuracy, a drop between 0.5% and 1.2% can be detected, and the gap is slowly increases with the number of inserted filters. This means the overfitting becomes stronger as more random filters are inserted into the network.

Figures from the bottom show the number of removed filters (TP+FP) and number of removed random filters (TP) among the 4 experiments. The first observation here is that filters in the first layer are more important than filters in the second layer: while the true positive rate is around 80% in the first layer, it is between 60 – 75% at the second layer. This means that more trained filters are removed from the second layer, but the accuracy values remain similar (or even better in case of the second layer). At the fully connected layers the true positive rate decreases below 50%. These layers contain more than 256 neurons but only a fraction of them contributes to the correct output. The pruning algorithm can “pick” almost randomly from the pool and the accuracy still remains near to the original accuracy.

In our second experiment we use the same network but instead of targeting only a single layer we insert randomly initialized filters into all the network layers. This problem is more challenging since filters have influence to each other through the layers. As presented in section 3.3, we apply sequential pruning: the algorithm prunes the layers one by one and repeats the process until the target size is not reached (Figure 3).

The results show that the algorithm can find more than 75% of the random filters in the convolutional layers and more than 50% of the neuron in the fully connected layers. While these values are similar with the results of the previous experiment, here the test accuracy drop increases to 1.5 – 3% (Figure 3, top).

### 4.3 Pruning the ResNet Architecture

Next, we are testing the ResNet (He et al., 2016) network. This is an efficient CNN architecture that applies residual blocks and “shortcut connections” for better propagation of the error signal. The residual

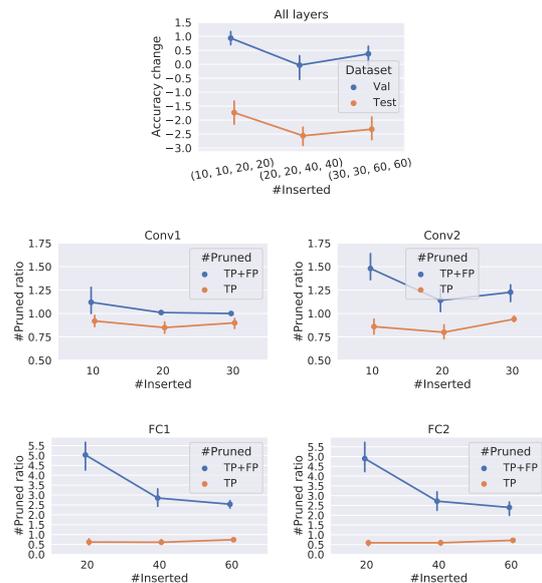


Figure 3: Up: Accuracy change of the pruned networks when different number of units are inserted into the four layers. Middle and down: true and false positive rates of the pruning at different layers.

block contains two sets of convolutional, batch normalization and ReLU layers, such that the output of a layer is fed into the input of the next layer. To prune the units in this residual block, we insert mask layers following the work of Sandor et al. (2020): the first mask layer is inserted after the first ReLU layer while the second mask layer is inserted before the shortcut connection.

**Training Details:** We experiment with the ResNet-32, 56 and 110 architectures. We train the networks following the work of He et al. (2016), with the following modifications: we change the initial 0.1 learning rate to 0.01 and 0.001 at epochs 100, and 150 and stop the training at 200 epochs. During training we apply cropping and horizontal flip as data augmentation. The network is trained on a 45K train set, a 5K validation set is used for training the parameterized probability distribution.

**Pruning Details:** Pruning is applied based on the algorithm presented in section 3.3. We start the process from the network’s first layer and calculate the probabilities layer by layer. In each layer, we train the probability distribution (Eq. 10) in 200 iteration such that in each iteration 50 masks are sampled and their corresponding scores are calculated. In each iteration the gradients are estimated – using Eq. (6) – and the parameters are updated – using Eq. (2). After 200 iterations units with small probability are turned off – here we follow the work of Sandor et al. (2020): we drop the least important units such that the accuracy

Table 2: Comparison of pruned ResNet with the results from the literature.

