Machine Learning in Cyber Security

Tutorial - Nov 9th, 2020

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Agenda

- Setting up a Python Scientific Environment
- Jupyter Notebook
- Scientific Computing (using <u>numpy</u>)
- Data Analysis (using <u>pandas</u>)
- Data Visualization (using <u>matplotlib</u>)

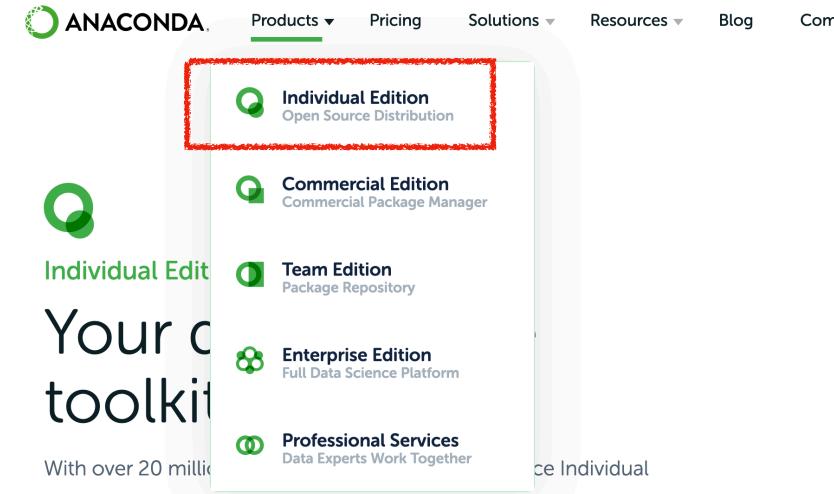
1. Setting up a Python Scientific Environment

- Download Anaconda
- Install:

```
$ chmod u+x /path/to/Anaconda<...>.sh
$ /path/to/Anaconda<...>.sh
```

- This should already contain most of dependencies required for the course
- If not:

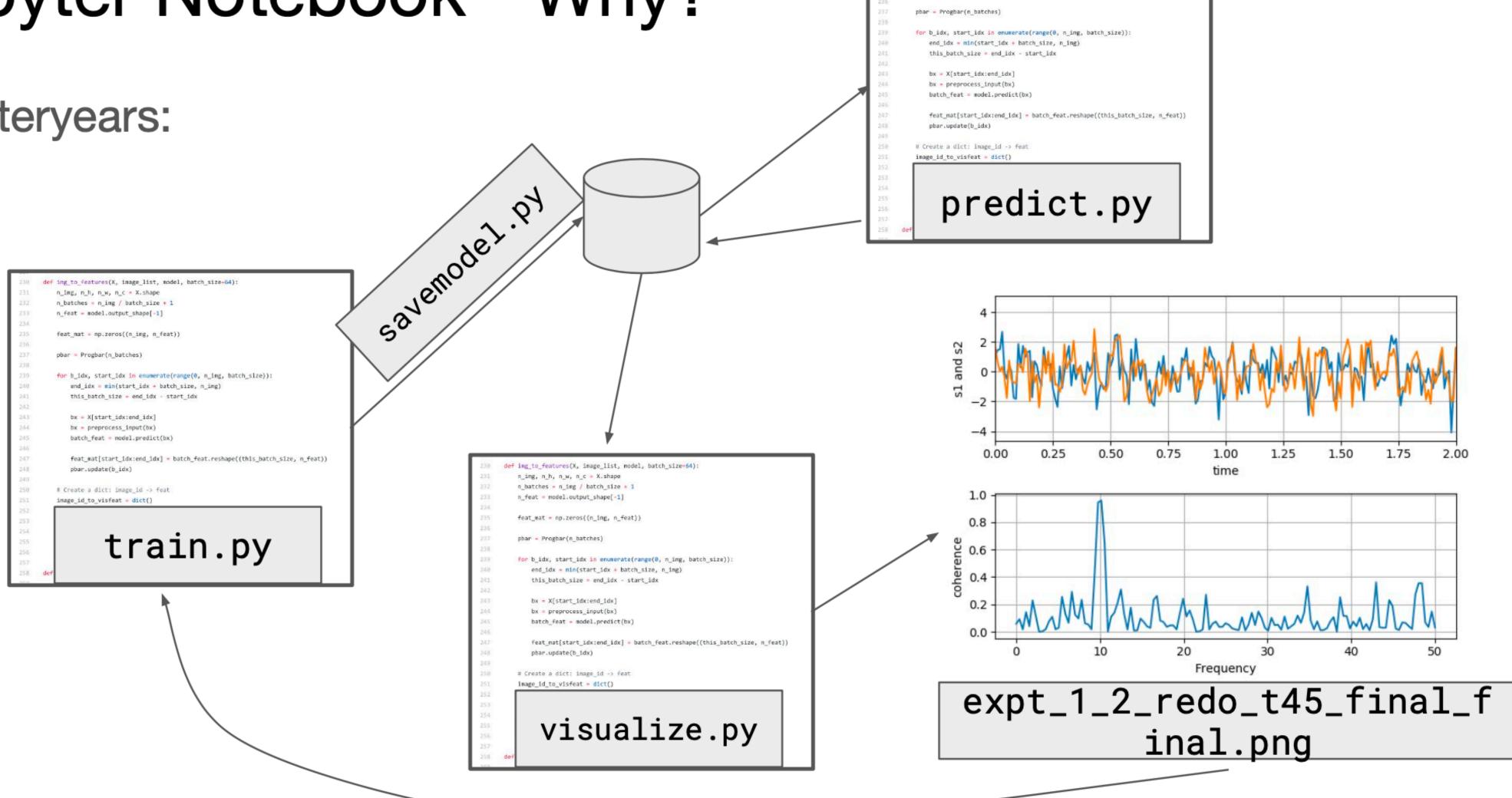
```
$ conda install <package-name> # or
$ pip install <package-name>
```



Edition (Distribution) is the easiest way to perform Python/R data science and machine learning on a single machine. Developed for solo practitioners, it is the toolkit that equips you to work with thousands of open-source packages and libraries.

2. Jupyter Notebook - Why?

In the yesteryears:



def ing_to_features(X, image_list, model, batch_size=64):

n_img, n_h, n_w, n_c = X.shape n_batches = n_img / batch_size = 1

n_feat = model.output_shape[-1]

feat_mat = np.zeros((n_img, n_feat))

2. Jupyter Notebook - Why?

Now:

42.ipynb

```
In [36]: n_epochs = 100
         batch_size = 150
         with tf.Session() as sess:
             init.run()
             for epoch in range(n_epochs):
                 for iteration in range(mnist.train.num_examples // batch_size):
                     X_batch, y_batch = mnist.train.next_batch(batch_size)
                     X_batch = X_batch.reshape((-1, n_steps, n_inputs))
                     sess.run(training_op, feed_dict={X: X_batch, y: y_batch})
                 acc_train = accuracy.eval(feed_dict={X: X_batch, y: y_batch})
                 acc_test = accuracy.eval(feed_dict={X: X_test, y: y_test})
                 print(epoch, "Train accuracy:", acc_train, "Test accuracy:", acc_test)
         0 Train accuracy: 0.93333334 Test accuracy: 0.9311
         1 Train accuracy: 0.96666664 Test accuracy: 0.9522
         2 Train accuracy: 0.97333336 Test accuracy: 0.9586
         3 Train accuracy: 0.96666664 Test accuracy: 0.9607
         4 Train accuracy: 0.97333336 Test accuracy: 0.9673
         5 Train accuracy: 0.98 Test accuracy: 0.9669
         6 Train accuracy: 0.97333336 Test accuracy: 0.9693
         7 Train accuracy: 0.96666664 Test accuracy: 0.968
         8 Train accuracy: 0.9533333 Test accuracy: 0.9723
         9 Train accuracy: 0.97333336 Test accuracy: 0.9683
         10 Train accuracy: 0.9866667 Test accuracy: 0.9734
         11 Train accuracy: 0.96666664 Test accuracy: 0.969
         12 Train accuracy: 0.9866667 Test accuracy: 0.9726
         13 Train accuracy: 0.9866667 Test accuracy: 0.9774
         14 Train accuracy: 0.98 Test accuracy: 0.9705
         15 Train accuracy: 0.9866667 Test accuracy: 0.976
         16 Train accuracy: 0.9866667 Test accuracy: 0.9739
         17 Train accuracy: 0.9866667 Test accuracy: 0.9709
         18 Train accuracy: 0.98 Test accuracy: 0.9736
         19 Train accuracy: 0.9866667 Test accuracy: 0.9775
```

Train models

```
In [14]: def plot_predictions(clf, axes):
             x0s = np.linspace(axes[0], axes[1], 100)
             x1s = np.linspace(axes[2], axes[3], 100)
             x0, x1 = np.meshgrid(x0s, x1s)
             X = np.c_{x0.ravel(), x1.ravel()}
             y_pred = clf.predict(X).reshape(x0.shape)
             y_decision = clf.decision_function(X).reshape(x0.shape)
             plt.contourf(x0, x1, y pred, cmap=plt.cm.brg, alpha=0.2)
             plt.contourf(x0, x1, y decision, cmap=plt.cm.brg, alpha=0.1)
         plot_predictions(polynomial_svm_clf, [-1.5, 2.5, -1, 1.5])
         plot_dataset(X, y, [-1.5, 2.5, -1, 1.5])
         save_fig("moons_polynomial_svc_plot")
         plt.show()
         Saving figure moons polynomial svc plot
             1.5
             1.0
          X_2^{0.5}
             0.0
            -0.5
               -1.5 -1.0 -0.5 0.0
                                        1.0 1.5 2.0 2.5
                                    x_1
```

Predict + Visualize

Computing features matches

- Matching
- d_i ∈ D_i → d_i ∈ D_i
- Matching is a nearest neighbor problem
- In a low dimensional space, can be efficiently performed using KD-Trees
- However, SIFT descriptors are 128-d vectors
- Alternatives
- Approximate Nearest Neighbors (as FLANN library)
- Brute force matching
- · Dimensionality reduction

Consider the two sets of descriptors, \mathbf{D}_i and \mathbf{D}_j , computed using a method as SIFT or SURF, for two images I_i and I_j . Matching can be performed using nearest neighbors, just selecting for each $\mathbf{d}_i \in \mathbf{D}_i$ the closest vector $\mathbf{d}_j \in \mathbf{D}_j$. Nearest neighbor queries can be efficiently done representing \mathbf{D}_i in a KD-Tree.

KD-Tree performance is close to brute force for vectors presenting large dimensions. SIFT vectors are 128-d and SURF ones are 64-d. Before matching using KD-Trees, a dimensionality reduction procedure, as PCA, is recommended. Sklearn and OpenCV provide PCA implementations.

```
In [6]: from sklearn.decomposition import PCA
    pca = PCA(n_components=10)

pca.fit(D_i)
    D_i = pca.transform(D_i)
    D_j = pca.transform(D_j)
```

- · Lowe recomends to compare the two nearest neighbors
- In a good match, there is a constrast between the two distances
- Descriptors with no proper match present similar distances between their closest neighbors
- Filtering
- d_i ∈ D_i should be assigned to just one d_i ∈ D_i

Interleave code, plots and explanations (markdown/tex)

Hands-on in Jupyter Notebook

- Create a new environment for the course
- conda create -n mlcysec python=3.7 numpy matplotlib jupyter nb_conda
- conda activate mlcysec
- Start Jupyter Notebook
- jupyter notebook
- Open browser
- http://localhost:8888

```
To access the notebook, open this file in a browser:
    file:///Users/wanghuipo/Library/Jupyter/runtime/nbserver-74986-open.html
Or copy and paste one of these URLs:
    http://localhost:8888/?token=d10b546ae3992c2010e05de7e25020431ea8027974373c30
```

or http://127.0.0.1:8888/?token=d10b546ae3992c2010e05de7e25020431ea8027974373c30