```
In [1]: # This Python 3 environment comes with many helpful analytics libraries
         installed
        # It is defined by the kaggle/python docker image: https://github.com/ka
        ggle/docker-python
        # For example, here's several helpful packages to load in
        import numpy as np # linear algebra
        import json
        from matplotlib import pyplot as plt
        from skimage import color
        from skimage.feature import hog
        from sklearn import svm
        from sklearn.metrics import classification report, accuracy score
        from sklearn.datasets import load_iris
        from sklearn.linear model import LogisticRegression
        import os.path
        import random
        from random import choice, sample
        import PIL
```

```
In [2]: import itertools
import numpy as np
import matplotlib.pyplot as plt

from sklearn import svm, datasets
from sklearn.model_selection import train_test_split
from sklearn.metrics import confusion_matrix
```

```
In [3]: from sklearn.datasets import load_iris
from sklearn.model_selection import cross_val_score
from sklearn.tree import DecisionTreeClassifier
```

In [4]: # Install a conda package in the current Jupyter kernel
import sys
!conda install --yes --prefix {sys.prefix} opency

Solving environment: done

Package Plan

environment location: /home/ec2-user/anaconda3/envs/python2

added / updated specs:

- opency

The following packages will be downloaded:

package	build		
libvpx-1.7.0	h439df22_0	2.4	MB
libopus-1.3	h7b6447c_0	631	KB
opencv-3.4.2	py27h6fd60c2_1	11	KB
jasper-2.0.14	h07fcdf6_1	1.1	MB
libglu-9.0.0	hf484d3e_1	377	KB
certifi-2018.11.29	py27_0	146	KB
ffmpeg-4.0	hcdf2ecd_0	73.7	MB
py-opencv-3.4.2	py27hb342d67_1	1.2	MB
freeglut-3.0.0	hf484d3e_5	251	KB
libopencv-3.4.2	hb342d67_1	40.4	MB
	Total:	120.2	MB

The following NEW packages will be INSTALLED:

```
ffmpeg: 4.0-hcdf2ecd_0
freeglut: 3.0.0-hf484d3e_5
jasper: 2.0.14-h07fcdf6_1
libglu: 9.0.0-hf484d3e_1
libopencv: 3.4.2-hb342d67_1
libopus: 1.3-h7b6447c_0
libvpx: 1.7.0-h439df22_0
py-opencv: 3.4.2-py27hb342d67_1
```

The following packages will be UPDATED:

```
certifi: 2018.8.24-py27_1 conda-forge --> 2018.11.29-py
27_0
    libstdcxx-ng: 7.2.0-hdf63c60_3 --> 8.2.0-hdf63c6
0_1
    opencv: 3.4.1-py27h6fd60c2_1 --> 3.4.2-py27h6fd60c2_1
    openssl: 1.0.2p-h470a237_0 conda-forge --> 1.0.2p-h14c39
75_0
```

The following packages will be DOWNGRADED:

ca-certificates: 2018.8.24-ha4d7672_0 conda-forge --> 2018.03.07-0

```
# | 100%
libopus-1.3
           631 KB
                 # | 100%
opency-3.4.2
           | 11 KB
                 # | 100%
jasper-2.0.14
           1.1 MB
                 # | 100%
libglu-9.0.0
           377 KB
                  # | 100%
                 certifi-2018.11.29
          146 KB
# | 100%
ffmpeg-4.0
           | 73.7 MB
                 # | 100%
                 py-opency-3.4.2
           1.2 MB
# | 100%
freeglut-3.0.0
                 251 KB
# | 100%
libopencv-3.4.2
           40.4 MB
                 # | 100%
Preparing transaction: done
Verifying transaction: done
Executing transaction: done
```

