Week 5 Progress Report

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Table of Contents

- 1. Learning and Using Principal Component Analysis (PCA)
- 2. Further Exploring the Random Forest Method
- 3. Trying more Neural Network Models
- 4. Future Steps and Goals

Principal Component Analysis (PCA)

- 1. Quick Understanding and Purpose
- 2. Visualizations and Metrics Received
- 3. How to further use data?

Initial Research and Understanding of PCA

- For high dimensional data, PCA is a method used to reduce the number of variables in data by extracting the important features from a large pool.
- PCA combines variables that are highly correlated together and form groups called 'principal components' that accounts for most variance in data
- Helps avoid overfitting by focusing on principal components instead of learning from non-important features
 - 'Denoising'
- In our dataset, find features (areas or pixels) that are most important in determining if it the melt pool will result in a good or bad part

Using PCA on Images

Principal Component Analysis (PCA) Application to images

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Can we use PCA for images?



- It took a while to realize (Turk, Pentland, 1991), but yes.
- Let us consider a 321×261 image.



- The image is considered as a very long 1D vector by concatenating image pixels column by column (or alternatively row by row), i.e. 321 × 261 = 83781.
- The huge number 83781 is the dimensionality of our vector space.
- The intensity variation is assumed in each pixel of the image.

- First, to know that it is valid to use PCA on images, I read PCA Application to images by Dr. Hlavac from Czech Technical University in Prague
- Asserts that images can be used in PCA, as they can be converted to a one-dimensional vector by row-by-row or column-by-column concatenation
- Like our dataset, a grayscale image is a matrix of values with pixels corresponding to a value (the image on the slide represents intensity in the photo, our dataset a pixel represents a corresponding temperature)

Achieving a Desirable Variance

Code for Graph on Next Slide

from sklearn.preprocessing import MinMaxScaler
scaler = MinMaxScaler()
data_rescaled = scaler.fit_transform(x_old)
pca2 = PCA().fit(data_rescaled)
plt.rcParams["figure.figsize"] = (30,30)

fig, ax = plt.subplots()
xi = np.arange(0, 2674, step=1)
y = np.cumsum(pca2.explained_variance_ratio_)

plt.ylim(0.0,1.1)
plt.plot(xi, y, marker='o', linestyle='--', color='b')

plt.xlabel('Number of Components')
plt.xticks(np.arange(0, 2674, step=40))
plt.ylabel('Cumulative variance (%)')
plt.title('The number of components needed to explain variance')

plt.axhline(y=0.95, color='r', linestyle='-')
plt.text(0.5, 0.85, '95% cut-off threshold', color = 'red', fontsize=16)

ax.grid(axis='x')

- PCA wants to account for the most variability possible in the dataset so that you can get unique features of the dataset
 - Choosing a number of components is important for PCA because it can yield a certain of variance
 - 95% variance seems to be a common for PCA models
- Instead of manually picking components, we can use visualization to figure out how many components to pick

Variance Formula

 $\sum (x_i - \bar{x})^2$ S^2 -

Choosing Components

from sklearn import decomposition
from sklearn.decomposition import PCA

pca = PCA(n_components=40)



Finding Important Features

Find important features

```
most_important_features = list()
for component in pca.components_:
    index = 0
    tempList = list()
    for feature in component:
    row = index // 250
    column = index % 250
    tempList.append((abs(feature) , (row, column)))
    index += 1
```

tempList.sort(reverse=True)
most_important_features.append([tempList[x] for x in range(10)])

- For each component, every feature has a magnitude of its corresponding values of its eigenvector
 - Bigger the magnitude, the more important it is
- To find these important components
 - Take the absolute value to get a magnitude
 - Sort by largest to smallest
 - Resulting in most important features being at top of list
- Added a (row, column) to see side-by-side how important each pixel is
- Took the top 10 most influential pixels for each component

Summary of PCA Results

Total X Ave = 108.21951219512195 Total Y Ave = 57.46585365853658

Component 6

[(0.09226205001661351, (191, 42)), (0.08857704421238168, (141, 42)), (0.07783892723328956, (1, 117)), (0.0743088303359767, (106, 112)), (0.06: X Mode = [(191, 1), (141, 1), (1, 1), (106, 1), (217, 1), (10, 1), (28, 1), (4, 1), (58, 1), (172, 1)] | Y Mode = [(42, 4)] X Ave = 92.8 | Y Ave = 92.1

Component 7

[(0.16035758617914145, (158, 42)), (0.1129284033191318, (170, 42)), (0.11148113374397745, (0, 42)), (0.1112104781257541, (75, 42)), (0.105116: X Mode = [(158, 1), (170, 1), (0, 1), (75, 1), (244, 1), (221, 1), (53, 1), (40, 1), (194, 1), (97, 1)] | Y Mode = [(42, 6)] X Ave = 125.2 | Y Ave = 54.0

Component 8

[(0.1812990210483322, (17, 42)), (0.1702388360333583, (25, 42)), (0.1615332465251595, (0, 42)), (0.1504909244751474, (182, 42)), (0.140310415! X Mode = [(17, 1), (25, 1), (0, 1), (182, 1), (43, 1), (171, 1), (188, 1), (75, 1), (53, 1), (111, 1)] | Y Mode = [(42, 8)] X Ave = 86.5 | Y Ave = 75.5

Component 9

[(0.1004579267898733, (213, 42)), (0.00376278149000422, (44, 42)), (0.08948275930191729, (9, 26)), (0.08888075600852027, (220, 214)), (0.08661 X Mode = [(213, 1), (44, 1), (9, 1), (220, 1), (0, 1), (209, 1), (207, 1), (43, 1), (64, 1), (53, 1)] | Y Mode = [(42, 5)] X Ave = 106.2 | Y Ave = 81.2

Component 10

[(0.16475976295418504, (217, 42)), (0.12881073712126304, (2, 42)), (0.11935895535266505, (31, 42)), (0.11705941189163369, (191, 42)), (0.1142: X Mode = [(217, 1), (2, 1), (31, 1), (191, 1), (0, 1), (140, 1), (182, 1), (42, 1), (101, 1), (170, 1)] | Y Mode = [(42, 9)] X Ave = 107.6 | Y Ave = 37.9

Component 11

[(0.15102827816400138, (138, 68)), (0.13027190458561, (109, 42)), (0.11129183941181499, (44, 42)), (0.10124925478676165, (76, 42)), (0.0976921 X Mode = [(138, 1), (109, 1), (44, 1), (76, 1), (13, 1), (111, 1), (201, 1), (140, 1), (30, 1), (82, 1)] | Y Mode = [(42, 8)] X Ave = 94.4 | Y Ave = 41.7

- For Modes (position, instances)
 - Position is either its x or y position
 - Instances is how many time that position occurred
- Can see the magnitude of an eigenvector (importance) next to its row, column coordinates
- Took x and y averages of top 10 most important features
- Took averages of the 40 componentes x and y averages
- y = 42, showed up a lot as a mode

Code for Summary of PCA Results

```
def findMode(important_component):
```

```
x_dict = dict()
y_dict = dict()
high_x_instances = 1
high_y_instances = 1
for feature in important_component:
    x_feat = feature[1][0]
    y_feat = feature[1][1]
    try:
        x_dict[x_feat] += 1
        if high_x_instances < x_dict[x_feat]:
        high_x_instances = x_dict[x_feat]
except:
        x_dict[x_feat] = 1</pre>
```

try: y_dict[y_feat] += 1 if high_y_instances < y_dict[y_feat]: high_y_instances = y_dict[y_feat] except: y_dict[y_feat] = 1 x_modes = list() y_modes = list() for key, value in x dict.items():

if value == high x instances:

for key, value in y dict.items():

return x modes, y modes

x modes.append((key, value))

```
def findAve(important_component):
    x_ave = 0
    y_ave = 0
    index = 0
    for feature in important_component:
        x_ave += feature[1][0]
        y_ave += feature[1][1]
        index += 1
    return x_ave/index, y_ave/index
```

```
# Show Results
component_num = 1
x_total_ave = 0
y_total_ave = 0
for component in most_important_features:
    print("Component {}".format(component_num))
    print(component)
    modes = findMode(component)
    aves = findAve(component)
    print("X Mode = {0} | Y Mode = {1}".format(modes[0], modes[1]))
    print("X Ave = {0} | Y Ave = {1}\n".format(aves[0], aves[1]))
    x_total_ave += aves[0]
    y_total_ave += aves[1]
    component_num += 1
print("Total X Ave = {0}".format(x total ave/component num))
```

print("Total Y Ave = {0}".format(y_total_ave/component_num))

Random Forest Exploration

- 1. Why further explore Random Forest Classifier
- 2. Understanding of Vectorizing a Matrix
- 3. More testing and results of Random Forest for various parameters

Why further explore Random Forest Classifier?

- Had very high accuracy (above 99% accuracy in classifying)
- Had very fast time compared to other ensemble methods
- Test to see if it can if the first trial was dumb luck or if it can be repeated
- Address whether vectorizing a matrix would lose relationships from the data

| RandomEorest(| lassifier | · - · | | 1 |
|-----------------------|--------------|--------|----------|---------|
| Accuracy : 0. | 998878923766 | 8162 | | |
| CV Score : 0. | 996153846153 | 8464 | | |
| AUC Score : | 0.0 | | | |
| | precision | recall | f1-score | support |
| 0 | 1.00 | 1.00 | 1.00 | 465 |
| 1 | 1.00 | 1.00 | 1.00 | 427 |
| accuracy | | | 1.00 | 892 |
| macro avg | 1.00 | 1.00 | 1.00 | 892 |
| weighted avg | 1.00 | 1.00 | 1.00 | 892 |
| [[464 1] [0 427]] | | | | |

Time Taken : 15.212837219238281 seconds

Vectorizing a Matrix Keeps Order and Relationship



Testing Random Forest Classifier

<u>Procedure</u>

- 100 Total Trials
- 25 Trials each for
 - 50/50 Train/Test Split
 - 60/40 Train/Test Split
 - 70/30 Train/Test Split
 - 80/20 Train/Test Split
- Every trial a new data split will be calculated to ensure random splits

<u>Outcomes</u>

For each type of train/test split over 25 trials:

- 1. Average Accuracy Score
- 2. Average Cross Validation Score
- 3. Average Time Taken per Trial

Code for Testing Random Forest Classifier

def splitData(split_num):

xTrain, xTest, yTrain, yTest = train_test_split(cropInData, outData, test_size = split_num, random_state = 0) dataset_size = len(xTrain) test_size = len(xTest) xTrain2 = np.array(xTrain) xTrain2 = np.expand_dims(xTrain2, -1) xTest2 = np.array(xTest) xTest2 = np.array(yTest) xTest2 = np.array(yTrain) yTest2 = np.array(yTrain) yTest2 = np.array(yTrain) yTest2 = np.array(yTest) X_train3 = xTrain2.reshape(dataset_size, -1) Y_train3 = yTrain2.reshape(dataset_size, -1) X_test3 = xTest2.reshape(test_size, -1) Y_test3 = yTest2.reshape(test_size, -1) return X_train3, Y_train3, X_test3, Y_test3

| ave_time_1 = 0 |
|----------------|
| ave_time_2 = 0 |
| ave_time_3 = 0 |
| ave_time_4 = 0 |
| accu_1 = 0 |
| accu_2 = 0 |
| accu_3 = 0 |
| accu_4 = 0 |
| cv1 = 0 |
| cv2 = 0 |
| cv3 = 0 |
| cv4 = 0 |

for x in range(100): model = RandomForestClassifier(n estimators =10) if x < 25: # 50/50 Train/Test Split x train, y train, x test, y test = splitData(.5) elif x < 50: # 60/40 Train/Test Split x train, y train, x test, y test = splitData(.4) elif x < 75: # 70/30 Train/Test Split x train, y train, x test, y test = splitData(.3) else: # 80/20 Train/Test Split x_train, y_train, x_test, y_test = splitData(.2) t0 = time.time()model.fit(x train,y train) y pred = model.predict(x test) proba = model.predict proba(x test) roc_score = roc_auc_score(y_test, proba[:,1]) cv score = cross val score(model, x train, y train, cv=10).mean() score = accuracy_score(y_test,y_pred) bin clf rep = classification report(y test, y pred, zero division=1) if x < 25: accu 1 += score ave_time_1 += time.time()-t0 cv1 += cv score elif x < 50: accu 2 += score ave time 2 += time.time()-t0 cv2 += cv score elif x < 75: accu 3 += score ave_time_3 += time.time()-t0 cv3 += cv score else: accu 4 += score ave time 4 += time.time()-t0 cv4 += cv score print("Trial {0} with accuracy of {1}\n".format(x+1, score))

Results for Testing Random Forest

50/50 Split

| Ave | Accuracy = | 0.9956931359353969 | |
|-----|------------|----------------------|---------|
| Ave | CV Score = | 0.9948781062942138 | |
| Ave | Time Taken | = 10.085623331069947 | seconds |
| | | | |

60/40 Split

Ave Accuracy = 0.9982169890664423 Ave CV Score = 0.9948219195279643 Ave Time Taken = 12.004673089981079 seconds

70/30 Split

Ave Accuracy = 0.997892376681614 Ave CV Score = 0.9959807692307691 Ave Time Taken = 13.799284038543702 seconds

80/20 Split

Ave Accuracy = 0.9975126050420168 Ave CV Score = 0.9963454242456479 Ave Time Taken = 15.548892192840576 seconds Highest average accuracy was 60/40 Train/Test split
 Accuracy of 99.821...%

- Lowest average accuracy was 50/50 Train/Test split
 - Accuracy of 99.569...%
- Time went up with the greater the split towards training
 - Logically, this makes sense as the model has more to data values to train on, the longer it will take

Visualizing a Decision Tree in a Random Forest Classifier



Convert to png using system command (requires Graphviz)
call(['dot', '-Tpng', 'tree.dot', '-o', 'tree.png', '-Gdpi=600'])

Display in jupyter notebook
Image(filename = 'tree.png')

Trying More Neural Networks

- 1. Overview of what parameters and architectures I was using
- 2. Overview of code and results of a currently successful model
- 3. Need for more testing

Trying different Models Tried

- A lot of time was spent editing different parameters from last weeks neural network (ave 79% test accuracy over 10 trials)
- Different parameters included:
 - Epochs
 - Batch Size
 - Learning Rate
 - Optimizers
 - Activation Functions
 - Many Hidden Layers
 - Few Hidden Layers
 - Different Types of Layer
 - Batch Normalization
 - Dropout Rates

A Promising Result

```
478/478 [======] - 0s 656us/step
    test loss, test acc: [0.0007525471131177035, 1.0]
1.
     478
    Actual value = 1 | Prediction = 0.9998002648353577 | Precition Rounded = 1.0
     478/478 [=======] - 0s 661us/step
     test loss, test acc: [0.0010028247639661553, 1.0]
2.
     478
     Actual value = 1 | Prediction = 0.9997267127037048 | Precition Rounded = 1.0
     Actual value = 0 | Prediction = 4.4352535041980445e-05 | Precition Rounded = 0.0
     478/478 [=========] - 0s 658us/step
     test loss, test acc: [0.0006307436167418364, 1.0]
3.
     478
     Actual value = 1 | Prediction = 0.9997641444206238 |
                                                        Precition Rounded = 1.0
     Actual value = 0 | Prediction = 0.0001633794599911198 | Precition Rounded = 0.0
     test loss, test acc: [0.0006265214261517053, 1.0]
4.
     478
     Actual value = 1 | Prediction = 0.999890923500061
                                                           Precition Rounded = 1.0
                                                     coronona | possible
      478/478 [======] - 0s 662us/step
      test loss, test acc: [0.0006233096327564472, 1.0]
5.
      478
      Actual value = 1 | Prediction = 0.9998434782028198 |
                                                      Precition Rounded = 1.0
```

5 Trials of

- 70/30 Train/Test Split
 - Every trial had a Ο different split
- Every trail had a test accuracy of 100%
- 20 Epochs
- Batch Size of 32

The Model's Architecture / Code

def model5():

model = Sequential()

Input Layer

model.add(Conv2D(32, kernel_size = (3, 3), activation='relu', input_shape=(250, 250, 1)))
model.add(MaxPooling2D(pool_size=(2,2)))
model.add(BatchNormalization())

Hidden 1

model.add(Conv2D(64, kernel_size=(3,3), activation='relu'))
model.add(MaxPooling2D(pool_size=(2,2)))
model.add(BatchNormalization())

Hidden 2

model.add(Conv2D(64, kernel_size=(3,3), activation='relu'))
model.add(MaxPooling2D(pool_size=(2,2)))
model.add(BatchNormalization())

Hidden 3

model.add(Conv2D(96, kernel_size=(3,3), activation='relu'))
model.add(MaxPooling2D(pool_size=(2,2)))
model.add(BatchNormalization())

Hidden 4

model.add(Conv2D(32, kernel_size=(3,3), activation='relu'))
model.add(MaxPooling2D(pool_size=(2,2)))
model.add(BatchNormalization())
model.add(Oropout(0.2))

Hidden 5

model.add(Flatten())
model.add(Dense(128, activation='relu'))

Output Layer
model.add(Dense(1, activation = 'sigmoid'))

Compile Model

sgd = SGD(lr = .01)
model.compile(loss = 'binary_crossentropy', optimizer = sgd, metrics = ['accuracy'])
model.summary()
return model

Model: "sequential_14"

| Layer (type) | Output | Shape | Param # |
|------------------------------|--------|---------------|---------|
| conv2d_54 (Conv2D) | (None, | 248, 248, 32) | 320 |
| max_pooling2d_50 (MaxPooling | (None, | 124, 124, 32) | 0 |
| batch_normalization_58 (Batc | (None, | 124, 124, 32) | 128 |
| conv2d_55 (Conv2D) | (None, | 122, 122, 64) | 18496 |
| max_pooling2d_51 (MaxPooling | (None, | 61, 61, 64) | 0 |
| batch_normalization_59 (Batc | (None, | 61, 61, 64) | 256 |
| conv2d_56 (Conv2D) | (None, | 59, 59, 64) | 36928 |
| max_pooling2d_52 (MaxPooling | (None, | 29, 29, 64) | 0 |
| batch_normalization_60 (Batc | (None, | 29, 29, 64) | 256 |
| conv2d_57 (Conv2D) | (None, | 27, 27, 96) | 55392 |
| max_pooling2d_53 (MaxPooling | (None, | 13, 13, 96) | 0 |
| batch_normalization_61 (Batc | (None, | 13, 13, 96) | 384 |
| conv2d_58 (Conv2D) | (None, | 11, 11, 32) | 27680 |
| max_pooling2d_54 (MaxPooling | (None, | 5, 5, 32) | 0 |
| batch_normalization_62 (Batc | (None, | 5, 5, 32) | 128 |
| dropout_26 (Dropout) | (None, | 5, 5, 32) | 0 |
| flatten_14 (Flatten) | (None, | 800) | 0 |
| dense_27 (Dense) | (None, | 128) | 102528 |
| dense_28 (Dense) | (None, | 1) | 129 |

Non-trainable params: 576

- Uses 'blocks' of Conv2D, MaxPooling2D, and Batch Normalization
- Finally flattens to a dense layer
- Uses SGD optimizer
- Every layer but the final layer uses relu
- Final layer uses sigmoid

Need to Further Test This Model

- Got first promising results for this model yesterday (6/24)
- Need to test:
 - Different train/test split ratios
 - More Epoch Sizes
 - Originally tested 5 epochs, that had lower accuracy scores
 - 20 Epochs was the text value tested, which yielded very good accuracy scores
 - Batch Size
- Essentially, test this model more thoroughly like the Random Forest Classifier

Future Steps and Goals



Álso Google Colab has Corgi Mode, it's great =)

- Test the new neural network model more thorough and give results
- What to do with PCA data / results?
- Suffering from Black Box Testing when construction Neural Networks
 - Advice?
- Ultimate Goal of Project Clarification
 - Should the final model be a neural network, or whatever ascertains a high classification accuracy?
- Any obvious next steps?