ResNet	Method	Accuracy (%)			↓(%)	
		Baseline	Pruned	Diff. ↓	FLOPs	Params.
32	SFP (He et al., 2018)	92.63	92.08	0.55	41.5	41.24*
	FPGM (He et al., 2019)	92.63	92.82	-0.19	<b>53.2</b>	<b>53.2*</b>
	LFE (Sandor et al., 2020)	92.97	92.42	0.55	46.4	49.35
	Ours	92.97	92.29	0.68	50.22	43.65
56	(Li et al., 2016)	93.04	93.06	-0.02	27.6	13.7
	SFP (He et al., 2018)	93.59	93.35	0.1	47.14	52.6*
	ThiNet (Luo et al., 2019)	93.8	92.98	0.82	49.78	49.67
	FPGM (He et al., 2019)	93.59	93.49	0.1	47.14	52.6*
	LFE (Sandor et al., 2020)	93.44	93.18	0.26	57.64	<b>68.14</b>
	Adapt-DCP (Liu et al., 2021)	93.74	93.77	-0.03	<b>68.48</b>	54.80
	Ours	93.44	93.08	0.36	64.22	57.79
110	(Li et al., 2016)	93.53	93.3	0.23	38.6	32.40
	SFP (He et al., 2018)	93.68	93.86	-0.18	40.8	40.72*
	FPGM (He et al., 2019)	93.68	93.85	-0.17	52.3	52.7*
	LFE (Sandor et al., 2020)	94.05	93.48	0.57	63.68	60.08
	Ours	94.05	93.45	0.6	<b>72.53</b>	<b>68.89</b>

\*Parameter drop percentage is not reported in the paper. These values are calculated from other available information (e.g. "40% of the filters are selected").

drop on the validation dataset is less than 1.0%. After a layer is pruned, we apply fine-tuning for 10 epochs.

Finally, when no more filters can be removed, we retrain the network for 100 epochs by setting the learning rate to 0.1 and decrease to 0.01 and 0.001 at epochs 40 and 60.

**Results:** We report the pruning results of the ResNet architecture in Table 2. The algorithm removes 43.65% of the parameters and 50.22% of the floating point operations (FLOPs) from the ResNet-32 architecture. While these values outperform the results of He et al. (2018) in terms of parameter and FLOPs reduction, remains below the results of He et al. (2019) and Sandor et al. (2020). While the pruning results are modest with the smaller ResNet, they are more promising with the ResNet-56 and ResNet-110 versions. We manage to remove 64.22% of the FLOPs and 57.79% of the parameters from the ResNet-56 with only 0.36 accuracy drop. These values outperform most of the results reported by the papers selected for comparison: only the FLOPs reduction result of Liu et al. (2021) and the parameter reduction result of Sandor et al. (2020) can outperform our algorithm. In case of the ResNet-110 architecture, our pruning algorithm removes more than two thirds of the floating point operations (72.53%) and the parameters (68.89%). These values surpass the results of the other papers with a significant margin.

## 5 CONCLUSIONS

In this paper, we described a structured pruning algorithm that approximates the importance probability of network units using Monte Carlo gradient estimation. To calculate the importance values, we introduced a function that scores the performance of different subnetworks. A subnetwork is defined as a binary masks that specifies the active and inactive units in the network. Given a set of score and their corresponding subnetwork – binary mask –, we maximize the expected score of the network by optimizing the probability distribution of the masks using estimated gradients from the Monte Carlo method. Based on the importance values, our method applies pruning on the network and produces a compressed model with parameters stored in smaller, dense matrices. We showed the effectiveness of our pruning algorithm on the CIFAR-10 dataset with a small VGG-like architecture as well as on different versions of the ResNet architecture. The experiments show that our algorithm has comparable results with current state-of-the-art pruning methods.

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