# **Lab Assignment Two**

# **Crops disease**

**Group: Lone Wolf** 

Member: Yang Shen

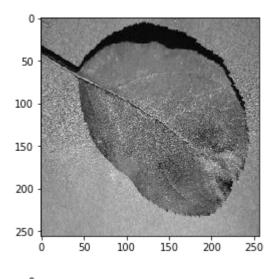
This dataset is containing the images of 10 different crops' leaves. For each crop, there are subfiles containing images about healthy and unhealthy leaves. The same leaf will take the photo from different angles. The main purpose of this dataset is to classify different leaves by their categories. We can know the name of the crop and whether the crop is getting the disease by analyzing the leaves photo.

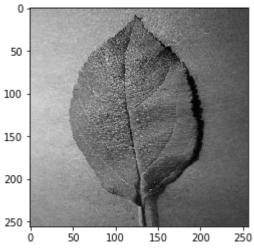
The third party who interested in this result is seeds company. They sell millions of seeds over a year. The quality of seeds is important for the company. Seeds company usually sell some anti-disease seeds in the market. But the performance of anti-disease seeds needs to keep tracking. Farmers can upload the photos to seeds company for daily bases. The seeds company can use our method to find out the category of leaves and what disease the crops got, or crops are healthy. So that they can give advice about how to deal with sick crops to farmers. Also, they can monitor the performance of their seeds.

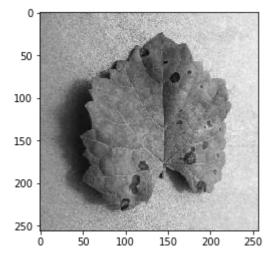
The performance of our model needs to be accurate in order to find out which disease is. The high true negative rate will help the company because they can save the workload on healthy crops. They do not need to send people to check for healthy crops. For the false positive rate, it does not need to be 0%. Because when the machine processes the images, if crops are sick, they will be check by a human. Because farmers need advice from a specialized person. So that if the false positive rate is around 20%, it means there will be 20% more workload for employees. I think 20% extra workload is affordable for seeds company. However, if false positive rate is high, it means that the company will mark the sick crops into healthy category. It will cause money to lose for farmers and cause the legal issue. However, if farmers upload multiple images for the same crop, only one of the images marked as sick, the whole crop will be marked as sick and send human to check. I think for real sick crops if farmer upload 10 images, one or two images are incorrectly marked as healthy and rest of images is marked as unhealthy. It is acceptable. So that I think false negative rate cannot high than 10%. The seeds company can tell farmers to upload more photos for the same crops to increase the chance to find out sick crops. I think set the minimum photo for one crop to 5 is good.

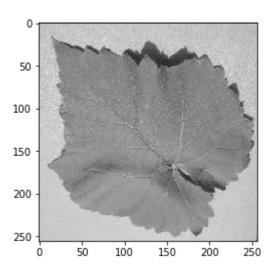
# In [1]:

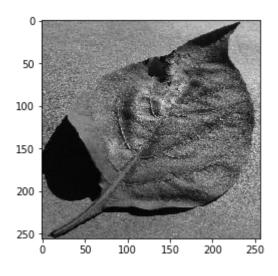
```
#https://www.youtube.com/watch?v=j-3vuBynn0E
%matplotlib inline
from matplotlib import pyplot as plt
import pandas as pd
import numpy as np
import os
import cv2
datadir = "C:/Users/jacks/Desktop/plant"
Category = ["Apple_Black_rot", "Apple_healthy", "Grape_Black_rot", "Grape_healthy", "Pepper_Bacteria
1 spot", "Pepper healthy", "Potato healthy", "Potato Late blight", "Tomato Bacterial spot", "Tomato h
ealthy"]
for i in Category:
    path = os. path. join(datadir, i) #path to the leaf data
    for img in os.listdir(path):
        img array = cv2. imread(os. path. join(path, img), cv2. IMREAD GRAYSCALE) # read all the leave
s file from each sub file
        plt. imshow(img array, cmap="gray") #change to black and white images
        plt.show()
        break
                #show some images not all
```