In [5]: import cv2

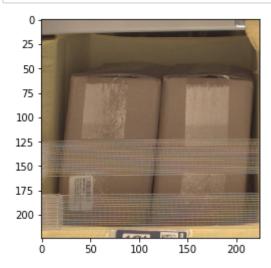
In [6]: # Input data files are available in the "../input/" directory.
For example, running this (by clicking run or pressing Shift+Enter) wi
11 list the files in the input directory
from subprocess import check_output
Any results you write to the current directory are saved as output.

```
N=range(500000)
 In [9]:
         num samples=5000
         dataset_index=random.sample(N,num_samples)
         #dataset index
         meta_dir = "amazon-bin/metadata/"
In [10]:
         img_dir = "amazon-bin/bin-images/"
In [11]:
         fname_img_vec=[]
          fname_meta_vec=[]
         for i in range(num samples):
             fname_img=str(dataset_index[i])+'.jpg'
             fname_meta=str(dataset_index[i])+'.json'
             jpg path = os.path.join(img dir,fname img)
             json_path = os.path.join(meta_dir,fname_meta)
             if os.path.isfile(jpg path) & os.path.isfile(json path):
                  fname img vec.append(fname img)
                  fname_meta_vec.append(fname_meta)
         print(np.shape(fname_img_vec))
         print(np.shape(fname_meta_vec))
          #fname meta vec
         (4911,)
         (4911,)
```

```
In [12]: # get label from
         num samples=len(fname img vec)
         #labels=np.zeros((num samples,1))
         labels=[]
         fname img vec less 5=[]
         fname_meta_vec_less_5=[]
         for i in range(num_samples):
             fname meta=fname meta vec[i]
             fname_img=fname_img_vec[i]
             json path = os.path.join(meta_dir,fname_meta)
             f = open(json path)
             dataset = json.load(f)
             expected_quantity=dataset['EXPECTED_QUANTITY']
             if expected quantity<=5:</pre>
                  #print(fname)
                  #print(expected quantity)
                  f.close()
                  #labels[i,0]=expected quantity
                  labels.append(expected quantity)
                  fname img vec less 5.append(fname img)
                  fname meta vec less 5.append(fname meta)
         labels=np.array(labels,ndmin=2)
         labels=labels.T
         print(np.shape(labels))
         print(np.shape(fname img vec less 5))
         print(np.shape(fname_meta_vec_less_5))
         (3296, 1)
```

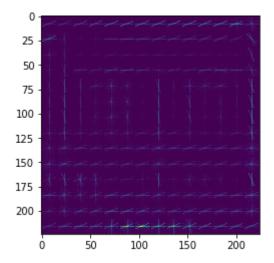
(3296, 1) (3296,) (3296,)

```
In [13]:
         fname=fname_img_vec_less_5[i]
         fname='542.jpg'
         fname
         jpg_path = os.path.join(img_dir,fname)
         # Convert Image to array
         img = PIL.Image.open(jpg_path)
         resized img = img.resize((224,224))
         plt.imshow(resized_img)
         arr = np.array(resized_img)
         plt.imshow(arr)
         data_gray = color.rgb2gray(arr)
         #plt.imshow(data gray)
         ppc = 16
         hog_images = []
         hog_features = []
         fd,hog_image = hog(data_gray, orientations=8, pixels_per_cell=(ppc,ppc),
         cells_per_block=(4, 4),block_norm= 'L2',visualise=True)
```



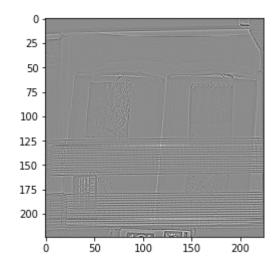
In [14]: plt.imshow(hog_image)

Out[14]: <matplotlib.image.AxesImage at 0x7fd90a88d510>



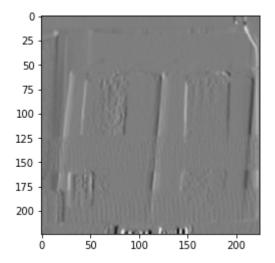
```
In [15]: fname_img_vec_less_5[i]
Out[15]: '453433.jpg'
In [16]: #img = cv2.imread(jpg_path,0)
    #resized_img = img.resize((224,224))
    laplacian = cv2.Laplacian(data_gray,cv2.CV_64F)
    sobelx = cv2.Sobel(data_gray,cv2.CV_64F,1,0,ksize=5)
    sobely = cv2.Sobel(data_gray,cv2.CV_64F,0,1,ksize=5)
    np.shape(img)
Out[16]: (591, 577, 3)
In [17]: plt.imshow(laplacian,cmap = 'gray')
```

Out[17]: <matplotlib.image.AxesImage at 0x7fd90a816350>



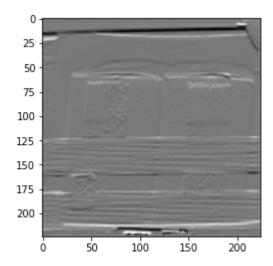
```
In [18]: plt.imshow(sobelx,cmap = 'gray')
```

Out[18]: <matplotlib.image.AxesImage at 0x7fd908128a10>



```
In [19]: plt.imshow(sobely,cmap = 'gray')
```

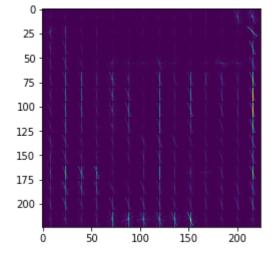
Out[19]: <matplotlib.image.AxesImage at 0x7fd9080a5110>



```
In [20]: ppc = 16
    hog_images = []
    hog_features = []
    fd,hog_image = hog(sobelx, orientations=8, pixels_per_cell=(ppc,ppc),cel
    ls_per_block=(4, 4),block_norm= 'L2',visualise=True)
```

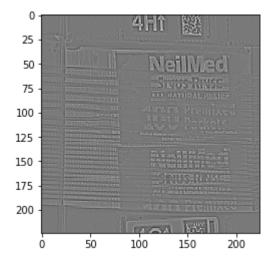
In [21]: plt.imshow(hog_image)

Out[21]: <matplotlib.image.AxesImage at 0x7fd90a90a1d0>



```
# do HOG on edges images, resize images on the fly
In [22]:
         hog features size=15488
         num samples=len(fname meta vec less_5)
         hog features matrix=np.zeros((num samples,hog features size))
         for i in range(num samples):
             fname=fname img vec less 5[i]
             fname
             jpg path = os.path.join(img dir,fname)
             # Convert Image to array
             img = PIL.Image.open(jpg_path)
             resized img = img.resize((224,224))
             # img = Image.open(jpg path).convert('RGB')
             # resized img = img.resize((224,224),img.BILINEAR)
             arr = np.array(resized img)
             #plt.imshow(arr)
             data_gray = color.rgb2gray(arr)
             #plt.imshow(data gray)
             ppc = 16
             hog images = []
             hog features = []
             laplacian = cv2.Laplacian(data gray,cv2.CV 64F)
             fd, hog image = hog(laplacian, orientations=8, pixels per cell=(ppc,p
         pc),cells_per_block=(4, 4),block_norm= 'L2',visualise=True)
             hog images.append(hog image)
             hog features.append(fd)
             hog_features = np.array(hog_features)
             hog features matrix[i,:]=hog features
             plt.imshow(laplacian,cmap = 'gray')
         np.shape(hog_features_matrix)
```

Out[22]: (3296, 15488)



```
In [23]:
         # do HOG on orginal images, resize images on the fly
         hog features size=15488
         num samples=len(fname meta vec less 5)
         hog features matrix=np.zeros((num samples,hog features size))
         for i in range(num samples):
             fname=fname img vec less 5[i]
             fname
             jpg path = os.path.join(img dir,fname)
             # Convert Image to array
             img = PIL.Image.open(jpg path)
             resized img = img.resize((224,224))
             # img = Image.open(jpg path).convert('RGB')
             # resized img = img.resize((224,224),img.BILINEAR)
             arr = np.array(resized img)
             #plt.imshow(arr)
             data gray = color.rgb2gray(arr)
             #plt.imshow(data gray)
             ppc = 16
             hog images = []
             hog features = []
             fd, hog image = hog(data gray, orientations=8, pixels per cell=(ppc,p
         pc),cells per block=(4, 4),block norm= 'L2',visualise=True)
             hog images.append(hog image)
             hog features.append(fd)
             hog features = np.array(hog features)
             hog features matrix[i,:]=hog features
         np.shape(hog features matrix)
```

Out[23]: "\n# do HOG on orginal images, resize images on the fly\nhog features s ize=15488\nnum_samples=len(fname_meta_vec_less_5)\nhog_features_matrix= np.zeros((num samples,hog features size))\nfor i in range(num sample fname=fname_img_vec_less_5[i]\n $s): \n$ fname\n jpg path = os.pa th.join(img_dir,fname)\n # Convert Image to array\n img = PIL.Ima ge.open(jpg path)\n resized img = img.resize((224,224))\n # img =# resized img = img.resize((22 Image.open(jpg path).convert('RGB')\n 4,224),img.BILINEAR)\n arr = np.array(resized_img)\n #plt.imshow (arr)\n data_gray = color.rgb2gray(arr)\n #plt.imshow(data gray) ppc = 16 n $hog_images = []\n$ hog_features = []\n image = hog(data gray, orientations=8, pixels per_cell=(ppc,ppc),cells_ per block=(4, 4),block norm= 'L2',visualise=True)\n hog images.appen d(hog image)\n hog features.append(fd)\n hog features = np.array (hog_features)\n hog_features_matrix[i,:]=hog_features\nnp.shape(hog _features_matrix)\n"

```
In [24]: data_frame = np.hstack((hog_features_matrix,labels))
    np.shape(data_frame)
    np.random.shuffle(data_frame)
```

```
In [25]: percentage = 80
   partition = int(num_samples*percentage/100)
```

```
In [26]: x train, x test = data frame[:partition,:-1], data frame[partition:,:-1
         y_train, y_test = data_frame[:partition,-1:].ravel() , data_frame[partit
         ion:,-1:].ravel()
         #clf = svm.SVC(kernel='poly',C=0.1)
         #clf.fit(x train,y train)
In [27]: # Logistic Regression
         #clf = LogisticRegression(random state=0, solver='lbfgs', multi class='mu
         ltinomial')
         #clf.fit(x train,y train)
In [28]: # Decision Tree
         #clf = DecisionTreeClassifier(random state=0)
         #clf.fit(x train,y train)
In [29]: | # svm tuning
         from sklearn import svm, grid search
         from sklearn.model selection import GridSearchCV
         def svc param_selection(X, y, nfolds):
             Cs = [0.001, 0.01, 0.1, 1, 10]
             gammas = [0.001, 0.01, 0.1, 1]
             param_grid = {'C': Cs, 'gamma' : gammas}
```

/home/ec2-user/anaconda3/envs/python2/lib/python2.7/site-packages/sklea rn/cross_validation.py:41: DeprecationWarning: This module was deprecat ed in version 0.18 in favor of the model_selection module into which al 1 the refactored classes and functions are moved. Also note that the in terface of the new CV iterators are different from that of this module. This module will be removed in 0.20.

grid search = GridSearchCV(svm.SVC(kernel='poly'), param grid, cv=nf

"This module will be removed in 0.20.", DeprecationWarning) /home/ec2-user/anaconda3/envs/python2/lib/python2.7/site-packages/sklea rn/grid_search.py:42: DeprecationWarning: This module was deprecated in version 0.18 in favor of the model_selection module into which all the refactored classes and functions are moved. This module will be removed in 0.20.

DeprecationWarning)

grid_search.fit(X, y)
grid search.best params

return grid search.best params

print('checkpoint')

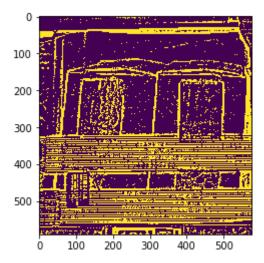
olds)

```
In [ ]: # SVM
        clf = svm.SVC(kernel='poly',C=C)
        clf.fit(x_train,y_train)
In [ ]: y pred = clf.predict(x_test)
        y pred
In [ ]: np.histogram(y pred)
In [ ]: print("Accuracy: "+str(accuracy_score(y_test, y_pred)))
        print('\n')
        print(classification_report(y_test, y_pred))
In [ ]:
In [ ]:
        gamma
In [ ]: def plot confusion_matrix(cm, classes,
                                   normalize=False,
                                   title='Confusion matrix',
                                   cmap=plt.cm.Blues):
            This function prints and plots the confusion matrix.
            Normalization can be applied by setting `normalize=True`.
            if normalize:
                cm = cm.astype('float') / cm.sum(axis=1)[:, np.newaxis]
                print("Normalized confusion matrix")
            else:
                print('Confusion matrix, without normalization')
            print(cm)
            plt.imshow(cm, interpolation='nearest', cmap=cmap)
            plt.title(title)
            plt.colorbar()
            tick_marks = np.arange(len(classes))
            plt.xticks(tick_marks, classes, rotation=45)
            plt.yticks(tick_marks, classes)
            fmt = '.2f' if normalize else 'd'
            thresh = cm.max() / 2.
            for i, j in itertools.product(range(cm.shape[0]), range(cm.shape[1
        ])):
                plt.text(j, i, format(cm[i, j], fmt),
                          horizontalalignment="center",
                          color="white" if cm[i, j] > thresh else "black")
            plt.ylabel('True label')
            plt.xlabel('Predicted label')
            plt.tight_layout()
```

```
In [ ]: cnf_matrix = confusion_matrix(y_test, y_pred)
         np.set printoptions(precision=2)
         cnf matrix
In [ ]:
         class_names=[0,1,2,3,4,5]
 In [ ]:
 In [ ]: # Plot non-normalized confusion matrix
         plt.figure()
         plot confusion matrix(cnf matrix, classes=class names,
                               title='Confusion matrix, without normalization')
In [ ]: # Plot normalized confusion matrix
         plt.figure()
         plot confusion matrix(cnf matrix, classes=class names, normalize=True,
                               title='Normalized confusion matrix')
In [ ]:
         1+1
         print('checkpoint')
In [ ]:
         # Tape removal algorithm
In [ ]:
In [14]:
         import cv2 as cv
In [22]: def show wait destroy(winname, img):
             cv.imshow(winname, img)
             cv.moveWindow(winname, 500, 0)
             cv.waitKey(0)
             cv.destroyWindow(winname)
In [15]: fname='542.jpg'
         fname
         jpg_path = os.path.join(img_dir,fname)
In [17]: src = cv.imread(jpg path, cv.IMREAD COLOR)
         gray = cv.cvtColor(src, cv.COLOR_BGR2GRAY)
In [20]: # Apply adaptiveThreshold at the bitwise not of gray, notice the ~ symbo
         gray = cv.bitwise not(gray)
         bw = cv.adaptiveThreshold(gray, 255, cv.ADAPTIVE THRESH MEAN C, cv.THRES
         H_BINARY, 15, -2)
```

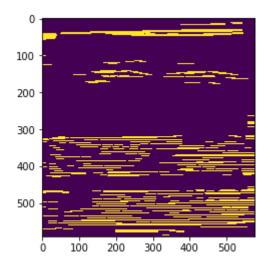
```
In [27]: plt.imshow(bw)
```

Out[27]: <matplotlib.image.AxesImage at 0x7fb5baf62710>



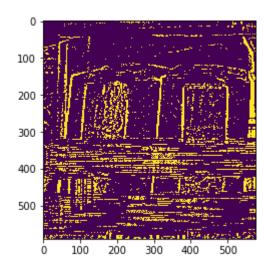
```
In [48]: plt.imshow(horizontal)
```

Out[48]: <matplotlib.image.AxesImage at 0x7fb5b8c12250>



```
In [57]: diff=bw-horizontal
   plt.imshow(diff)
```

Out[57]: <matplotlib.image.AxesImage at 0x7fb5b8569e10>



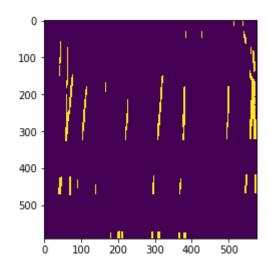
```
In [49]: # Specify size on vertical axis
    rows = vertical.shape[0]
    verticalsize = rows / 30

# Create structure element for extracting vertical lines through morphol
    ogy operations
    verticalStructure = cv.getStructuringElement(cv.MORPH_RECT, (1, vertical
        size))

# Apply morphology operations
    vertical = cv.erode(vertical, verticalStructure)
    vertical = cv.dilate(vertical, verticalStructure)
```

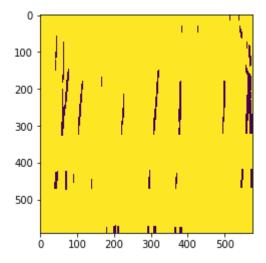
```
In [50]: plt.imshow(vertical)
```

Out[50]: <matplotlib.image.AxesImage at 0x7fb5b8b73e90>

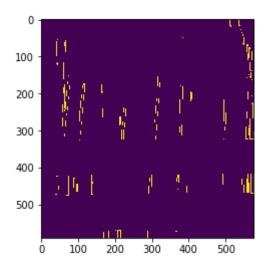


```
In [51]: vertical = cv.bitwise_not(vertical)
    plt.imshow(vertical)
```

Out[51]: <matplotlib.image.AxesImage at 0x7fb5b872dc90>

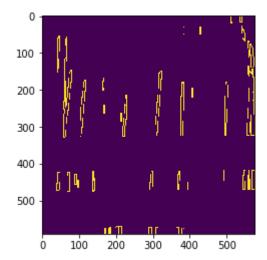


Out[52]: <matplotlib.image.AxesImage at 0x7fb5b871db10>



```
In [53]: # Step 2
kernel = np.ones((2, 2), np.uint8)
edges = cv.dilate(edges, kernel)
plt.imshow(edges)
```

Out[53]: <matplotlib.image.AxesImage at 0x7fb5b868c850>



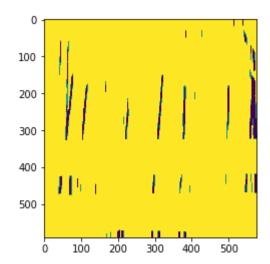
```
In [54]: # Step 3
    smooth = np.copy(vertical)

# Step 4
    smooth = cv.blur(smooth, (2, 2))

# Step 5
    (rows, cols) = np.where(edges != 0)
    vertical[rows, cols] = smooth[rows, cols]
```

```
In [55]: plt.imshow(vertical)
```

Out[55]: <matplotlib.image.AxesImage at 0x7fb5b86030d0>



In []: