Random Forest, PCA and KNN-MNIST dataset

Siyuan Zhang

1 Resources

Student: Siyuan Zhang Language: Python

Code: https://github.com/Siyuan-gwu/Compare-Random Forest-PCA-KNN-MNIST-

dataset

Instruction: The instruction to run the code is on github (readme)

Resource:

1. https://www.kaggle.com/sflender/comparing-random-forest-pca-and-knn

2. https://iq.opengenus.org/algorithm-principal-component-analysis-pca/

import numpy as np

import pandas as pd

import seaborn as sb

import os

import matplotlib.pyplot as plt, matplotlib.image as mpimg

from sklearn.ensemble import RandomForestClassifier

from sklearn.model_selection import train_test_split

from sklearn.neighbors import KNeighborsClassifier

from sklearn.decomposition import PCA

import pylab

2 Dataset details

There are two datasets, training data and testing data

- 1. The training data contains 28000 images
- 2. Each image has 784 pixels in total, each single pixel-value indicates the lightness or darkness of that pixel
- 3. The first column is the actual digit of the image
- 4. The testing data is the same with training data, but without "label" column to represent the actual number

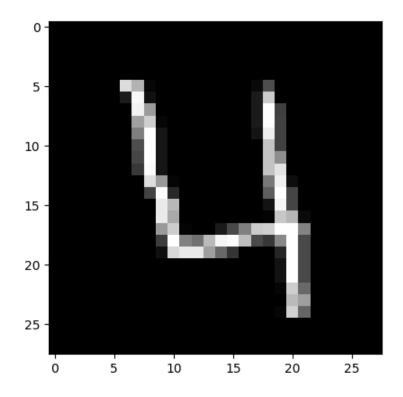
Inspect the Dataset

#read the data									
train = pd.read_csv('train.csv')									
print (train.head(5))									
	label	pixel0	pixel1	pixel2		pixel780	pixel781	pixel782	pixel783
0	1	0	0	0		0	0	0	0
1	0	0	0	0		0	0	0	0
2	1	0	0	0		0	0	0	0
3	4	0	0	0		0	0	0	0
4	0	0	0	0		0	0	0	0

[5 rows x 785 columns]

Print an example of the number

```
#plot a number "4"
img = train.values[3]
img = img.reshape(28,28)
plt.imshow(img, cmap='gray')
plt.show()
```



Data pre-processing

Drop the label of train dataset.

```
train = train.drop("label", 1)
```

Do the data normalization, if the pixel larger than one, then change it to one. Otherwise, do nothing.

Check the performance of random forest classifier

```
def random_forest():
  # loading training data
  print('Loading training data')
  X_tr = train.values[:, 1:].astype(float)
  y_tr = train.values[:, 0]
  scores = list()
  scores_std = list()
  print('Start learning...')
  n_trees = [10, 15, 20, 25, 30, 40, 50, 70, 100, 150]
  for n_tree in n_trees:
    print(n_tree)
    recognizer = RandomForestClassifier(n_tree)
     score = cross_val_score(recognizer, X_tr, y_tr)
    scores.append(np.mean(score))
     scores_std.append(np.std(score))
  sc_array = np.array(scores)
  std_array = np.array(scores_std)
  print('Score: ', sc_array)
  print('Std : ', std_array)
  plt.figure(figsize=(4,3))
  plt.plot(n_trees, scores)
  plt.plot(n_trees, sc_array + std_array, 'b--')
  plt.plot(n_trees, sc_array - std_array, 'b--')
```

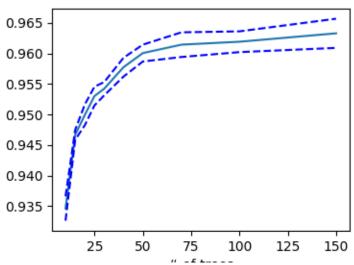
```
plt.ylabel('CV score')

plt.xlabel('# of trees')

plt.savefig('cv_trees.png')

plt.show()
```

Result:



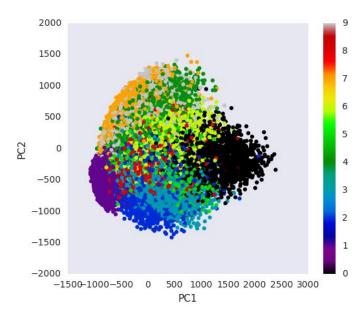
The x-axis is the number of trees
The y-axis is the rate of accuracy

Try on PCA and KNN (we use component = 2)

```
#try on PCA
pca = PCA(n_components=2)
pca.fit(train)
transform = pca.transform(train)

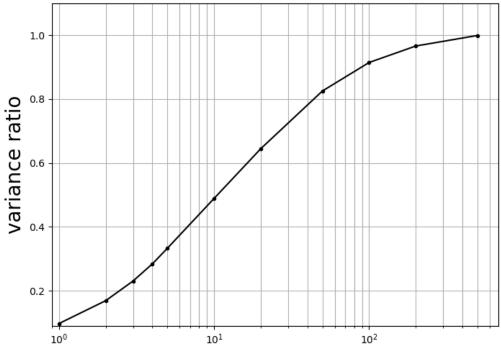
plt.figure()
plt.scatter(transform[:,0],transform[:,1], s=20, c = target, cmap = "nipy_spectral", edgecolor = "None")
plt.colorbar()
plt.clim(0,9)

plt.xlabel("PC1")
plt.ylabel("PC2")
```



We notice that PCA separates the feature space into visible clusters already for 2 components. Next, I am going to check how many components are needed to capture most of the variance in the data.

```
n_components_array=([1,2,3,4,5,10,20,50,100,200,500])
vr = np.zeros(len(n_components_array))
i=0;
for n_components in n_components_array:
  pca = PCA(n_components=n_components)
  pca.fit(train)
  vr[i] = sum(pca.explained_variance_ratio_)
  i=i+1
plt.figure(figsize=(8,4))
plt.plot(n_components_array,vr,'k.-')
plt.xscale("log")
plt.ylim(9e-2,1.1)
plt.xlim(0.9)
plt.grid(which="both")
plt.xlabel("number of PCA components",size=20)
plt.ylabel("variance ratio",size=20)
plt.show()
```



number of PCA components

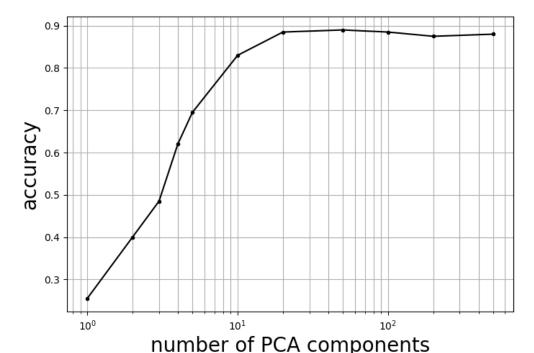
We notice that ~100 PCA components are needed to capture ~90% of the variance in the data. But I cannot know the prediction result from this plot. Next, I would train a KNN classifier on the PCA output.

```
clf = KNeighborsClassifier()
n_components_array=([1,2,3,4,5,10,20,50,100,200,500])
score_array = np.zeros(len(n_components_array))
i=0

for n_components in n_components_array:
    pca = PCA(n_components=n_components)
    pca.fit(train)
    transform = pca.transform(train.iloc[0:1000])
    score_array[i] = evaluate_classifier(clf, transform, target.iloc[0:1000], 0.8)
    i=i+1

plt.figure(figsize=(8,4))
plt.plot(n_components_array,score_array,'k.-')
plt.xscale('log')
plt.xlabel("number of PCA components", size=20)
plt.ylabel("accuracy", size=20)
```

```
plt.grid(which="both")
plt.show()
```



We notice that the accuracy seems to saturate at \sim 90% for $>\sim$ 20 PCA components. In fact, the accuracy even seems to drop for much larger numbers, even though a larger number of PCA components captures more of the variance in the data. Maybe the drop of the accuracy is probably due to overfitting.

Result

Submit the knn+PCA and Random Forest test result on Kaggle

Get the result score: 0.95

```
#PCA and KNN test result

pca = PCA(n_components=50)

pca.fit(train)

transform_train = pca.transform(train)

transform_test = pca.transform(test)

clf = KNeighborsClassifier()

clf.fit(transform_train, target)

results = clf.predict(transform_test)

np.savetxt("result_knn_pca.csv',

np.c_[range(1, len(test) + 1), results],

delimiter=',',

header='ImageId,LabeI',

comments=",

fmt='%d')
```

Get the result score: 0.964

It seems that the PCA and KNN accuracy is higher.

3 Algorithm Description

PCA algorithm

Principal component analysis (PCA) is a technique to bring out strong patterns in a dataset by suppressing variations. It is used to clean data sets to make it easy to explore and analyze. The algorithm of Principal Component Analysis is based on a few mathematical ideas namely:

- 1. Variance and Covariance
- 2. Eigen Vectors and Eigen values

Algorithm steps:

- 1. Get your data
- 2. Give your data a structure
- 3. Standardize your data
- 4. Get covariance of matrix Z
- 5. Calculate Eigen Vectors and Eigen Values
- 6. Sort the Eigen Vectors

- 7. Calculate the new features
- 8. Drop unimportant features from the set

4 Runtime

For PCA:

Suppose we have n data points, and each represented with p features. Covariances matrix computation is $O(p^2*n)$, its eigen-value decomposition is $O(p^3)$. So, the time complexity of PCA is $O(p^2*n+p^3)$