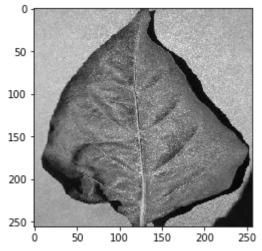


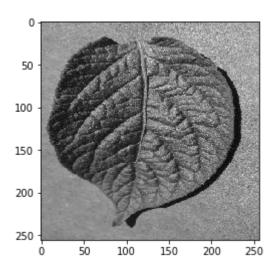


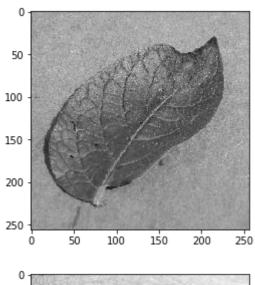


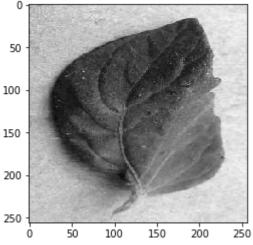


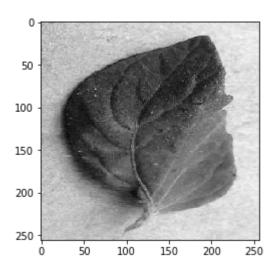










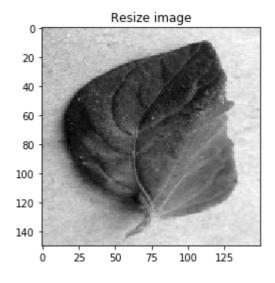


# In [2]:

```
print('original size of the images:',img_array.shape)
print('======')

#show the resize of images
imagesize = 150
resize_array = cv2.resize(img_array, (imagesize,imagesize))
plt.imshow(resize_array, cmap = 'gray')
plt.title('Resize image')
plt.show()
```

original size of the images: (256, 256)



Ther original images are 256 X 256, resize to 150 X 150 is also clear to see the image, and it will reduce the total size of array.

# In [3]:

```
images=[] #just reimport the data from the files and resize the images as well
def creat_data():
    for i in Category:
        path = os. path. join(datadir, i)
        class_num = Category. index(i) #set a index for the column which will help me to set the
    names for different leaves
    for img in os. listdir(path):
        img_array = cv2. imread(os. path. join(path, img), cv2. IMREAD_GRAYSCALE)
        resize_array = cv2. resize(img_array, (imagesize, imagesize))
        images. append([resize_array, class_num]) # append the 1-d images data and the column index
creat_data()
```

# In [4]:

```
#create the array contain the images data with 1-D image features
X = []
y = []
#seprate the 1-d images data and column values
for features, label in images:
    X. append(features)
    y. append(label)

names = y. copy()
#reshape the array into the correct format, which each row represent one images
X_images = np. array(X). reshape(-1, imagesize, imagesize)
X = np. array(X_images). reshape(len(images), 22500)
_, h, w = X_images. shape
```

#### In [5]:

```
#https://stackoverflow.com/questions/2582138/finding-and-replacing-elements-in-a-list-python
# replace the index value to the names of leaf
item to replace = 0
replacement value = "Apple Black rot" #name of leaf
indices to replace = [i for i, x in enumerate(names) if x == item to replace]
for i in indices to replace:
    names[i] = replacement_value
item to replace = 1
replacement value = "Apple healthy"
indices to replace = [i for i, x in enumerate(names) if x == item to replace]
for i in indices to replace:
    names[i] = replacement_value
item to replace = 2
replacement value = "Grape Black rot"
indices_to_replace = [i for i, x in enumerate(names) if x==item_to_replace]
for i in indices_to_replace:
    names[i] = replacement value
item_to_replace = 3
replacement value = "Grape healthy"
indices_to_replace = [i for i, x in enumerate(names) if x==item_to_replace]
for i in indices to replace:
    names[i] = replacement value
item_to replace = 4
replacement_value = "Pepper_Bacterial_spot"
indices_to_replace = [i for i, x in enumerate(names) if x==item_to_replace]
for i in indices to replace:
    names[i] = replacement value
item to replace = 5
replacement value = "Pepper healthy"
indices to replace = [i for i, x in enumerate(names) if x == item to replace]
for i in indices to replace:
    names[i] = replacement value
item to replace = 6
replacement value = "Potato healthy"
indices_to_replace = [i for i, x in enumerate(names) if x==item_to_replace]
for i in indices to replace:
    names[i] = replacement_value
item to replace = 7
replacement_value = "Potato_Late_blight"
indices to replace = [i for i, x in enumerate(names) if x == item to replace]
for i in indices to replace:
    names[i] = replacement_value
item to replace = 8
```

```
replacement_value = "Tomato_Bacterial_spot"
indices_to_replace = [i for i, x in enumerate(names) if x==item_to_replace]

for i in indices_to_replace:
    names[i] = replacement_value

item_to_replace = 9
replacement_value = "Tomato_healthy"
indices_to_replace = [i for i, x in enumerate(names) if x==item_to_replace]

for i in indices_to_replace:
    names[i] = replacement_value
```

# In [6]:

```
n_samples, n_features = X. shape
_, h, w = X_images. shape

print("n_samples: {}".format(n_samples))
print("n_features: {}".format(n_features))

print("Image Sizes {} by {}".format(h, w))
print (150*150) # the size of the images are the size of the feature vectors
```

n\_samples: 4231 n\_features: 22500 Image Sizes 150 by 150 22500

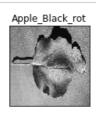
#### In [7]:

```
# a helper plotting function
#since the data read by file, here is all leaves from first file
def plot_gallery(X_images, titles, h, w, n_row=3, n_col=6):
    """Helper function to plot a gallery of portraits"""
    plt.figure(figsize=(1.7 * n_col, 2.3 * n_row))
    plt.subplots_adjust(bottom=0, left=.01, right=.99, top=.90, hspace=.35)
    for i in range(n_row * n_col):
        plt.subplot(n_row, n_col, i + 1)
        plt.imshow(X_images[i].reshape((h, w)), cmap=plt.cm.gray)
        plt.title(names[i], size=12)
        plt.xticks(())
        plt.yticks(())
```





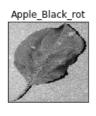






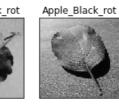


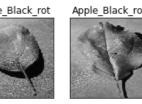


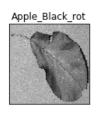


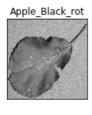


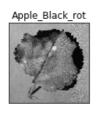


















# **PCA**

# In [8]:

```
from sklearn.decomposition import PCA

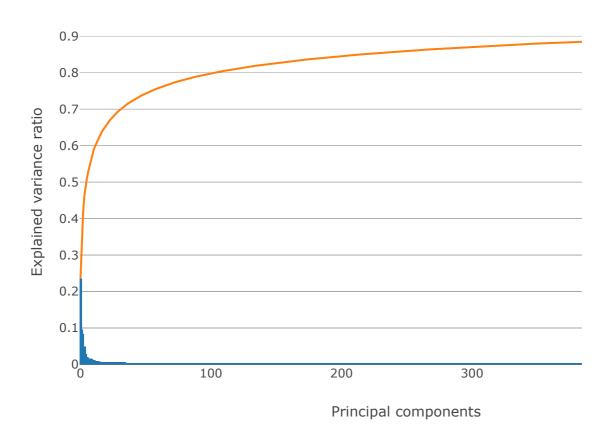
n_components = 500
print ("Extracting the top %d eigenleaves from %d leaves" % (
    n_components, X. shape[0]))

pca = PCA(n_components=n_components)
%time pca.fit(X.copy())
eigenleaves = pca.components_.reshape((n_components, h, w))
```

Extracting the top 500 eigenleaves from  $4231\ leaves$  Wall time:  $7.\,83\ s$ 

# In [9]:

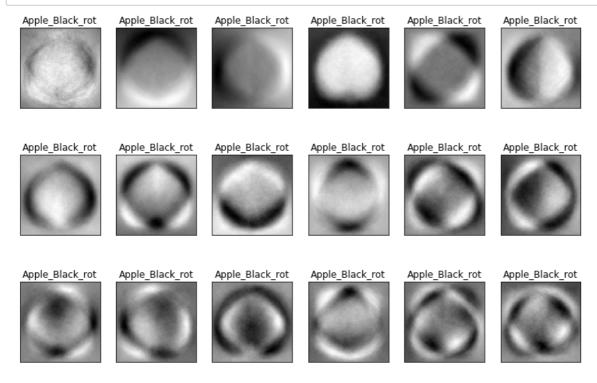
```
def plot explained variance(pca):
    import plotly
    from plotly.graph_objs import Bar, Line
    from plotly.graph_objs import Scatter, Layout
    from plotly.graph objs.scatter import Marker
    from plotly.graph_objs.layout import XAxis, YAxis
    plotly. offline. init notebook mode() # run at the start of every notebook
    explained_var = pca.explained_variance_ratio_
    cum var exp = np.cumsum(explained var)
   plotly. offline. iplot({
        "data": [Bar(y=explained_var, name='individual explained variance'),
                Scatter(y=cum_var_exp, name='cumulative explained variance')
        "layout": Layout(xaxis=XAxis(title='Principal components'), yaxis=YAxis(title='Explained
variance ratio'))
   })
pca = PCA(n_components)
X pca = pca. fit(X)
plot explained variance (pca)
```



I do the Full PCA analysis which is Linear Reduction. Then plotting the number of components by the explained variance, I found that with 500 components, 90% of the variance will be explained.

#### In [10]:

```
eigenleaves_titles = ["eigenleaves %d" % i for i in range(eigenleaves.shape[0])] plot_gallery(eigenleaves, eigenleaves_titles, h, w)
```



# In [11]:

```
def reconstruct_image(trans_obj, org_features):
    low_rep = trans_obj. transform(org_features)
    rec_image = trans_obj. inverse_transform(low_rep)
    return low_rep, rec_image
```

#### In [12]:

```
#https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsClassifier.html
#use KNeighbors Classifier and build to see actual classification performance
from sklearn.neighbors import KNeighborsClassifier
low_dimensional_representation, freconstructed_image = reconstruct_image(pca, X. reshape(4231, -1))
neigh = KNeighborsClassifier(n_neighbors=3, n_jobs=-1)
neigh.fit(freconstructed_image, names)
%time fpred = neigh.predict(freconstructed_image)
```

Wall time: 1min 40s

# In [13]:

```
import numpy as np
faccuracy = 0
for i in range(len(fpred)):
    faccuracy = float(sum(fpred[i] == names[i] for i in range(len(names)))) / len(names)
print('accuracy of the classifier is: ',faccuracy)
```

accuracy of the classifier is: 0.6960529425667691

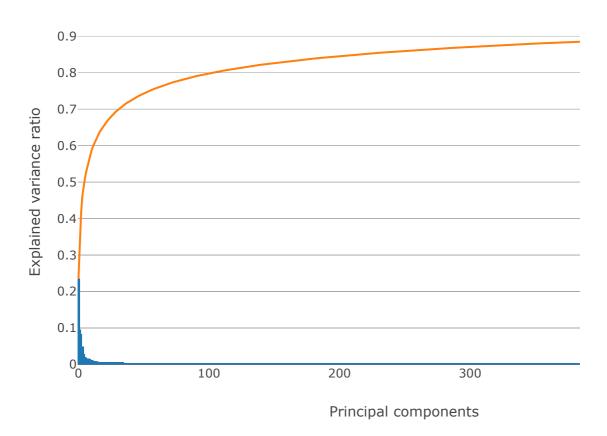
# **Randomized PCA**

# In [10]:

Extracting the top 500 eigenleaves from 4231 leaves Wall time:  $7.86~\mathrm{s}$ 

# In [11]:

```
def plot explained variance(rpca):
    import plotly
    from plotly.graph_objs import Bar, Line
    from plotly.graph_objs import Scatter, Layout
    from plotly.graph objs.scatter import Marker
    from plotly.graph_objs.layout import XAxis, YAxis
    plotly. offline. init notebook mode() # run at the start of every notebook
    explained_var = rpca.explained_variance_ratio_
    cum var exp = np.cumsum(explained var)
    plotly. offline. iplot({
        "data": [Bar(y=explained_var, name='individual explained variance'),
                 Scatter(y=cum_var_exp, name='cumulative explained variance')
        "layout": Layout(xaxis=XAxis(title='Principal components'), yaxis=YAxis(title='Explained
variance ratio'))
   })
rpca = PCA(n_components)
X \text{ rpca} = \text{rpca. fit}(X)
plot explained variance (rpca)
```



I do the Randomized PCA analysis which is Linear Reduction. Then plotting the number of components by the explained variance, I found that with 500 components, 90% of the variance will be explained. This result is pretty similar to the full pca result.

```
In [16]:
```

```
def reconstruct_image(trans_obj, org_features):
    low_rep = trans_obj. transform(org_features)
    rec_image = trans_obj. inverse_transform(low_rep)
    return low_rep, rec_image
```

#### In [17]:

```
#https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsClassifier.html
from sklearn.neighbors import KNeighborsClassifier
low_dimensional_representation, rreconstructed_image = reconstruct_image(rpca, X. reshape(4231, -1))
neigh = KNeighborsClassifier(n_neighbors=3, n_jobs=-1)
neigh.fit(rreconstructed_image, names)
%time rpred = neigh.predict(rreconstructed_image)
```

Wall time: 1min 38s

# In [18]:

```
import numpy as np

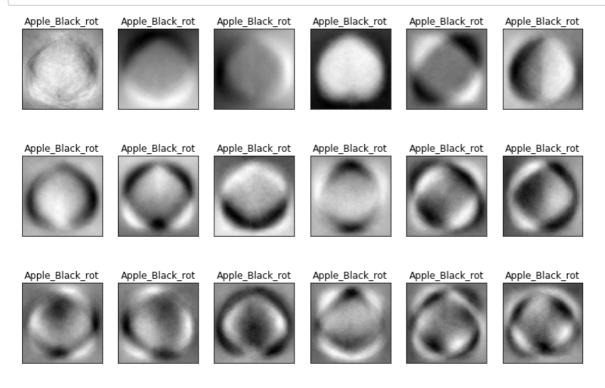
raccuracy = 0
for i in range(len(rpred)):
    raccuracy = float(sum(rpred[i] == names[i] for i in range(len(names)))) / len(names)

print('accuracy of the classifier is : ',raccuracy)
```

accuracy of the classifier is: 0.69463483809974

# In [12]:

eigenleaves\_titles = ["eigenleaves %d" % i for i in range(eigenleaves.shape[0])]
plot\_gallery(eigenleaves, eigenleaves\_titles, h, w)



The eigenvalues of the leaf is similar to the result from the full pca method. We will see the reconstructed images compare to each other in kernel pca plot.

# **Kernel PCA**

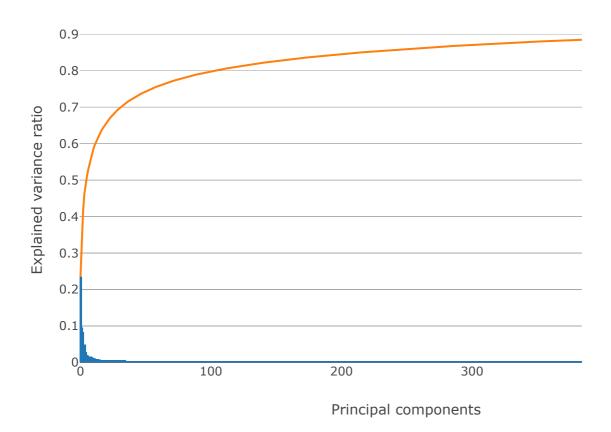
```
In [13]:
```

#### In [27]:

```
import pickle
kpca = pickle.load(open('C:/Users/jacks/Desktop/plant/large_data/kpca.p', 'rb'))
```

# In [21]:

```
def plot explained variance(kpca):
    import plotly
    from plotly.graph_objs import Bar, Line
    from plotly.graph_objs import Scatter, Layout
    from plotly.graph objs.scatter import Marker
    from plotly.graph_objs.layout import XAxis, YAxis
    plotly. offline. init notebook mode() # run at the start of every notebook
    explained_var = kpca.explained_variance_ratio_
    cum var exp = np.cumsum(explained var)
    plotly. offline. iplot({
        "data": [Bar(y=explained_var, name='individual explained variance'),
                 Scatter(y=cum_var_exp, name='cumulative explained variance')
        "layout": Layout(xaxis=XAxis(title='Principal components'), yaxis=YAxis(title='Explained
variance ratio'))
   })
kpca = PCA(n_components)
X \text{ kpca} = \text{kpca. fit}(X)
plot explained variance(kpca)
```



I do the kernel PCA analysis which is non-Linear Reduction. Then plotting the number of components by the explained variance, I found that with 500 components, 90% of the variance will be explained. This is the same as full pca and randomized pca.

# In [42]:

```
#https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsClassifier.html
from sklearn.neighbors import KNeighborsClassifier
low_dimensional_representation, kreconstructed_image = reconstruct_image(kpca, X. reshape(4231, -1))
neigh = KNeighborsClassifier(n_neighbors=3, n_jobs=-1)
neigh.fit(kreconstructed_image, names)
%time kpred = neigh.predict(kreconstructed_image)
```

Wall time: 1min 7s

#### In [43]:

```
import numpy as np

kaccuracy = 0
for i in range(len(kpred)):
    kaccuracy = float(sum(kpred[i] == names[i] for i in range(len(names)))) / len(names)

print('accuracy of the classifier is: ',kaccuracy)
```

accuracy of the classifier is: 0.5398251004490664

# In [14]:

```
from ipywidgets import widgets # make this interactive!
import warnings
# warnings. simplefilter('ignore', DeprecationWarning)
# warnings. simplefilter("always", DeprecationWarning)
def plt reconstruct(idx to reconstruct):
    idx_to_reconstruct = np. round(idx_to_reconstruct)
    reconstructed_image = pca.inverse_transform(pca.transform(X[idx_to_reconstruct].reshape(1, -
1)))
    reconstructed image rpca = rpca.inverse transform(rpca.transform(X[idx to reconstruct].resha
pe(1, -1))
    reconstructed image kpca = kpca.inverse transform(kpca.transform(X[idx to reconstruct].resha
pe(1, -1))
    plt. figure (figsize=(15, 7))
    plt. subplot (1, 4, 1)
    plt.imshow(X[idx_to_reconstruct].reshape((h, w)), cmap=plt.cm.gray)
    plt.title(names[idx to reconstruct])
    plt.grid()
    plt. subplot (1, 4, 2)
    plt. imshow(reconstructed image. reshape((h, w)), cmap=plt. cm. gray)
    plt.title('Full PCA')
    plt.grid()
    plt. subplot (1, 4, 3)
    plt. imshow(reconstructed image rpca. reshape((h, w)), cmap=plt. cm. gray)
    plt.title('Randomized PCA')
    plt.grid()
    plt. subplot (1, 4, 4)
    plt. imshow(reconstructed image kpca. reshape((h, w)), cmap=plt. cm. gray)
    plt.title('Kernel PCA')
    plt.grid()
widgets.interact(plt_reconstruct, idx_to_reconstruct=(0, n_samples-1, 1), __manual=True)
```

#### Out[14]:

```
<function main .plt reconstruct(idx to reconstruct)>
```

#### In [3]:

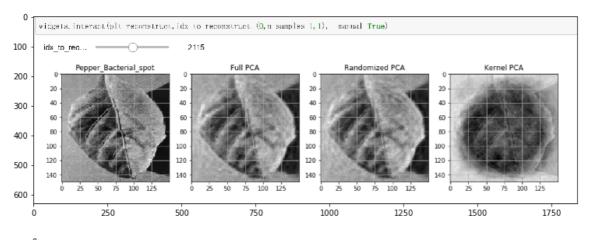
```
#screen shot because widget will not show on other computer
from PIL import Image

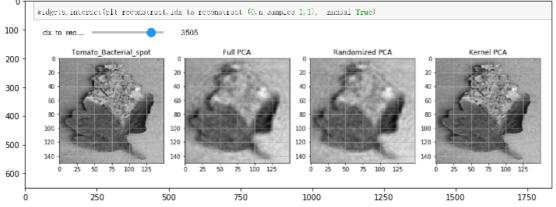
plt.figure(figsize=(20, 15))

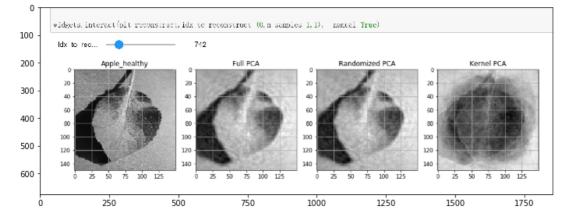
img = Image.open('wigpca-1.png')
plt.subplot(3, 1, 1)
plt.imshow(img)
img1 = Image.open('wigpca-2.png')
plt.subplot(3, 1, 2)
plt.imshow(img1)
plt.subplot(3, 1, 3)
img2 = Image.open('wigpca-3.png')
plt.imshow(img2)
```

# Out[3]:

<matplotlib.image.AxesImage at 0x292506b6358>







After showing all the performances of three PCA method, I found that for the most images, full pca and randomized pca have pretty similar result. For Kernel PCA, sometimes it have the best performance of reflecting the original image. For the most time, kernel pca have similar result comparing to other two method. However, there are some cases that the kernel PCA performance is very bad. Just like professor talked in the class, when kernel pca do well, it does so well. When it does not, it is so bad.

#### In [45]:

```
print ('Full pca accuracy of the classifier is : ', faccuracy)
print ('Randomized pca accuracy of the classifier is : ', raccuracy)
print ('Kernel pcaaccuracy of the classifier is : ', kaccuracy)
```

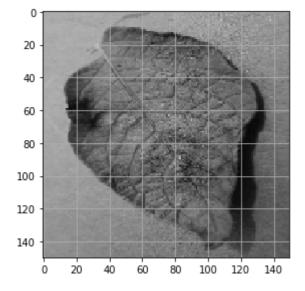
Full pca accuracy of the classifier is : 0.6960529425667691 Randomized pca accuracy of the classifier is : 0.69463483809974 Kernel pcaaccuracy of the classifier is : 0.5398251004490664

After comparing the accuracy for each method, the full pac and randomized have a little bit difference. I personally prefer full pca as the best method for this dataset.

# **DAISY feature extraction**

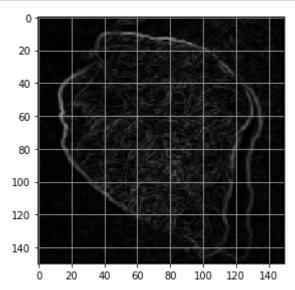
#### In [15]:

```
from skimage.io import imshow
idx_to_reconstruct = 222
idx_to_reconstruct = int(np.random.rand(1)*len(X))
img = X[idx_to_reconstruct].reshape((h, w))
imshow(img)
plt.grid()
```



# In [16]:

```
from skimage.filters import sobel_h, sobel_v
gradient_mag = np.sqrt(sobel_v(img)**2 + sobel_h(img)**2)
imshow(gradient_mag)
plt.grid()
```



#### In [20]:

```
# now let's understand how to use it
from skimage.feature import daisy
features = daisy(img, step=10, radius=10, rings=2, histograms=4, orientations=8, visualize=False
)
print(features.shape)
print(features.shape[0]*features.shape[1]*features.shape[2])
```

(13, 13, 72) 12168

#### In [21]:

```
# create a function to tak in the row of the matric and return a new feature
def apply_daisy(row, shape):
    feat = daisy(row.reshape(shape), step=10, radius=10, rings=2, histograms=6, orientations=8, v
isualize=False)
    return feat.reshape((-1))
```

#### In [22]:

```
# apply to entire data, row by row,
# takes about a minute to run
%time daisy_features = np.apply_along_axis(apply_daisy, 1, X, (h,w))
print(daisy_features.shape)
```

```
Wall time: 1min 48s (4231, 17576)
```

#### In [23]:

```
from sklearn.metrics.pairwise import pairwise_distances
# find the pairwise distance between all the different image features
%time dist_matrix = pairwise_distances(daisy_features)
```

Wall time: 2.02 s

#### In [24]:

```
import copy
from ipywidgets import fixed
# put it together inside a nice widget
def closest_image(dmat, idx1):
    distances = copy.deepcopy(dmat[idx1,:]) # get all image diatances
    distances[idx1] = np.infty # dont pick the same image!
    idx2 = np. argmin(distances)
    distances[idx2] = np.infty
    idx3 = np. argmin (distances)
    plt. figure (figsize=(10, 16))
    plt. subplot (1, 3, 1)
    imshow(X[idx1].reshape((h,w)))
    plt.title("Original Image "+names[idx1])
    plt.grid()
    plt. subplot (1, 3, 2)
    imshow(X[idx2].reshape((h,w)))
    plt.title("Closest Image "+names[idx2])
    plt.grid()
    plt. subplot (1, 3, 3)
    imshow(X[idx3].reshape((h, w)))
    plt.title("Next Closest Image "+names[idx3])
    plt.grid()
widgets.interact(closest_image, idx1=(0, n_samples-1, 1), dmat=fixed(dist_matrix), __manual=True)
```

# Out[24]:

```
<function main .closest image(dmat, idx1)>
```

# In [6]:

```
#screen shot because widget will not show on other computer

from PIL import Image

plt.figure(figsize=(20,15))

img = Image.open('dsy2 (2).png')

plt.subplot(3,1,1)

plt.imshow(img)

img1 = Image.open('dsy2 (1).png')

plt.subplot(3,1,2)

plt.imshow(img1)

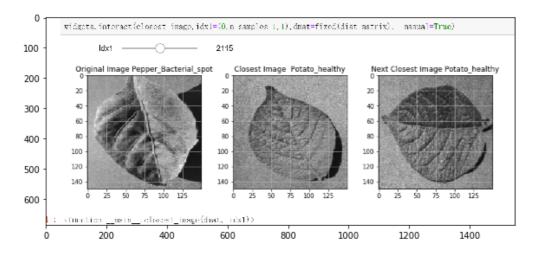
plt.subplot(3,1,3)

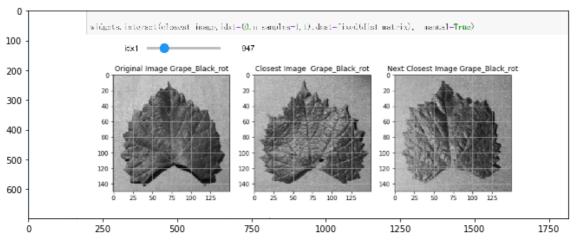
img2 = Image.open('dsy3.png')

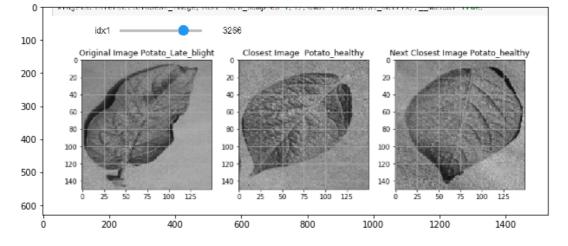
plt.imshow(img2)
```

#### Out[6]:

<matplotlib.image.AxesImage at 0x2924fd6e208>







We can see that for some images, it is pretty accurate. But there are some cloest images are not.

# In [33]:

```
#https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsClassifier.html
from sklearn.neighbors import KNeighborsClassifier
neigh = KNeighborsClassifier(n_neighbors=3, n_jobs=-1)
neigh.fit(daisy_features, names)
%time dpred = neigh.predict(daisy_features)
```

Wall time: 1min 18s

#### In [34]:

```
import numpy as np

daccuracy = 0
for i in range(len(dpred)):
    daccuracy = float(sum(dpred[i] == names[i] for i in range(len(names)))) / len(names)

print('Daisy accuracy of the classifier is: ', daccuracy)
```

Daisy accuracy of the classifier is: 0.5376979437485228

The accuracy for daisy feature extraction is only 54%, which is not good enough. It does not perform well.

# key point matching

# In [7]:

# In [8]:

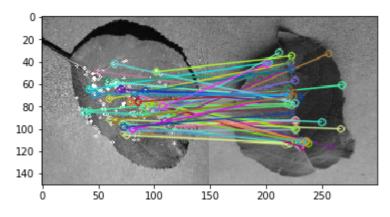
```
#http://answers.opencv.org/question/207758/drawkeypoints-bug/
def key_point(vis, keypoints, color = (255, 0, 0)):
    for i in keypoints:
        x, y = i.pt
        cv2.circle(vis, (int(x), int(y)), 1, color)
```

#### In [9]:

```
#talked to other group about this (mumber:beicheng hu)
#https://docs.opencv.org/3.0-beta/doc/py_tutorials/py_feature2d/py_matcher/py_matcher.html
from skimage.io import imshow
from skimage import img_as_ubyte
import cv2
import numpy as np
from matplotlib import pyplot as pit
orb = cv2.0RB_create()
idx_to_reconstruct = 0
#http://scikit-image.org/docs/dev/api/skimage.html
#Convert an image to unsigned byte format, with values in [0, 255]
img0 = img_as_ubyte(np.array(X[idx_to_reconstruct].reshape((150,150)))) #reshape image, prepare
for analyzing
kp0, des0 = orb.detectAndCompute(img0, None) #keypoints and descriptor
key_point(img0, kp0, color=(255,0,0))
```

#### In [10]:

```
bf = cv2.BFMatcher(cv2.NORM HAMMING, crossCheck=True)
#https://www.programcreek.com/python/example/89342/cv2.drawMatchesKnn
X \text{ kp2} = []
X_{des2} = []
for j in range(len(keyimg)):
    key point (img0, kp0, color=(255, 0, 0))
    img_j = keyimg[j] #prepare the right side of pair
    img_j = img_as_ubyte(img_j)
    kp2, des2 = orb. detectAndCompute(img j, None) #keypoints and descriptor
    X kp2. append (kp2) #appending all keypoints
    X des2. append (des2) #appending all descritor
    matches = bf. match(des0, X_des2[j])
    #https://stackoverflow.com/questions/8966538/syntax-behind-sortedkey-lambda
    matches = sorted(matches, key = lambda x:x.distance) #sort the matches by distance
#https://docs.opencv.org/3.0-beta/modules/features2d/doc/drawing function of keypoints and match
es. html
    imgs = cv2. drawMatches(img0, kp0, img_j, X_kp2[j], matches, img_j, flags=2) #draw the pair keymat
ching plot
    #plt. imshow(imgs), plt. show()
                                   will be 400 images for 400 pairs if running here
plt.imshow(imgs), plt.show() #only show the best match
```

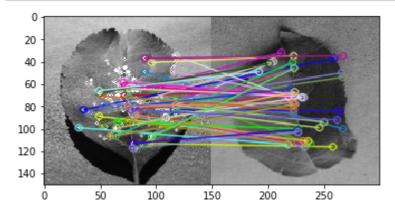


#### Out[10]:

(<matplotlib.image.AxesImage at 0x284ff229d68>, None)

# In [11]:

```
#show the second image with other pairs
orb = cv2. ORB create()
idx to reconstruct = 1
img0 = img as ubyte(np.array(X[idx to reconstruct].reshape((150,150))))
kp0, des0 = orb. detectAndCompute(img0, None)
key_point(img0, kp0, color=(255, 0, 0))
bf = cv2.BFMatcher(cv2.NORM_HAMMING, crossCheck=True)
X \text{ kp2} = []
X des2 = []
#https://docs.opencv.org/3.0-beta/doc/py tutorials/py feature2d/py matcher/py matcher.html
#https://www.programcreek.com/python/example/89342/cv2.drawMatchesKnn
for j in range (len (keyimg)):
    key_point(img0, kp0, color=(255, 0, 0))
    img j = keyimg[j]
    img_j = img_as_ubyte(img j)
    kp2, des2 = orb. detectAndCompute(img j, None)
    X_kp2. append (kp2) #appending all keypoints
    X des2. append (des2)
    matches = bf.match(des0, X_des2[j])
    #https://stackoverflow.com/questions/8966538/syntax-behind-sortedkey-lambda
    matches = sorted(matches, key = lambda x:x.distance) #sort the matches by distance
#https://docs.opencv.org/3.0-beta/modules/features2d/doc/drawing_function_of_keypoints_and_match
    imgs = cv2. drawMatches(img0, kp0, img_j, X_kp2[j], matches, img_j, flags=2)
plt. imshow(imgs), plt. show() #only show the best match
```

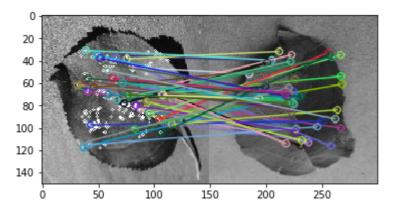


#### Out[11]:

(<matplotlib.image.AxesImage at 0x284ff28f6d8>, None)

#### In [12]:

```
#show the third image with other pairs
orb = cv2. ORB create()
idx to reconstruct = 2
img0 = img as ubyte(np.array(X[idx to reconstruct].reshape((150,150))))
kp0, des0 = orb. detectAndCompute(img0, None)
key_point(img0, kp0, color=(255, 0, 0))
bf = cv2.BFMatcher(cv2.NORM_HAMMING, crossCheck=True)
X \text{ kp2} = []
X des2 = []
#https://docs.opencv.org/3.0-beta/doc/py tutorials/py feature2d/py matcher/py matcher.html
#https://www.programcreek.com/python/example/89342/cv2.drawMatchesKnn
for j in range (len (keyimg)):
    key_point(img0, kp0, color=(255, 0, 0))
    img j = keyimg[j]
    img_j = img_as_ubyte(img j)
    kp2, des2 = orb. detectAndCompute(img j, None)
    X_kp2. append (kp2) #appending all keypoints
    X des2. append (des2)
    matches = bf.match(des0, X_des2[j])
    #https://stackoverflow.com/questions/8966538/syntax-behind-sortedkey-lambda
    matches = sorted(matches, key = lambda x:x.distance) #sort the matches by distance
#https://docs.opencv.org/3.0-beta/modules/features2d/doc/drawing_function_of_keypoints_and_match
    imgs = cv2. drawMatches(img0, kp0, img_j, X_kp2[j], matches, img_j, flags=2)
plt. imshow(imgs), plt. show() #only show the best match
```



# Out[12]:

(<matplotlib.image.AxesImage at 0x284ff0ad438>, None)

#### In [ ]: