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Abstract

Scientific documents often use LATEX for typesetting. While numerous packages and templates exist, it makes sense to create a new one. Just because.

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1 TECHNOLOGY

In this chapter I'm describing the technology that I used to implement the set of utilities and experiments described in section ??? This chapter is organized as follows. In 1.1 I'm describing what is TensorFlow and how its computational graphs work. In 1.2 I'm describing Keras, an high-level API for building Machine Learning models without working with a low-level library like Tensor-Flow. In 1.3 I'm describing CleverHans, a library to generate adversarial examples against a given model. In 1.4 I'm describing scikit-learn, a library for building and training Machine Learning models.

1.1 TensorFlow

TensorFlow is a C++ framework for Machine Learning released by Google. It uses a programming model called Data Flow that aims to allow distributed and parallel computations. While this *paradigm* makes TensorFlow suitable even for research and production environments, it can be quite daunting to use when tinkering. In fact, computational graphs are built and only later executed. This is counter-intuitive at first. For example, in the following snippet of code

```
>>> import tensorflow as tf
>>> symbol = tf.constant(42)
>>> symbol
<tf.Tensor 'Const:0' shape=() dtype=int32>
```

symbol doesn't contain a reference to the integer 42. Instead it contains a reference to a node of the computational graph (which will always output 42). In general until you don't run the computational graph it's hard to determine what's going to be the value of a tensor. That's slowing down exploratory analysis.

1.1.1 ABOUT COMPUTATIONAL GRAPHS

TensorFlow computational graphs are a fundamental part of the framework. A computational graph is a directed graph representing a computation involving tensors. A tensor in TensorFlow is a multi-dimensional array: it can be a scalar, a matrix, a batch of RGB images (which is a 4D vector), ... Each node represents an operation on tensors, while edges represent tensors. (This distinction between nodes and edges is more important from a formal standpoint than a practical one: actually thinking of tensors as nodes makes no difference to the best of my knowledge.) Figure 1.1 shows a very simple computational graph. It's the one associated to an addition of two tf.float32's (i.e. two tensors made of 32-bit float numbers). The first two nodes a and b outputs

a tensor each. Those are used to feed the add node that will output the tensor resulting from the sum of a and b.

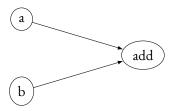


Figure 1.1: An easy computational graph

A more interesting example would be to implement a neural network using the computational graph. The neural network we want to implement consists of 4 input neurons and 4 output neurons. A graphical representation of that is given in Figure 1.2.

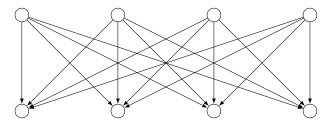


Figure 1.2: A simple neural network

We'll need the weights and the bias and do

$$X * W + b$$

where $W \in \mathbb{R}^{4 \times 4}$ and $b \in \mathbb{R}^4$, while X is a *batch* of input vectors in \mathbb{R}^4 . If you imagine the input is a flattened image of 2 by 2 pixels, this neural network learns how to classify 4 different classes of images.

Using the TensorFlow Python API to build the related computational graph, we'll do

```
>>> import tensorflow as tf
>>> W = tf.random_normal(shape=(4, 4))
>>> b = tf.random_normal(shape=(4,))
>>> X = tf.placeholder(shape=(None, 4), dtype=tf.float32)
>>> logits = tf.matmul(X, W) + b
>>> probabilities = tf.nn.softmax(logits)
>>>
>>> probabilities
<tf.Tensor 'Softmax:0' shape=(?, 4) dtype=float32>
```

Note the tf.nn.softmax function is applied to the result of the matrix multiplication. This is done for various reasons: it introduces non-linearity allowing the neural network to learn to compute

any function – what's known as the "Universal approximation theorem"; also it normalizes the results of the neural network in [0,1] allowing to interpret the output of the neural network in terms of *probabilities* or confidence levels.

To get results out of the graph you need what's called a session in TensorFlow. Basically a session *powers on* the graph, allowing you to run the computational graph with your own data and get actual tensors of actual numbers out of it. In the example above we can use a session to get probabilities out of our neural network for a batch of 2×2 images.

```
>>> import numpy as np
>>> ten_two_by_two_images_batch = np.random.rand(10, 4)
>>> ten_two_by_two_images_batch
array([[0.54485176, 0.33854871, 0.45185129, 0.79884188],
      [0.41204776, 0.23552753, 0.04101023, 0.47883844],
      [0.25544491, 0.7610509, 0.49307137, 0.6098213],
      [0.02545156, 0.70459456, 0.22067103, 0.64743811],
      [0.92359354, 0.96497353, 0.45790538, 0.49380769],
      [0.13330072, 0.22947966, 0.02996348, 0.69954114],
      [0.38397249, 0.30473362, 0.87023559, 0.90153084],
      [0.77056319, 0.94843128, 0.39095345, 0.50572861],
      [0.90112077, 0.19240995, 0.48437166, 0.46200152],
      [0.98589042, 0.2013479 , 0.86091217, 0.55886214]])
>>> with tf.Session() as session:
       # we're _feeding_ the X placeholder with an actual numpy array
        session.run(probabilities, feed_dict={X: ten_two_by_two_images_batch})
array([[0.03751625, 0.8651346, 0.05245555, 0.04489357],
      [0.11873736, 0.6301607, 0.17646323, 0.07463871],
      [0.06252376, 0.82338244, 0.07284217, 0.04125158],
      [0.12086256, 0.6911012, 0.14362463, 0.04441159],
      [0.03662292, 0.8575108, 0.04834893, 0.05751737],
      [0.12984137, 0.63064647, 0.1834405, 0.05607158],
      [0.01711418, 0.93650085, 0.02116078, 0.02522417],
      [0.04886489, 0.83023334, 0.06332602, 0.05757581],
      [0.033136, 0.84808177, 0.05061754, 0.06816475],
      [0.01250888, 0.9262628 , 0.01797607, 0.0432523 ]], dtype=float32)
                   ^^^^^^ the last image is of the second class with a confidence of 92%
```

Of course building neural networks is pretty useless if you can't train them. In fact the probabilities above are totally non-sense: they're only based on the initial weights and bias randomly extracted from a normal distribution. There's no training set, no training phase yet. To train a network in TensorFlow we need a tf.Variable. A variable in TensorFlow is a tensor whose value persists across sessions and that can be modified. This allows you to basically add parameters to your computational graphs, allowing you to compute different functions with the same graph. That, combined with a learning procedure that modifies the tf.Variables allows you to train your model.

To build a tf. Variable in TensorFlow you can just wrap an existing tensor. In the example above, the part of the graph that you want to parameterize are the weights matrix W and the bias vector b.

```
W = tf.Variable(W)
b = tf.Variable(b)
logits = tf.matmul(X, W) + b # redefine logits: using different tensors now
probabilities = tf.nn.softmax(logits)
```

To perform training we now need a training set. A training set in this example woule consist of lots of 2×2 images and an ID associated to each image representing the class of that image. *Training* basically means minimizing a loss function between the provided IDs of the training set and the IDs computed by the neural network. Minimization happens thanks to the parameters of our neural network: the w and b tf. Variables.

```
y_true = tf.placeholder(shape=(None,), dtype=tf.float32)
cross_entropy = tf.nn.softmax_cross_entropy_with_logits_v2(logits=logits, labels=y_true)
```

where *cross-entropy* is our loss function that we want to minimize.

To minimize the cross-entropy one can use the Gradient descent algorithm – an optimization procedure that works iteratively. TensorFlow implements that algorithm via a class exposing .minimize(): it returns a tensor that each time you run it in a session will modify the graph's tf. Variables, trying to lower the value of the loss function.

```
from tf.train import GradientDescentOptimizer as SGD
optimizer = SGD(learning_rate=0.5).minimize(cross_entropy)
```

So a typical training phase would consist of

```
with tf.Session() as session:
    session.run(tf.global_variables_initializer()) # needed.

n_training_steps = 10
for i in range(n_training_steps):
    images, classes = next_batch()
    # fictitious function returning batches from the training set

with tf.Session() as session:
    session.run(optimizer, feed_dict={X: images, y_true: classes})
```

As you increase n_training_steps, wand b will be modified pushing the value of the loss function towards 0. That makes the model learn how to correctly classify images similar to those in the training set.

1.2 Keras

Keras is a high level library for building Machine Learning models. It consists of a simple API and bindings for a backend of choice, most notably TensorFlow. When you use the library, you're encouraged to use *layers* instead of plain tensors. In fact the whole concept of TensorFlow tensors is hidden away by the library abstractions. To implement your model you stack layers – basically you build a list of layers. For example, to implement a simple neural network one would stack a Dense layer of 4 hidden neurons and an Activation layer for the Softmax function.

```
from keras.models import Sequential
model = Sequential([
   Dense(batch_input_shape=(None, 4), units=4),
   Activation('softmax')
])
```

Under the hood, a Layer is simply a callable object whose __call()_ method takes a TensorFlow tensor X and manipulates it. For example a simple custom one would be

```
from keras.engine.base_layer import Layer
import tensorflow as tf

class Add42(Layer):
    def __init__(self):
        self.fortytwo = tf.constant(42)

def __call__(self, X):
    return tf.add(X, self.fortytwo)
```

(Of course in a real layer instead of self.fortytwo one would have trainable TensorFlow variables.)

While the idea is beginner friendly – as it allows people to immediately concentrate on model's architecture instead of thinking about tensors – at the time of this writing the abstraction can become leaky when things goes wrong, needing the programmer to know about TensorFlow computational graphs to debug her program successfully.

1.3 CLEVERHANS

CleverHans is a library written by Google for building adversarial examples. It implements a variety of attacks against neural networks (FGS, T-FGS, Carlini and Wagner, ...) and it's compatible with models built with Keras or plain TensorFlow. It uses the same computational graphs of TensorFlow to generate adversarial examples; again, as you use the library more and more understanding computational graphs becomes a necessity.

1.4 Scikit-learn

Scikit-learn is a Python library written by Google providing a number of models, learning algorithms and other utilities for Machine Learning. It's perfect for fast prototyping as it uses a simple and consistent API and handles numpy arrays or even Python lists.

I've used scikit-learn to borrow a couple of decomposition algorithms (PCA, FastICA, ...) without having to implement them. This required a bit of reasoning as reusing scikit-learn algorithms on TensorFlow graphs was not straightforward.

1.5 GOOGLE CLOUD PLATFORM

Google Cloud Platform is a set of services that allows people to *borrow* computational power from Google. In our case, we used Google Cloud Platform to get access to a GPU for few dollars. That made running experiments a lot faster in many cases.

2 Training Models

Before running experiments on how ML models can be *shielded* against adversarial examples we had to create and train them.

2.1 MOTIVATION

As we're training a variety of models it's useful to find a strategy to find an appropriate number of epochs to train the model for. Fixing a number of epochs (say 500) and train all the models for that same number of epochs feels *cheesy*: some models can learn faster, other slower, and you can end up comparing models that are at different stage in their learning phase.

What we did was trying to *understand* when the model has done learning and stop training there. We used two approaches/heuristics that are described in the following sections.

Note that trying to make the model learn better we reduced the learning rate on plateau: if for a fixed number of epochs the accuracy on the validation set does no longer improve, the learning rate is reduced. This is known to help models to move on when the gradient descent algorithm is stuck somewhere.

To make our experiments here we used a fully-connected neural network with 2 layers of 100 hidden neurons each and 10 output neurons – *FC-100-100-10*. This structure is the same described in ???, which is the paper we got inspiration from. ??? trained FC-100-100-10 for 500 epochs and we assume they chose that number as it gave them the best results on the test set.

What we want to do is to empirically prove that we can achieve the same accuracy on the test set without training FC-100-100-10 for exactly 500 epochs but instead stopping training when the model stops learning according to at least one of our heuristics.

2.2 EARLYSTOPPING

EarlyStopping is one of the two heuristics. It's already implemented by Keras as a Callback. It checks if over a specified number of epochs (the *patience*) a specified metrics stopped improving. When that happens the model is asked to stop training there. We set a *patience* of 60 epochs as on Google Cloud Platform 1 epochs runs is 1 second; a patience of 60 epochs means that the model didn't improve for a whole minute.

2.3 STOPONSTABLEWEIGHTS

StopOnStableWeights is the second heuristic. It has to be implemented from scratch. This heuristic collects data on model's weights over a specified number of epochs called the *patience*. Then, the same weight over different epochs is checked: if it has a small standard deviation maybe it's getting

stable. If the maximum of these standard deviation is below a certain threshold we conclude the model's weights are getting stable, hence the model is stopping learning.

2.4 Data obtained

We obtained the following data:

- Model trained for 1000 epochs: accuracy on the test set of 97.71%
- Model trained for 500 epochs: accuracy on the test set of 97.55%
- Model trained for 500 epochs, reducing learning rate on plateau after 30 epochs: accuracy
 on the test set of 97.47%
- Model trained for -1 epochs, stopping after accuracy on the validation set stopped improving for 60 epochs: accuracy on the test set of 97.44% (trained for 424 epochs)
- Model trained for -1 epochs, stopping as model weights had a maximum standard deviation
 of 0.5 over 60 epochs: accuracy on the test set of 97.66% (trained for 614 epochs)
- Model trained for -1 epochs, stopping after accuracy on the validation set stopped improving for 60 epochs, reducing learning rate on plateau after 30 epochs: accuracy on the test set of 97.18% (trained for 510 epochs)
- Model trained for -1 epochs, stopping as model weights had a maximum standard deviation
 of 0.5 over 60 epochs, reducing learning rate on plateau after 30 epochs: accuracy on the
 test set of 97.6% (trained for 514 epochs)

2.5 Conclusion

From the above results the best approach seems to be using the *StopOnStableWeights* heuristics with a patience of 60 epochs, reducing the learning rate on plateau after 30 epochs. In the following chapters we'll use this setting for training models, even when they're no longer the FC-100-100-10 used here.

3 Reconstruction or retraining?

As we're testing different filter techniques it's important that we understand how we're implementing these techniques in our models as defenses.

3.1 Motivation

There are at least two different fashions to integrate a filter/decomposition technique in our models. We can either add a layer to a pre-trained model and use it only *after* training, or we can add a layer to a pre-trained model and re-train the network after this layer has been added. In the latter case, the model has been trained only on filtered input, while in the former it does not. Following lexicon used in ???, we named the first technique as *reconstruction* and the second one *retraining*. There's another alternative which is to initialize the model, add the filter layer then train the network while it's still freshly initialized. It basically consists in making the network never to see unfiltered input. This seemed to make things so much better for the attacker than both reconstruction and retraining that we just avoided to compare it with the other approaches. Instead when it comes to reconstruction and retraining who is better is not that clear. We did some measurements to better understand it.

3.2 How did we perform the comparison

We built and train models using both reconstruction and retraining. We had now two families of models. For example, for FC-100-100-10 with a filter layer implementing PCA retaining 80 components we had one model trained using reconstruction and another trained using retraining. We attacked each model using Fast Gradient Sign obtaining a curve saying what was the adversarial success score as η increases – where η is the freedom given to the attacker: the more the freedom the easier is to forge an input. Then, we paired each model trained using reconstruction with the same model trained using retraining. We computed the average value of the curve obtained subtracting the curve for retraining from the curve for reconstruction. At last we computed the average for the set of pairs, getting what is the average difference between reconstruction and retraining.

3.3 Data obtained

We obtained that retraining is only 1% more effective than reconstruction when it comes to protect models from adversarial input.

3.4 Conclusion

While implementing retraining defense is not hard we concluded the difference is not significant enough and decided to stick with reconstruction which is a clearer solution – it allows us to think about filter techniques without worrying about how that impacts on training.

ACRONYMS

PCA Principal component analysis

SNF Smith normal form TDA Topological data analysis

GLOSSARY