

BACHELOR THESIS

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Adversarial Examples Generation for Deep Neural Networks

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Introduction

The effort to automate various processes in our lives has always been one of the key concepts of scientific research. The automation of mechanical work has already been solved to great extent by advances in engineering. On the contrary, automated processing of information has been solved just partially. Well defined tasks i.e., tasks with fully defined behaviour can be solved and the challenge lies just in effectivity of the solution. On the other hand, models solving tasks based on raw real word data, human level input and output or some degree of fuzziness are yet to be found or, if they exist, suffer from several shortcomings. One of those shortcomings is vulnerability to adversarial examples i.e., artificially created inputs misinterpreted by those models. In this thesis, we will cover the task of generating adversarial examples for the models used for classification in computer vision.

TODO 800 chars artificial intelligence -¿ machine learning -¿ sota deep learning

TODO 800 computer vision and classification tasks

1. Preliminaries

In this chapter we will present theoretical background of various subjects relevant to this writing. We will give a compact overview of the task of image classification in computer vision with concrete examples of available datasets. We will cover various artificial intelligence paradigms with emphasis on deep learning for image classification and image generation, and evolutionary algorithms as a space search method. Adversarial example generation methods will be discussed in the following, separate chapter for their importance with respect to this work.

1.1 Classification Tasks in Computer Vision

The field of computer vision (CV), sometimes percieved as a part of the artificial intelligence, examines the machine processing of imaging data. The basics of the field were laid down in the 1960s, the golden years of artificial intelligence, with ambitious goal to build systems with universal understanding of visual concepts. A multitude of subtasks of this ultimate goal were solved to a large extent; however the task as a whole remains to be solved at the time of writing. The computer vision is experiencing boom once again with the advent of deep learning and graphics processing unit (GPU) accelerated computation. See Szeliski [2010] for in depth introduction to the field of computer vision.

Based on the previous approaches in CV, such as the use of convolutional filters, in conjunction with increasing computational power and advancements in machine learning, such as reinvention of gradient backpropagation, the convolutional neural networks emerged (see section 1.2). Those models outperformed former state of the art methods i.e., support vector machines with RBF kernels or scale-invariant feature transform with Fisher vectors, in terms of accurracy (Krizhevsky et al. [2012a]) and enabled wider range of tasks to be solved. The tasks that are being solved by convolutional neural networks in CV are image classification, object detection, semantic segmentation, image generation, style transfer, image captioning, etc. We will elaborate more on image classification tasks, available datasets and respective state of the art results.

1.1.1 Task Definition

The task of image classification lies in assigning one and only one label l from the set of possible output labels L to each input image I from the space of all possible images \mathbf{I} of which we often think as a unit hypercube i.e., $[0,1]^{h \cdot w \cdot c}$ with h, w, c being input images height, width and number of channels (depth) respectively. Task with input varying in shape can be transformed to the canonical classification task by adding preprocessing step which unifies shape e.g., reshaping using bilinear interpolation or conversion to common colour space (see Szeliski [2010]), or by building a multitude of solutions – each targeted at specific shape of input data. Several derived tasks e.g., hierarchical classification with labels forming tree-like structure, or multilabel classification with examples belonging to multiple categories exist, but are out of scope of this writing.

Figure 1.1: MNIST samples

1.1.2 MNIST like tasks

The whole family of datasets for benchmarking of image classification solutions has been created. For the sake of interchangeability all of them share the same input size of $28 \times 28 \times 1$ (rectangular grayscale image) and often same dataset size and structure with 60,000 training and 10,000 testing examples. Those datasets, particularly the original MNIST itself, became widely used benchmarking datasets for they are rather small in terms of the number of examples and their shape, enabling fast prototyping and experimentation.

MNIST The dataset comprises of handwritten digits – a selected subset of the National Institute of Standards and Technology [2010] database. The database consists of approximately 800,000 handwritten alphanumeric characters written by 3600 writers, namely high school students and the United States Census Bureau employees. The subset has been chosen so that the amount of samples written by students compared to the Census Bureau employees is equal in each dataset split. Moreover the sets of writers chosen for training samples and testing samples are disjoint. The chosen samples are normalized with the following steps – each character is reshaped to fit $20 \times 20px$ patch preserving the aspect ratio and the center of mass of this patch is aligned with the center of larger $28 \times 28px$ patch creating final sample (see fig 1.1). The classification accuracy of over 99% was reached by Lecun et al. [1998] and their LeNet model. Due to increasing capabilities of machine learning approaches, MNIST has become obsolete and is no longer a valid benchmarking dataset for performance comparison nowadays.

EMNIST The Extended MNIST (EMNIST) dataset contains alphanumeric characters and it is seen as an extension of the MNIST dataset. It comes from the same database and was created by mimicking the steps used for MNIST sample preprocessing. However, the procedure differs sligtly and consequently EMNIST is not a superset of MNIST. The dataset comes in several layouts. The EMNIST Complete consists of approximately 700,000 training samples and 115,000 testing samples each classified to one of 62 classes (10 digits, 26 lower-case and 26 uppercase letters). The EMNIST Merge is a variation of the complete EMNIST with visually ambigous letter classes e.g., C, K, P, S merged, leading to 47 classes in total. The EMNIST Balanced comprises of 112,800 training and 18,800 testing samples in 47 classes (using merging), with equal sample frequency across classes. Appart from those datasets, there are EMNIST Digits, EMNIST Letters and EMNIST MNIST, dataset sharing the layout of original MNIST dataset (see Cohen



Figure 1.2: Fashion MNIST samples

et al. [2017]). Despite being substantially harder task than MNIST, EMNIST is not widely used.

Fashion MNIST The Fashion MNIST by Xiao et al. [2017a] shares the layout of MNIST dataset, having the same number of same shaped train and test examples as well as the same number of classes. The dataset comprises of grayscale thumbnails of pieces of clothes in ten classes T-shirt/top, Trouser, Pullover, Dress, Coat, Sandal, Shirt, Sneaker, Bag, Ankle boot (see fig 1.2). This dataset seems to be promising replacement of the original MNIST as it is defining harder and more computer vision relevant task (images of real world objects rather than manmade symbols), while preserving moderate hardware requirements and model complexity demands.

1.1.3 Photo datasets

Due to the need to classify real world imagery i.e., data with substantial amount of noise, lacking quality or having non-trivial composition such as multiple objects in the scene, varying viewing angles or conditions, or heterogenous background), the datasets of real world photographs were created. Those datasets range from collections of thousands of low resolution thumbnails in dozens of classes to millions of reasonably large pictures in hierarchical systems of classes.

CIFAR The CIFAR dataset comes in two versions, namely CIFAR-10 and CIFAR-100, named after number of classes present. Both datasets share the same example shape i.e., 32×32 (rectangular color image) and dataset layout comprising of 50,000 training and 10,000 testing images with balanced class occurence. CIFAR-10 consists of following classes – airplane, automobile, bird, cat, deer, dog, frog, horse, ship, truck (see fig 1.3), while CIFAR-100 introduces hierarchical system of 20 superclasses e.g., fish, small mammals, tress, large carnivores each subdivided into 5 classes e.g., fish – aquarium fish, flatfish, ray, shark, trout (see Krizhevsky [2009]).

ImageNet ImageNet is large image database, providing labeled data for image classification and object detection. The images are labeled according to WordNet® hierarchy (see Miller [1995]) and the whole database aim to provide approximately 1000 images for each class. ImageNet serves as a benchmark for current state of the art image classification architectures, which became popular being winning submisions to ILSVRC challenge (challenge based on ImageNet



Figure 1.3: CIFAR 10 samples

dataset, see Russakovsky et al. [2015]) – consisting of 1.2 million images in 1000 classes. Current state of the art results achieved by *NASNet* reach 3.8% top-5 error and 17.3% top-1 error (see Zoph et al. [2017]).

1.2 Deep Learning in Image Classification

The deep learning is a branch of machine learning characterized by utilization of multilayered neural networks. Due to the complexity of deep networks and solutions tailored to solve tasks with fuzzy real world data, deep learning approaches are often not backed up by strong methematical theory. However they are proving successful in solving those tasks and provide current state of the art solutions in the field of computer vision, speech recognition and synthesis, machine translation, etc. See Goodfellow et al. [2016] for in depth introduction into the deep learning.

1.2.1 Machine Learning

Machine learning is an artificial intelligence paradigm based on the concept of knoledge extraction from observed states or experience, followed by application of obtained knowledge to new observations.

A computer program is said to learn from experience E with respect to some class of tasks T and performance measure P, if its performance at tasks in T, as measured by P, improves with experience E. Mitchell [1997]

More formally the machine learning solution, often referred to as model M, can be percieved as the approximation of probability distibution P, optionaly conditional, of random variable \mathbf{x} , representing the task being solved. The model is often implemented by a parametric differentiable composite function M_{θ} and the process of finding the solution for a given task is called training and lies in finding the optimal set of parameters θ . The process of training relies on training data i.e., set of observed states, samples of random variable \mathbf{x} . The optimal set of parameters is searched for using maximum likelihood estimate principle i.e., the estimate of likelihood that the probability distribution of model $P_{M_{\theta}}$ matches the probability distribution of random variable \mathbf{x} (see Flach [2012]). This process is implemented as a minimization of loss function (see section 1.2.2), often using

gradient descent algorithm (section 1.2.3). Given the characteristics of the training data, machine learning tasks can be split into the following two categories – supervised and unsuperivsed

Training data of the *supervised* task take form of (x, y) i.e., input – output or querry – answer example pairs. The goal is to approximate the conditional probability distribution P(y|x) (classification, encoding). In case of reinforcement learning, the dataset itself is not available, and the ground truth comes in form of full description of the dynamic environment in which the model operates.

Training data of the unsupervised task take form of singletons x i.e., samples of unknown distribution. The the goal is usually to find unknown structure (clustering), be able to generate new samples i.e., approximate the probability distribution P(x), or transform the data to some latent space with usefull properties (compression, encoding, feature extraction).

1.2.2 Loss Functions

Given the probability distribution \bar{P} of the training data – samples of unknown distribution P, and probability distribution $\bar{P}_{M_{\theta}}$, the loss function quantifies the likelihood of $\bar{P}_{M_{\theta}}$ matching the distribution P, using the estimation \bar{P} . Frequently used loss functions are Kullback-Leibler divergence, often referred to as cross-entropy, used for the purpose of classification tasks; mean-squared error for regression tasks, or more advanced loss functions such as conditional random fields (CRF) loss or connectionist temporal classification (CTC) loss. Only the two former will be discussed in detail for their relevance to this writing.

Kullback-Leibler divergence For the discrete probability distributions, the *KL-divergence* has the following form.

$$D_{KL}(\bar{P} \parallel \bar{P}_{M_{\theta}}) = -\sum_{i} \log \bar{P}(i) \frac{\bar{P}_{M_{\theta}}}{\bar{P}(i)}$$

$$\tag{1.1}$$

It holds that KL-divergence is always non-negative and equal to zero if and only if $\bar{P} = \bar{P}_{M_{\theta}}$. It is worth to note that KL-divergence is not symmetric, hence it is not a distance measure.

Mean squared error The *mean-squared error* (MSE) loss function is widely used in regression tasks as it measures the sum of variance and squared bias of the model error. It takes the following form.

$$MSE(\bar{P}, \bar{P}_{M_{\theta}}) = \frac{1}{n} \sum_{i=1}^{n} (\bar{P}(i) - \bar{P}_{M_{\theta}})^{2}$$
(1.2)

It holds that MSE is always non-negative and equal to zero if and only if $\bar{P} = \bar{P}_{M_{\theta}}$.

1.2.3 Optimizers

The optimizer controls the process of finding optimum of the loss function. Unlike classical methods of mathematical optimization working with fully defined functions (or probability distributions they generate), optimizers in machine learning

work with finite set of samples of unknown distribution. As a result, finding the global optimum on the training set is not the preferable goal, as it brings the issue of overfitting – relying on characteristics specific to the training data \bar{P} which are not representative in context of the whole unknown distribution P. To fight overfitting, methods of regularization are often employed (see section 1.2.4). Due to the hardware limitations, it is often not possible for the optimizer to work with the whole training set. As a result, sampling of dataset is employed in the form of batching, and the process of training is performed in epochs – iterations over the whole training set. Optimizers frequently used during training of deep learning models are stochastic gradient descent (optionally with momentum or Nesterov momentum), RMSProp, AdaGrad, Adam, etc.

Gradient Descent The gradient descent is a space search method. It navigates the space of the parameters of the machine learning model by iteratively taking steps in the opposite direction of the gradient of the loss function, hence minimizing it. The step size is driven by *learning rate* parameter. The algorithm does not guarantee finding global optimum. When using batching, we talk about stochastic gradient descent for we have access only to the approximation of the loss function in each step.

```
Algorithm 1 Stochastic gradient descent [optionally with Nesterov momentum]
```

```
\alpha \leftarrow \text{learning rate}
[\beta \leftarrow \text{momentum rate}, v \leftarrow 0]
\mathbf{repeat}
\text{Sample a minibatch of } m \text{ training examples } (x^{(i)}, y^{(i)})
[\theta \leftarrow \theta + \beta v]
g \leftarrow \frac{1}{m} \nabla_{\theta} \sum_{i} L(M_{\theta}(x^{(i)}), y^{(i)})
[v \leftarrow \beta v - \alpha g]
\theta \leftarrow \theta - \alpha g
\mathbf{until early stopping criterion is met}
```

The SGD with *Nesterov momentum* of rate β (algorithm 1.2.3) runs as follows. We set the initial momentum to zero, then each step, until we have reached the goal, we first change parameters according to the accumulated momentum, we estimate gradient of the loss function, then we update the momentum and perform step according to the estimated gradient.

Adam The Adam optimizer by Kingma and Ba [2014] is one of optimizers derived from basic SGD. It addresses the shortcomings of the stochastic gradient descent, mainly high variance in loss function gradient estimation and inability to effectively navigate certain landscapes. Adam keeps track of the first and the second moment estimates of the loss function gradient, using exponential moving average.

Adam (algorithm 1.2.3) estimates the mean of gradient in the same way as SGD with momentum. Moreover, the second moment estimate provides adaptive learning rate for each parameter in θ resulting in faster convergence during training and more stable behaviour.

Algorithm 2 Adam optimizer

```
\begin{array}{l} \alpha \leftarrow \text{learning rate} \\ \beta_1 \leftarrow \text{mean decay rate}, \beta_2 \leftarrow \text{variance decay rate} \\ s \leftarrow 0, r \leftarrow 0, t \leftarrow 0 \\ \textbf{repeat} \\ g \leftarrow \frac{1}{m} \nabla_{\theta} \sum_{i} L(M_{\theta}(x^{(i)}), y^{(i)}) \\ t \leftarrow t + 1 \\ s \leftarrow \beta_1 s + (1 - \beta_1) g \\ r \leftarrow \beta_2 r + (1 - \beta_2) g^2 \\ \hat{s} \leftarrow s/(1 - \beta_1^t) \\ \hat{r} \leftarrow r/(1 - \beta_2^t) \\ \theta \leftarrow \theta - \frac{\alpha}{\sqrt{\hat{r} + \epsilon}} \hat{s} \\ \textbf{until early stopping criterion is met} \end{array}
```

1.2.4 Regularization

As stated before, the issue of overfitting is adressed by means of regularization – methods, that improve model generalization ability (performance on unseen data), while not neccessarily changing the performance on training data. Frequently used regularization algorithms are: early stopping, p-norm regularization, dataset augmentation, ensembling, dropout, etc. See Goodfellow et al. [2016] for in depth overview.

The early stopping is a policy terminating training process if model performance on validation set visibly stalls or worsens. The p-norm regularization adds additional term to the loss function computing p-norm of model parameters, forcing them to maintain mean close to zero and low variance; l2 and l1norms are often used. The dataset augmentation stands for careful application of transformations to training data providing mean of enlarging training set, and consequently improving model performance; shifting, flipping, random cropping, addition of random noise are broadly used augmentation methods, they need to be tailored to task specifically, to preserve label of augmented example. Ensembling lies in training multiple models with different initialization, different training process or even with different architectures and employing a voting scheme for those models. This often leads to better results; however this approach comes with significant computational power demands. The dropout masks internal featuremaps (see section 1.2.5) randomly during the training process, forcing the layers to perform well even with noisy inputs, pushing the decision boundary between classes far from presented examples.

1.2.5 Layers

Neural networks are often thought of as layered structures with layers being transormations of tensors. The deep neural networks are composed of tenths or even hundreds of those layers. Various types of layers serve for feature extraction, non-linear transformation, normalization, reshaping, etc. We will refer to tensors being passed between layers as featuremaps or hidden representation, slices of featuremaps along the last axis will be called channels. See Goodfellow et al. [2016] for more.

Dense The most basic and arguably the first layer invented is a dense layer, computing biased linear combination of inputs. Given the input vector x_{in} of length s_{in} , parameters W as weight matrix of shape $s_{out} \times s_{in}$ and b bias vector of length s_{out} , the output of dense layer of size s_{out} corresponds to (1.3).

$$L_{W,b}(x_{in}) = Wx_{in} + b \tag{1.3}$$

Activation functions Linear combinations itself are not able to form strong enough models. Cybenko [1989] proved, that added non-linearity, in their case sigmoid (1.4) function, theoretically enables construction of universal approximators. This theorem, often referred to as the universal approximation theorem, has been later proven for other non-linear functions e.g., tanh, rectified linear unit ReLU (1.5) and its modifications, etc. (see Sonoda and Murata [2015]). Those are widely used in machine learning as an activation functions.

$$\sigma(x) = \frac{1}{1 + e^{-x}} \tag{1.4}$$

$$f(x) = \begin{cases} x, & \text{for } x > 0\\ 0, & \text{for } x \le 0 \end{cases}$$
 (1.5)

Activation function as a neural net layer is then straightworward elementwise application of it to the input featuremap.

Convolution A multitude of machine learning tasks process data e.g., images, natural language samples or audio, with inherent spatial properties – mainly context locality. Convolutional layers use fixed size sliding window (kernel) to perform discrete convolution, in order to globally extract local context. Given the input tensor X_{in} of shape (w, h, d_{in}) , the 2-dimensional convolution (Conv2D) of output depth d_{out} , with kernel tensor K of shape $(w_K, h_K, d_{out}, d_{in})$ and bias vector b of length d_{out} corresponds to (1.6).

Conv2D
$$(X_{in})_{x,y} = \sum_{i=1}^{w_K} \sum_{j=1}^{h_K} W_{i,j} X_{in_{x+i-\lceil w_K/2 \rceil, y+j-\lceil h_K/2 \rceil}} + b$$
 (1.6)

As stated in (1.6), each output value $X_{out_{x,y}}$ is computed as a sum of the elemntwise multiplication of convolution kernel with equally shaped part of input featuremap centered around the position corresponding to $X_{out_{x,y}}$. N-dimensional convolution for different n can be computed analogically. As the convolution crops the featuremap proportionally to the size of the kernel, padding of the input tensor may be used as a preprocessing step, to maintain constant hidden representation size. To perform featuremap downsampling, we can compute convolution only in evenly spaced positions, the distance between those positions is referred to as stride.

Pooling Pooling represents another way of downsampling. Unlike strided convolution, pooling is often parameterless and instead of performing convolution with trainable kernel, it applies given function to sliding window (channel-wise). Average and maximum are frequently used pooling functions – layers are then called max-pooling and avg-pooling respectively.

Normalization This type of layer fights internal covariate shift, undesirable effect of model hidden representations not having stable probability distribution, by centering and scaling featuremaps to have zero mean and unit variance. Usage of normalization often results in faster and more stable training as trainable layers need not compensate for, possibly significant, changes in properties of outputs of preceding layers. Normalitation layer keeps track of first and second moment estimate often using exponential moving average. There is a multitude of normalization layers varying in a way those moments are computed. Normalization yields moderate regularization effect. According to the span of normalization, we recognize three types of it: batch, layer and group.

The batch normalization takes place over all but last axis (channels) of featuremap. The gain of employing batch normalization depends on training batch size, as bigger batches benefit more (see Ioffe and Szegedy [2015]). The layer normalization takes place over all but first axis (examples) of featuremap. Unlike batch normalization, the performance is not compromised when using smaller batch size. Finally, the group normalization takes place over multiple channels per example, forming multiple groups of features. This effectively forces the features of similar moments, ideally features with similar semantics, to share the same group (see Wu and He [2018]).

1.2.6 Known Architectures

The advent of deep learning and its soaring popularity has been boosted by outstanding results achived with deep learning models, beating then state of the art methods by a large margin. Arguably AlexNet was the first architecture to show the potential of convolutional neural networks, followed by gradually deeper architectures VGGNet, ResNet, DenseNet, etc. Apart from image classification, deep learning models dominated image segmentation, natural language processing tasks e.g., machine translation, speech recognition, synthesis etc. Architectures for these tasks are out of scope of this writing.

AlexNet AlexNet by Krizhevsky et al. [2012b] is a convolutional neural net using convolutions of size $11 \times 11, 5 \times 5$ and 3×3 , max-pooling for downsampling, ReLU as an activation function and dropout after dense layers for regularization. It has 8 layers consisting of 62.3 million trainable parameters. SGD with momentum is used for training. The AlexNet scored 15.3% top-5 accuracy in ILSVRC 2012 competition.

VGGNet With twice as many layers as AlexNet and more then double parameter count, VGGNet by Simonyan and Zisserman [2014] is another well known architecture. It uses exclusively 3×3 convolutions, max-pooling for downsampling, ReLU as an activation function and dropout for regularization after dense layers. SGD with momentum is used for training. With 16 layers and 138 million parameters, VGGNet scored top-5 accuracy of 7.3%.

ResNet Trying to train even deeper models, the issue of vanishing gradients arose. He et al. [2015] came up with residual shortcuts, connections bypassing

convolutions, which enabled gradient flow. With that improvement, their ResNet-152 model, with 152 layers and parameter count of 60.2 millions, scored top-5 accuracy of 3.6%. It uses 3×3 convolutions, average pooling, ReLU activation and dropout after dense layers. Moreover batch normalization between each convolution and activation is used. It is trained using SGD with momentum.

DenseNet Further exploiting the performance gain of residual shortcuts usage, Huang et al. [2016] created DenseNet architecture. Residual connections do not bypass just one layer at a time, but connects every preceding layer to the current one, forming a so called $dense\ block$, those blocks are interconnected with $transition\ layers$, which perform downscaling of featuremaps. Several improvements as bottlenecking and feature compression using 1×1 convolutions are introduced. It uses 3×3 convolutions, ReLU activation, dropout and batch normalization between each convolution and activation, and SGD with Nesterov momentum for training.

1.3 Evolutionary Algorithms as Space Search Algorithm

Evolutionary algorithms represent one of heuristic search algorithms and are inspired by the process of natural evolution. Following the work of Darwin [1859], we can think of evolution as of the process of gradual improvement of individuals of some population, with respect to a measure of fitness, given by environment the population lives in. The key principle is, according to Darwin, the natural selection i.e., the survival of the fittest, which together with process of reproduction and heredity leads to propagation and spread of promising traits in succeeding generations. Formalising those principles, we obtain heuristic search algorithm by simulating natural evolution. For in depth introduction see Engelbrecht [2007].

Assume a task with objective O on space S. Then population P is a subset of space to be searched $(P \subset S)$, with individual i being an element of population $(i \in P)$, hence element of search space S i.e., possible solution. Fitness function $F: i \mapsto \mathbb{R}$ is a real valued measure of fitness depending on individual, often function of the objective function O of the task to be solved i.e., $F: O(i) \mapsto \mathbb{R}$. As an operator $Op: P_{in_1}, P_{in_2}, \ldots, P_{in_n} \mapsto P_{out}$ we define a function converting one or more populations to a new one, often in element-wise (per individual) manner. Epoch is then a one step of evolutionary algorithm consisting of application of sequence of operators to current population P_t , producing new population $P_{t+1} = Op_n(Op_{n-1}(\ldots Op_1(P_t)))$.

The evolutionary algorithm stops if the stopping criterion is met i.e., if some individual of current population, or the whole population itself, reach sufficient performance with respect to objective O.

It is worth to note that population and consequently individuals are often not sampled from space S directly but rather conveniently encoded in a format suitable for evolutionary algorithm, the representation of individual often takes form of array of features, often referred to as genome with individual features called traits or genes.

1.3.1 Operators

The ability of evolutionary algorithms to effectively search given space lies in application of suitable operators. Those operators can be divided into three categories – reproduction operators e.g., cross-over which combines several individuals into new one; mutation operators, which add random perturbations to individuals, with the idea of exploring genes not present in current population; and finally selection operators mimicking the natural selection based on fitness of individuals.

Assuming the population P of individuals being vectors of genes $i \in G_1 \times G_2 \times \cdots \times G_l$ of length l, with G_k being the *domain* of gene g_k . The following paragraphs elaborate on several most frequently used operators.

N-point cross-over takes two distinct individuals i, j producing offspring u, v by randomly partitioning parent vectors into n+1 segments, and swapping odd ones. Crossover with n+1=l is called the universal crossover and instead of partitioning the genome to segments and swapping them based on parity, it decides for each gene, whether to swap it or not randomly. For individuals being multidimensional matrices, the partitioning can be defined analogically, by partitioning each axis separately. Various modifications of cross-over are used based on the type of genes. For real valued ones, averaging (optionally weighted) instead of swapping is sometimes used.

In case of random mutation, each gene g_k of individual i is mutated, with probability p_{gene} by drawing a sample from given distribution – this may be uniform distribution over gene domain G_k (unbiased mutation), normal distribution centered in g_k with given variance in case of real valued genes (biased mutation), etc.

Roulette wheel selection represents one of stochastic selection operators. Given population P of individuals i_1, i_2, \ldots, i_n we can construct an imaginary roulette wheel, where each individual is represented by a slot of size proportional to its fitness, normalized by aggregate fitness of the whole population. Selected population is then sampled from this roulette wheel. As any individual may be chosen multiple times, small populations suffer from high variance in roulette wheel sampling. This issue is addressed by stochastic universal sampling algorithm.

Another stochastic selection operator is tournament selection. Given target size of output population s_{out} and tournament size s_t , the output population is selected by running s_out independent tournaments i.e., taking the fittest individual (winner) from random input population sample of size s_t . In case of fitness function not corresponding to real fitness of the individual in terms of proportionality and scaling, tournament selection outperforms roulette wheel selection, as it depends only on the ordering of the fitness of the individuals.

2. Adversarial Examples for Deep Learning Models

Performance of deep learning models is not perfect and certain amount of input data gets missclassified naturally due to the insufficient accuracy of models. However Szegedy et al. [2013] showed that misclassification can be achieved artificially by adding small perturbations designed to fool the model to the previously correctly classified examples. Those inputs are called adversarial examples. We will focus on adversarial examples for image classification task. The task of generating adversarial example for input x, model M_{θ} , such that $M_{\theta}(x) = l$ is the correct label, lies in finding the perturbation η , such that $M_{\theta}(x + \eta) = M_{\theta}(x') = l'$, with l' being target class for the adversarial attack.

2.1 Taxonomy

We can characterise adversarial attacks by following, mutually independent, properties i.e., specificity, scope, adversary's knowledge and perturbation constraints.

The specifity of adversarial attack is defined by the target class l'. In case of targeted attack, the target class l' corresponds to one specific classification category. On the other hand, in case of non-targeted attack, l' corresponds to all but the ground truth class l and adversarial example is than any $x' = x + \eta$ such that $M_{\theta}(x') \neq l$.

The scope of adversarial attack lies in number of original examples to be attacked by single η . Singleton attack stands for attack on single example x such that $M_{\theta}(x + \eta) = l'$. Multi attack stands for attack on arbitrary number of examples $x \in X$ of the same class such that $\forall x \in X : M_{\theta}(x + \eta) = l'$. Finally, universal attack seeks single η such that $\forall x : M_{\theta}(x) = l \implies M_{\theta}(x + \eta) = l'$

We distinguish between three cases of adversary's knowledge. White-box attack assumes full knowledge of targeted model i.e., architecture, weights and training data as well as parameters of the optimizer being used for training. On the other hand black-box attack assumes no additional information about targeted model, appart from the classification predictions and respective probabilities. Finally Surrogate attack treats the targeted model as a black-box, yet has full access to substitute model i.e., model sharing some characteristics of targe model e.g., task it solves, training data, architecture, output probability distribution, etc.

Regarding the perturbation constraints of the attack, we distinguish the following three cases. Unconstrained attack with no limitations on perturbation η , constrained attack, with limited intensity of perturbation η allowed, with respect to some measure, and finally optimized, where the perturbation intensity is minimized. Frequently used intensity measures are l1 and l2 norms, or PASS by Rozsa et al. [2016] and SSIM by Wang et al. [2004] – visual similarity measures.

2.2 Gradient based methods

Thanks to inherent property of deep learning models – differentiability, gradient based methods are easy to be used and perform well in white-box and reasonably well in surrogate scenarios. More of those methods have been invented, namely L-BFGS using Broyden–Fletcher–Goldfarb–Shanno optimization algorithm, fast gradient sign method and its modifications, feature adversary, one-pixel attack, etc.

Fast Gradient Sign Method The fast gradient sign method computes gradients of loss function of the target model with respect to target example x and class l'. The sgn of gradients is taken and multiplied by ϵ , the step size and subtracted from x (targeted attack) – this way we move the target image in direction of rising target class prediction (2.1). In case of non-targeted attack, gradients are computed with respect to source class l and instead of subtraction we use adition to move the target image in direction of falling source class prediction.

$$x' = x - \epsilon sgn(\nabla_{\theta} L(M_{\theta}(x), l'))$$
(2.1)

The FGSM can be used iteratively with $x_t = x'_{t-1}$. Optional clipping after each step may take place in case of constrained attack. Versions of FGSM with momentum exists – generating the adversarial example the same way as the training of deep learning models optimizes the parameters.

2.3 Generative models

Appart from methods based on backpropagation, usage of deep generative models for adversarial example generation has been researched. Zhao et al. [2017] trained generative adversarial network on task dataset mapping random latent space to images – generator G and invertor mapping images to latent space of generator I – inverter. The adversarial examples are generated by finding z' close to z = I(x), such that $M_{\theta}(G(z')) = l'$. This approach generates adversarial examples more natural to human observer as we are perturbing the hidden representation and not the final output.

3. Our Solution

In our solution we focus on design of a black-box adversarial attack, for TODO reason. We will shortly elaborate on available black-box attack strategies and then we will present and discuss our pure evolutionary algorithm based approach and finally suggest a hybrid method employing evolutionary algorithm with surrogate model as an informed operator.

3.1 Related Work

Regarding the black-box adversarial attacks for deep learning models, only a few methods has been proposed so far. Papernot et al. [2016] suggest an approach converting black-box attack to surrogate attack, by synthesizing dataset for the surrogate model to be trained on. Narodytska and Kasiviswanathan [2017] performs local-search of the neighbourhood of targeted image to estimate the gradient of targeted model. This estimate is later use to guide the generation of adversarial example. The former approach is later extended by Ilyas et al. [2018], who employ natural evolutionary strategies to estimate target model gradients with respect to targeted image. The adversarial example is generated using the gradient descent algorithm on the estimated gradient. Finally, Vidnerová and Neruda [2016] used evolutionary algorithms to perform adversarial attack mostly on shallow machine learning models or models based – we will base our solution on their approach, evaluating the method on state of the art deep learning models and proposing a multitude of improvements to the evolutionary algorithm based strategy.

3.2 Pure Evolutionary Algorithms

As stated above, the method we will be improving is based on use of evolutionary algorithm. Vidnerová and Neruda [2016] propose following configuration of the evolutionary algorithm. For the targeted model M_{θ} , with input $[0,1]^{h\times w}$. The individual i is encoded as a vector of pixel intensities ($i \in [0,1]^{hw}$). The two-point crossover is selected as a reproduction operator and for the mutation, the biased random mutation with Gaussian distribution is employed. Selection is performed using tournament selection of size 3 and relies on fitness function which is an arithmetical average of Euclidean distance between source image and individual (adversarial example intensity) and Euclidean distance between prediction of individual under M_{θ} and targeted predictions. The evaluation, performed on several shallow models trained on MNIST dataset, runs 10,000 generations of evolutionary algorithm with population size of 50, allowing for at most 500,000 accesses of the targeted model.

We propose the change of encoding if the individual to 2D matrix to maintain spatial relations of data—images. Those relations are further exploited by selected operators, namely cross-over. The suitability of this change is supported by the fact, that vast majority of deep learning image classifiers is based on use of convolutional layers, which extracts the local context of processed data. We

also experimented with perturbations smaller in terms of shape, which where added to the targeted image using bilinear interpolation, while proving moderatly successful, they did not outperformed full sized scenario.

The use of two point cross-over is preserved with necessary changes to support data of higher dimensionality. In our case the cross-over swaps randomly selected rectangular regions of parent individuals i.e., image crops, to produce offspring. We experimented with the use of universal crossover without success – we assume, that the degradation in performance is again caused by the inherent properties of convolutional layers, where the partial change of area covered by convolutional kernel yields unstable results in convolutional layer activation, while the change of whole rectangular region maintains the local response of the layer.

The idea of biased mutation is preserved as well, but we change the probability distribution to binomial distribution i.e., discrete probability distribution, maintaining the discrete characteristics of intensity of pixels. As a result, biased mutation driven by binomial distribution mimics shifting of the intensity of the pixel by several shades of gray, while at the same time maintaining the beneficial properties of normal distribution.

For the optimization we use two objectives – the score of prediction of targeted class, which needs to overcome 0.5 for successful attack; and MSE of individual with respect to targeted image, to quantify the intensity of perturbation. We choose MSE over Euclidean distance for it account more for big pixel-wise differences.

As a result, the selection operator is changed to selection suitable for multiobjective optimization, namely non-dominated sorting algorithm. For that matter, we choose NSGA-II with crowding distance, which uses concept of Paretooptimality to construct a so called non-dominated fronts – gropus of mutually non-dominated individuals. Furthermore NSGA-II employs secondary sorting algorithm (in our case crowding distance) allowing for more stable selection of individuals (see Deb et al. [2002] for more). We assume this approach superior to average of objectives for the issues with different scaling of prediction error and perturbation intensity. Appart from that, we suggest the use of clipping on the intensity of perturbation if it overcome certain admissible value, combining the constrained and optimized adversarial attack paradigm.

Finally the dual clipping takes place. Firstly, to maintain the individual in the range of a unit cube (the searched space) and secondly to maintaint the pixel-wise perturbation in the allowed range – this is employed to reject individuals with few significantly shifted pixels, as those do not show sufficient visual similarity to targeted image.

We evaluate our solution on the Fashion MNIST dataset, allowing each run of evolutionary algorithm to perform at most 65000 evaluations of targeted model M_{θ} , we chose this amount to get comparable results with white-box and surrogate approaches trained on training set of Fashion MNIST with size 60,000. See section 4 for results and comparison.

3.3 Hybrid Methods

4. Experiments

In order to empirically assess the performance of proposed solutions we have carried out a multitude of experiments on the Fashion MNIST dataset. This particular dataset was chosen for it is simple enough to enable thorough examination of various scenarios using k-fold cross-validation and measurements of results on statistically significant number of examples (often whole Fashion MNIST test set), yet more complex then MNIST and the like (1.1.2). Following the results of Fashion MNIST benchmark (Xiao et al. [2017b]) we have chosen two high scoring model architectures of varying nature – namely:

SimpleNet Simple wide shallow network with three convolutional layers of depth 32, 64 and 128 respectively and one hidden dense layer of size 128. Each convolutional layer uses square kernel of size 3 and stride 1 with same padding and is followed by batch normalization and ReLU activation. Max pooling with kernel of size 2 and stride 2 is added in-between each pair of convolutions for subsampling. Flattened output of last convolutional layer is followed by ReLU activated hidden dense layer and final output layer of size corresponding to the number of classes. Dropout with rate 0.3 is added after each max pooling and in front of hidden and output dense layers. Model is trained with Adam optimizer (Kingma and Ba [2014]) with default (recommended) parameters for 128 epochs. Initial learning rate 0.001 is divided by 5 in 50% and 75% of training process. To the best of our knowledge the name SimpleNet does not refer to any well-known model architecture and was chosen for easier distinction of used architectures when refering to them. Whole model has approximately 900k trainable parameters.

DenseNet Specifically DenseNet-BC of depth 52 with growth rate k=8 and 0.5 compression factor, representing deep narrow convolutional network. Dropout of rate 0.2 is used as proposed by authors as well as recommended training parameters i.e., SGD with Nesterov momentum of 0.9 and learning rate of 0.1 divided by 10 in 50% and 75% of training process which takes, again, 128 epochs. Whole model has approximately 120k trainable parameters. See Huang et al. [2016].

Both models use weight decay of 0.0001 as additional regularization method as well as standard input preprocessing - mean subtraction and standard deviation division with channel-wise precomputed moments on training data (TODO citation). Batch size of 128 is used during training and no early stopping method is employed.

Training of both architectures is carried out using stratified 10-fold cross-validation resulting in 20 target models trained with varying random initialization on differing training sets. From the test set accuracy boxplot 4.1, we can see that densenet outperforms simplenet by approximately 0.8%. The confusion matrices of median models are very similar for both architectures.

Experiments overview In the following sections we are going to examine following approaches towards generating adversarial examples for image classifiers.

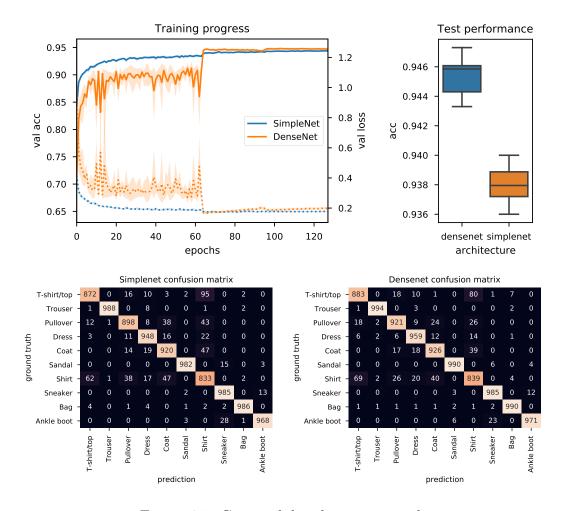


Figure 4.1: Cross validated training results

Firstly, white-box fast gradient sign method attack serving as both implementation sanity check and baseline is carried out. Next the transferability of adversarial examples is examined using surrogate models. Those approaches include transfering adversarial examples between models trained on the same dataset – the most naive approach, training surrogate model on outputs of target model and finally training focused surrogate models as a binary classifiers for specific classification category in one-vs-all manner. For each approach based on use of surrogate models, the transferability of adversarial examples between models of the same architecture as well as models of differing architectures is assessed. Appart from pure gradient methods, evolutionary algorithms are employed as a space search method, generating adversarial examples for target model. Finally using both gradient based methods and heuristic space search together, the performance of hybrid methods is measured.

For some experiments one *SimpleNet* and one *DenseNet* model is chosen, taking median models with respect to accuracy, we will refer to them as *median SimpleNet/DenseNet*. In the same manner a single class is chosen, median in terms of model prediction accuracy and attack performance, for use in some experiments – this proves to be the *dress* class for Fashion MNIST dataset.

Each approach is tested with either non-targeted and targeted singleton attack to each class under following conditions – attack is constrained to a maximum of 0.1 pixel-wise difference between targeted and adversarial examples, which corresponds approximately to 26 shades of gray difference in either direction.

4.1 Fast Gradient Sign Method Approaches

Iterative FGSM is applied to each example in Fashion MNIST test set. FGSM with step size of $\frac{1}{255}$ is used, which corresponds to shift by one shade of grey in resulting adversarial example. The following experiments compare adversarial attack performance for various model architectures, target model knowledge available to adversary and character istics of the attack itself.

4.1.1 White-Box Attack

The white-box iterative FGSM is run against the target models directly. Following box plots show the distribution of the number of steps (y-axis) needed to be taken to obtain adversarial example from source label (x-axis) to target (plot title), those results are collected on the whole Fashion MNIST test set using cross validated target models i.e., for each class – architecture pair, the median step count for each test set example is taken, resulting in 1000 datapoints. Only valid datapoints i.e., those representing successfully generated adversarial examples are used to render each box. Presented heatmaps than show success rate of attack for each source – target label pair for given architecture (plot title). Blank cell in heatmap represents maximal possible value i.e., 1000.

Results Following from presented results 4.2 we can see that white box attack against single example proves to be successful in both targeted and non-targeted scenarios for each examined architecture. *Simplenet* architecture proves to be slightly more resilient both in terms of success rate and iterative FGSM step

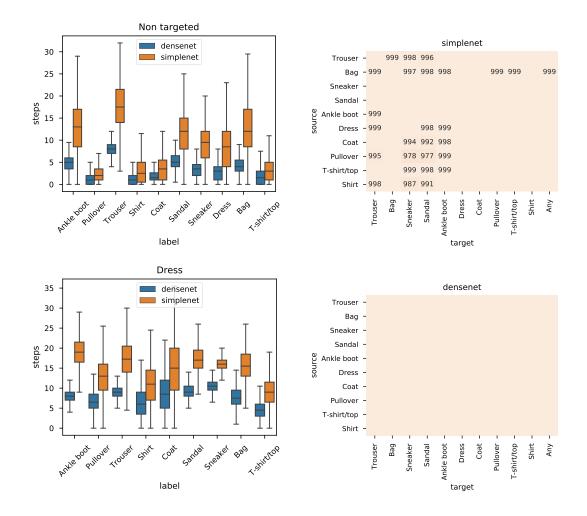


Figure 4.2: FGSM White-box

count. Consulting confusion matrix of targeted models, more confused classes are easier to attack in non-targeted scenario. In targeted attack the distribution of step count approximately corresponds to the distribution of targeted class confusion (column *dress* in confusion matrices).

4.1.2 Surrogate attack

Following experiment explores the transferability of adversarial examples using surrogate models. We compare the performance of following four scenarios.

plain surrogate model is model trained on the same dataset (not necessarily using the same train, validation splitting) 4.3

full surrogate model is trained to fit output probability distribution of targeted model using whole training dataset 4.4

reduced is similar to *full* surrogate, except for being trained only on a fraction of training data to mimic limited ability to call targeted model (when obtaining output probability distribution) 4.5

binary surrogate model is similar to *full* surrogate in terms of data size, but is trained as a binary classifier between one chosen class and all other classes,

allowing for targeted attack to selected class and non-targeted attack from it 4.6

Results Appart from varying approaches towards surrogate training, we compare performance of surrogates being of the same architecture versus differing architecture from the targeted model. From attached plots (4.3, 4.4, 4.5, 4.6), we can see that

TODO turn bullet list into paragraphs??, evidence for feature reuse?

- SimpleNet is more resilient to attacks than DenseNet (left column of plots scoring lower values than right column), we assume that this might be caused by extensive feature reuse in case of DenseNet tampering with some feature influence prediction of multiple classes (those dependent on the modified feature)
- it is easier to generate adversarial examples among visually similar classes types of shoes (sneaker, sandal, ankle boot), clothes covering the chest (dress, coat, pullover, t-shirt/top), while class unlike any other (trouser) is resilient to adversarial attack; from this point of view, bag category might seem as an outlier, being very easily attacked we think this might be again caused by feature reuse, as from the point of visual similarity, it is close to both shoes and pieces of clothes
- asymmetry in performance of cross architecture attacks is interesting phenomenon, with *DenseNet* target model attacked by *SimpleNet* surrogate being far more successful than its counterpart; same architecture attack, on the other hand proves to be successful in both cases, outperforming cross architecture attacks; this again may be caused by feature reuse with *DenseNet* with more categore independent features unable to attack *SimpleNet* with more category dependent features
- comparable performance of plain surrogates (trained on the whole dataset, unaware of target model distribution) and reduced surrogates (trained on reduced dataset, aware of target model) both inferior to full surrogates (trained on the whole dataset, aware of target model) which confirms that both increasing amount of data and increasing knowledge about targeted model improve success rate of adversarial attack

TODO place plots before analysis? to backreference?? Experiments will be carried out for those scenarios:

- EA (black-box, pure EA)
- hybrid (surrogate, EA + pre-trained surrogate)
- hybrid (surrogate, EA + on-demand-trained surrogate)

Each combination of following options will be tested:

- targeted vs non-targeted
- single image vs multi-image vs generalization (unseen images)

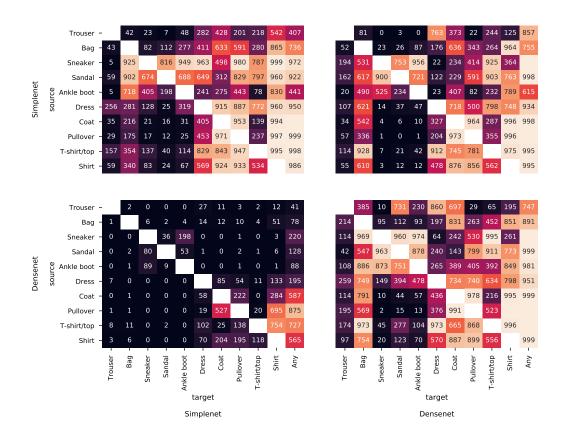


Figure 4.3: Plain surrogate success rate

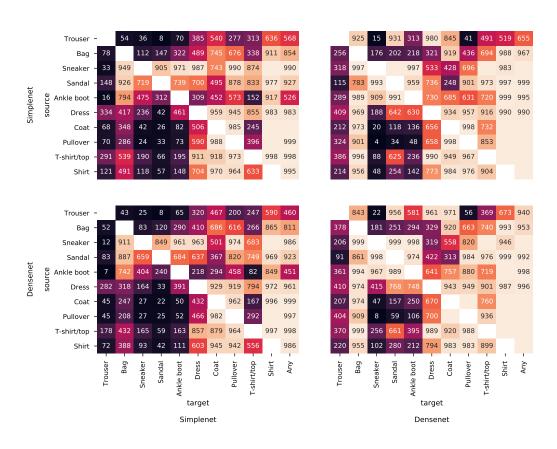


Figure 4.4: Full surrogate success rate

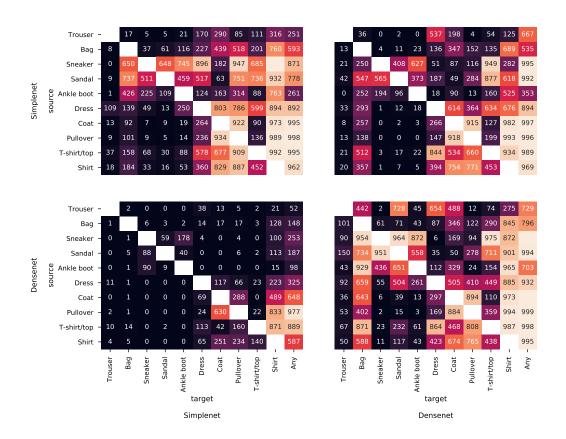


Figure 4.5: Reduced surrogate success rate

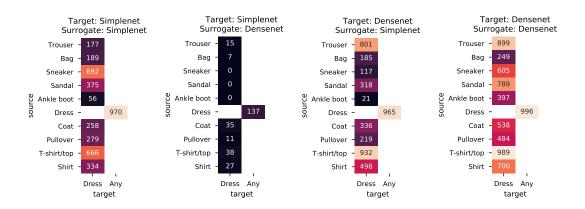


Figure 4.6: Binary surrogate success rate

Each experiment will be cross-validated using 10-fold cross-validation.

Architectures - Simple CNN - 3 cnn layers with max-pooling and one hidden dense layer - DenseNet-BC-8-52

Datasets - Fashion MNIST - Cifar-10 - bude-li čas median simple_cnn model - $\dot{\iota}$ fold_7 median dense_net model - $\dot{\iota}$ fold_0

experiments on test set - unseen data - common scenario in real world FGSM white-box, target model - confidence bound 0.5, 0.95 (for surrogate) - whole test set, targeted/non-targeted, single-image - multi-image - whole class - outcomes (metrics) ?? - median/mean/std/quantiles of step count for iterative FGSM ? - median/mean/std/quantiles of added noise? (MSE?) - rate of success (will be close to 1) - time? - some examples? which ones (unsuccessful ones if any?) show visual FGSM surrogate (different seed) - cross-validation folds cross comparison (whole/partial?) - 2 vs 20 (wiht 0.95 bound) - median model - median of accuracy - multi-image - whole class FGSM surrogate (teacher-student) - 2 median modely * 10 fold * 2 architectures - whole dataset (resplitted randomly) -1/10th of dataset (resplitted randomly) - multi-image - whole class FGSM surrogate (binary teacher-student) - 2 median-model * 10 fold * 2 architectures * 2 big small data (just one median class) - (maybe if really bad - balance dataset) EA (black-box, pure EA) - 10 random (really random) samples from test set - multiimage - vs median models hybrid - pre-trained surrogate - 2 median-surrogates * 2 binary/full * 10 random sample * ... hybrid - on-demand surrogate - 2 binary/full * 10 random sample * 2 * policies

4.2 Evolutionary Algorithms

4.3 Hybrid methods

Conclusion

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List of Tables

List of Abbreviations

TODO??

A. Evgena Framework User Documentation

TODO??