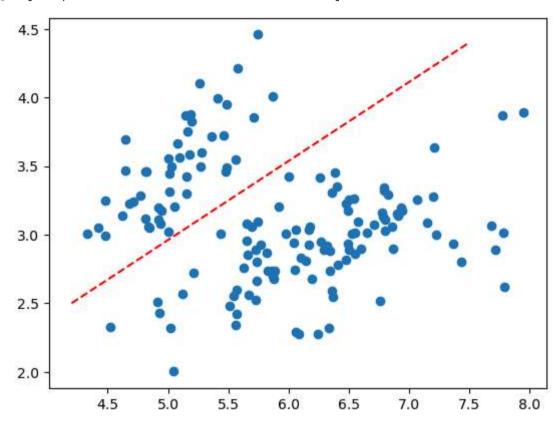
1 Clustering

```
import numpy as np
import matplotlib.pyplot as plt
data = np.loadtxt('Data/iris.txt')
iris = data[:,0:2:1]
plt.scatter(iris[:,0],iris[:,1])
plt.plot([4.2,7.5],[2.5,4.4],"r",linestyle='--')
```

Out[1]: [<matplotlib.lines.Line2D at 0x1490f84a4d0>]



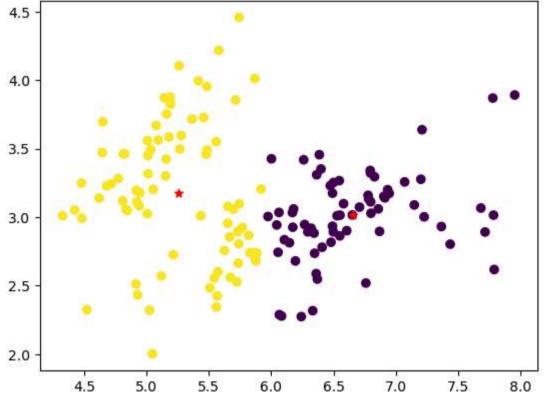
I think there are 2 clusters here, and I draw a red dot line to split them.

```
In [2]: import mltools.plot as plot
import mltools as ml

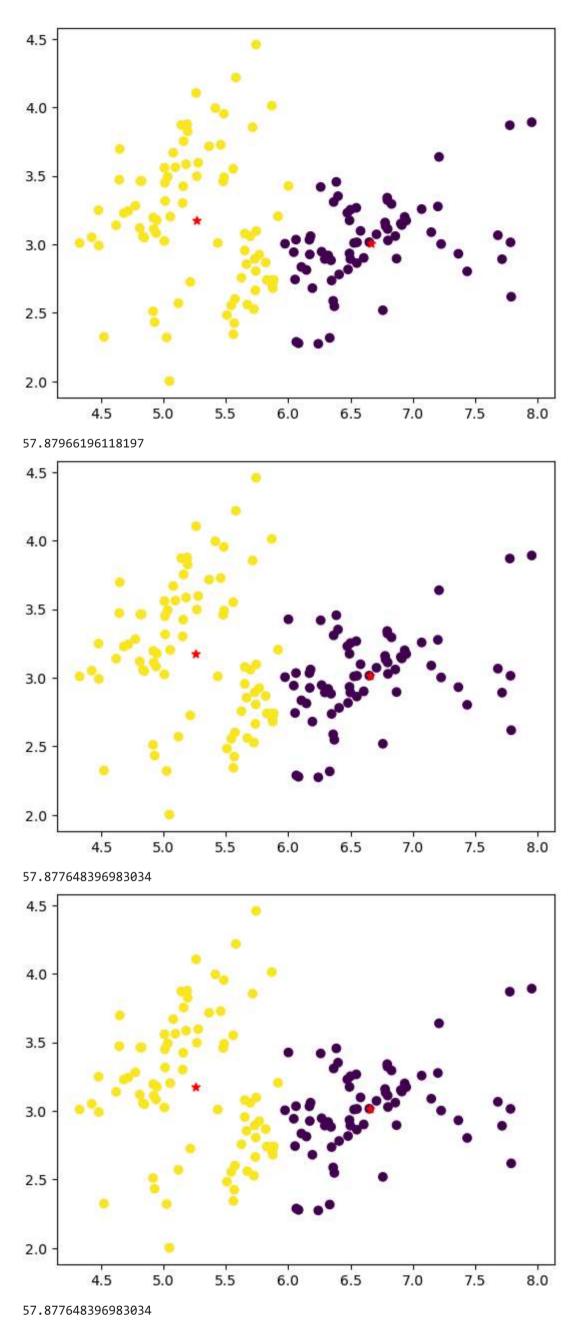
In [48]:

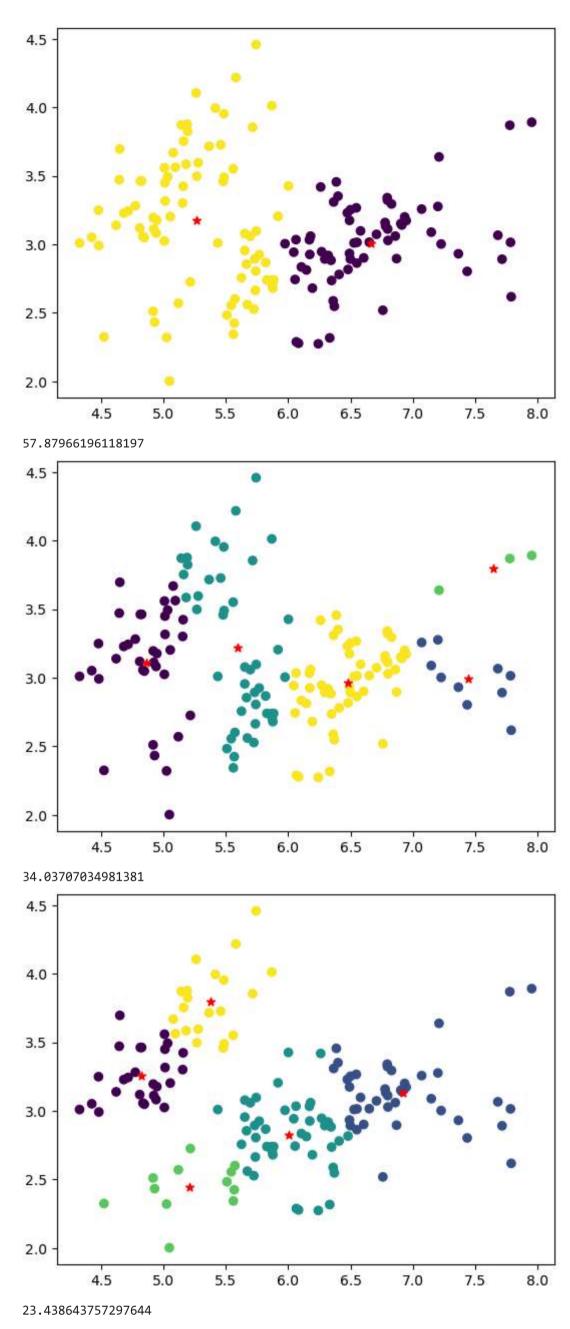
def test_K_of_model(k):
    for i in range(5):
        z,c,s = ml.cluster.kmeans(iris,K=k)
        plt.scatter(iris[:,0],iris[:,1],c=z)
        plt.scatter(c[:,0],c[:,1],marker='*',color ="red")
        plt.show()
        print(s)
```

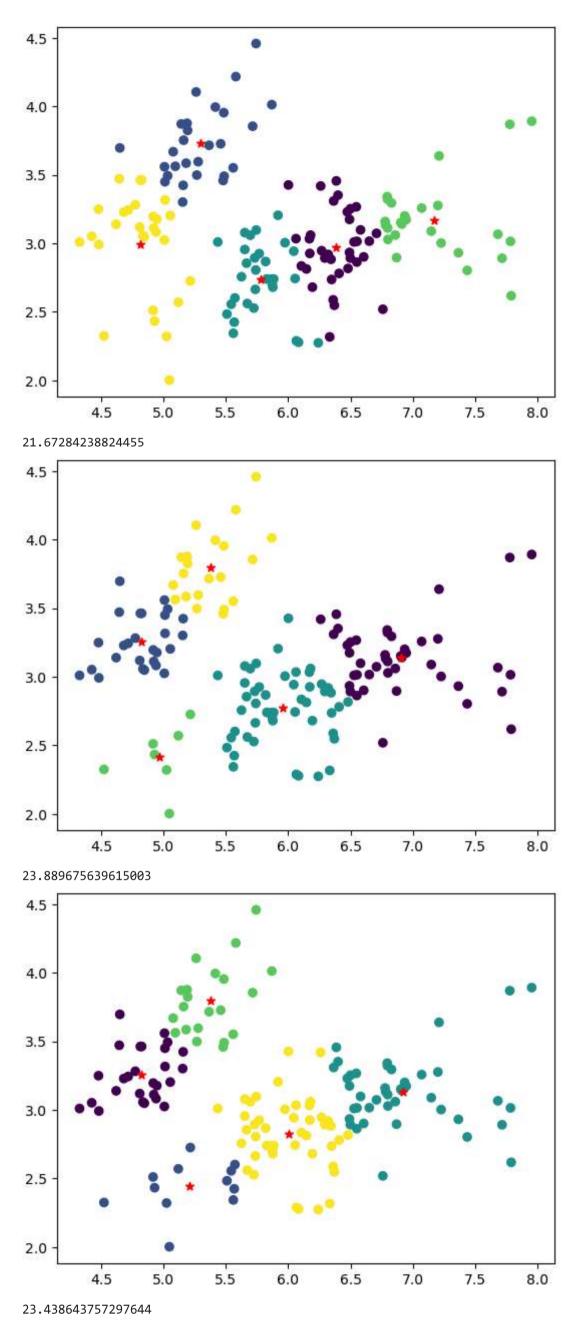
In [49]: for k in [2,5,20]:
 test_K_of_model(k)

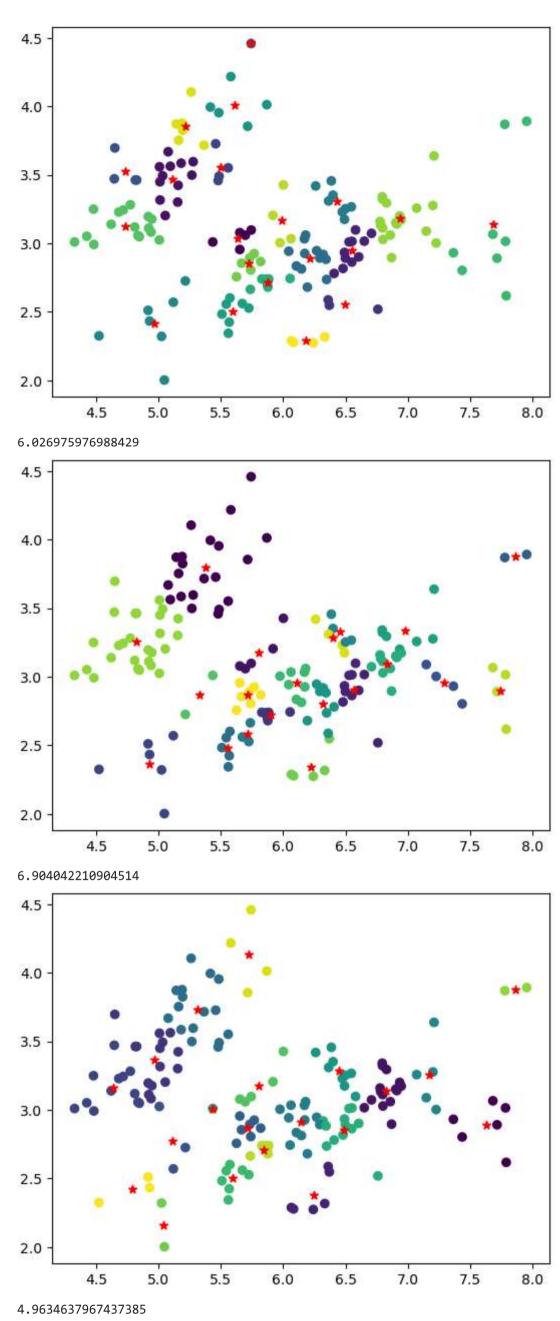


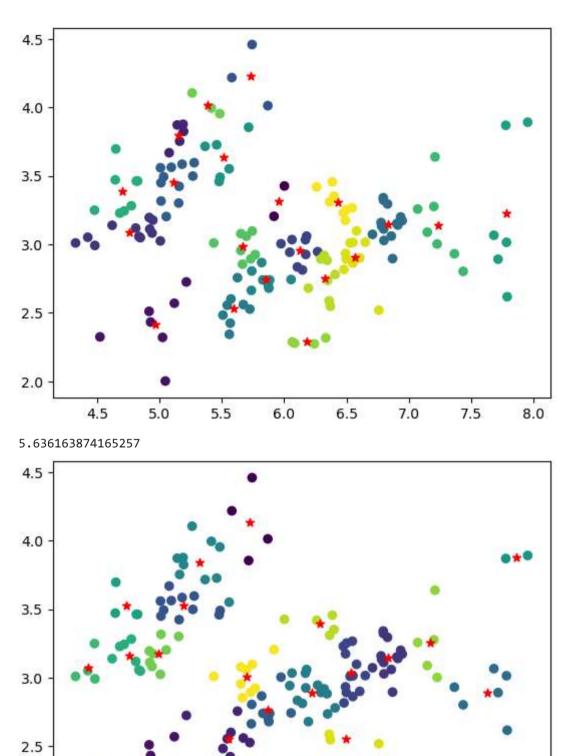
57.877648396983034











4.26953908350041

4.5

2.0

```
In [4]: def agglomerative_with_K(k):
    z,j=ml.cluster.agglomerative(iris,k,"max")

plt.title("Complete Link (max distance) of {} clusters".format(k))
    plot.plotClassify2D(None,iris,z)
    #plt.scatter(iris[:,0],iris[:,1],c=z,s = z)
    plt.show()
    z,j=ml.cluster.agglomerative(iris,k,"min")
    plt.title("Single Link (min distance) of {} clusters".format(k))
    #plt.scatter(iris[:,0],iris[:,1],c=z,s = z)
    plot.plotClassify2D(None,iris,z)

plt.show()
```

7.5

8.0

7.0

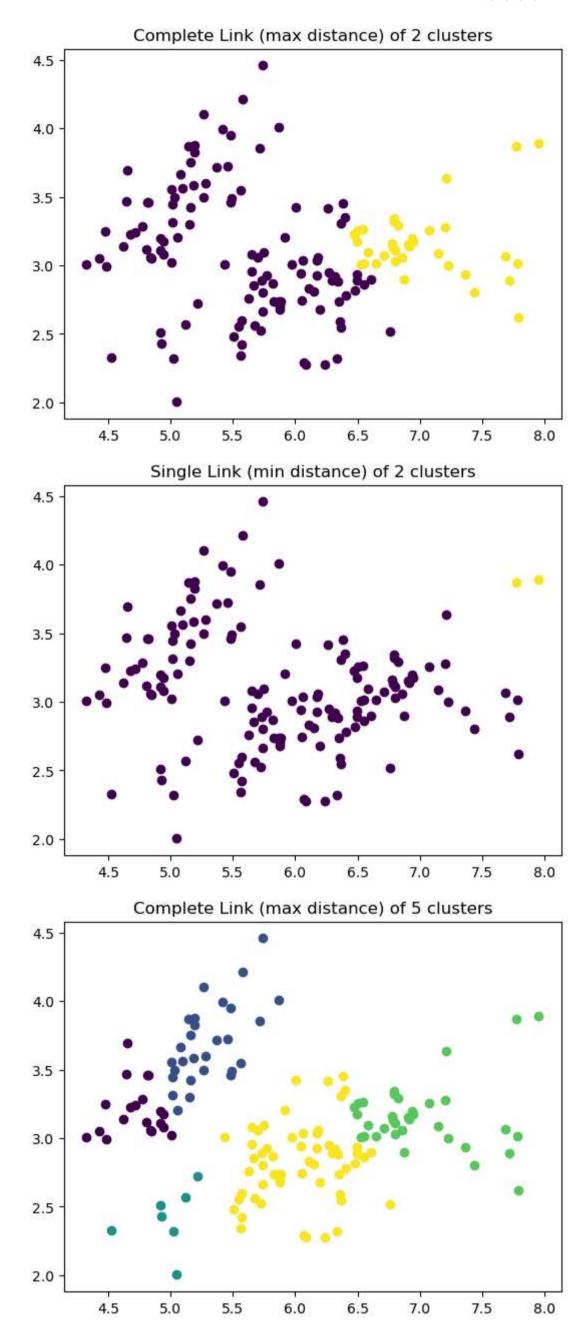
6.5

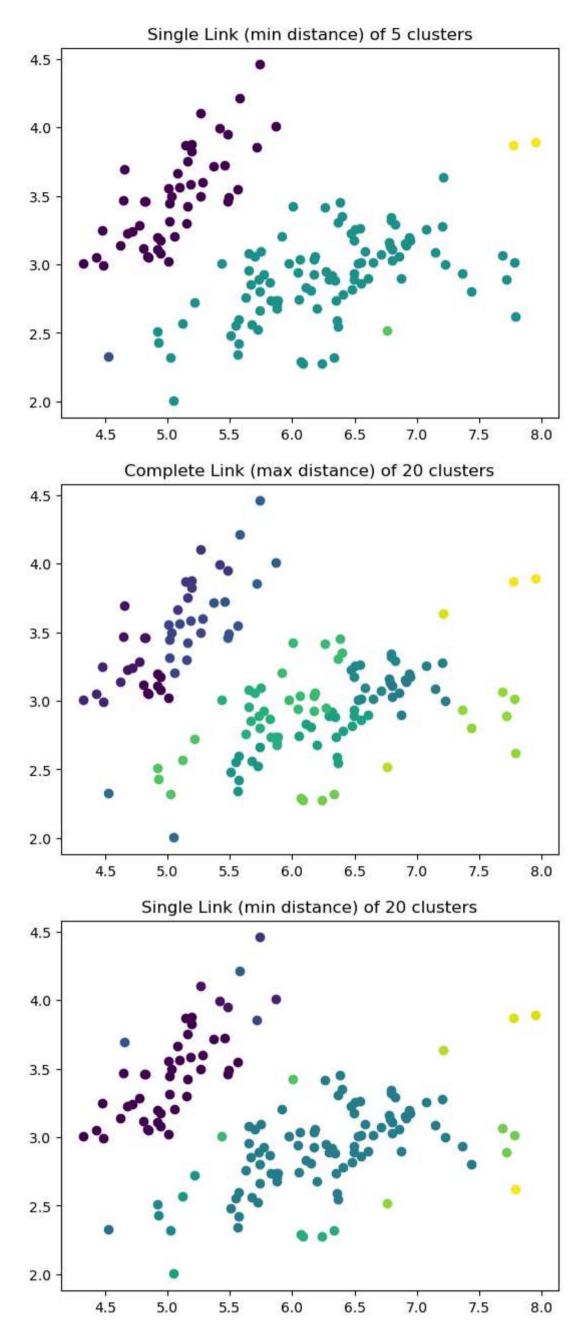
6.0

5.5

In [5]: for k in [2,5,20]:
 agglomerative_with_K(k)

5.0



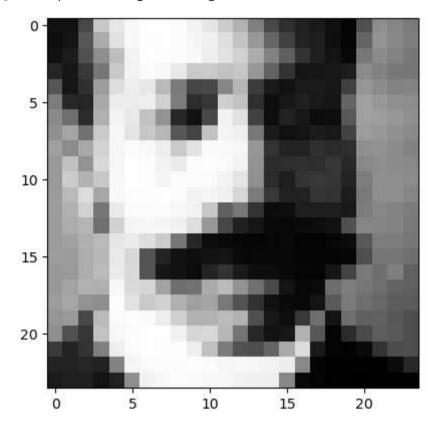


I think the similarity of these two clustering models are that the performance gets better as the number of K improved. But we can hardly find same solution in the two models. In Kmeans, the results is stable when number of cluster is low, and agglomerative models can easily be influenced by outliners. But when the number of cluster improves, the agglomerative models outperforms the Kmeans models.

2 EigenFaces

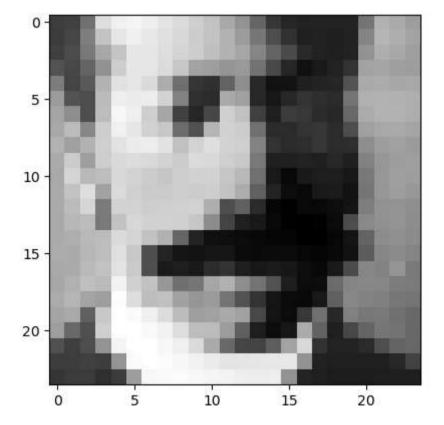
```
import numpy as np
import matplotlib.pyplot as plt
X = np.genfromtxt("data/faces.txt", delimiter=None) # load face dataset
plt.figure()
i = 1
img = np.reshape(X[i,:],(24,24))
plt.imshow( img.T , cmap="gray")
```

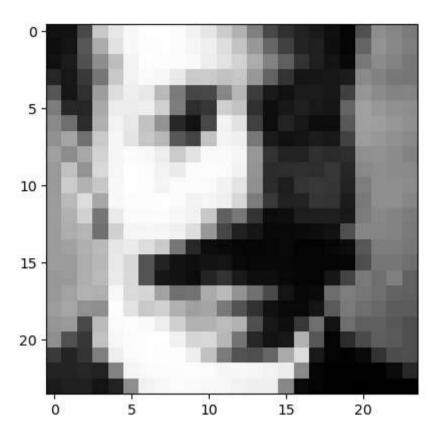
Out[2]: <matplotlib.image.AxesImage at 0x14911a9fdc0>



```
In [6]: mu = np.mean(X, axis=0)
X0 = X - mu
img = np.reshape(X0[i,:],(24,24))
plt.imshow( img.T , cmap="gray")
plt.figure()
img2 = np.reshape(X[i,:],(24,24))
plt.imshow( img2.T , cmap="gray")
```

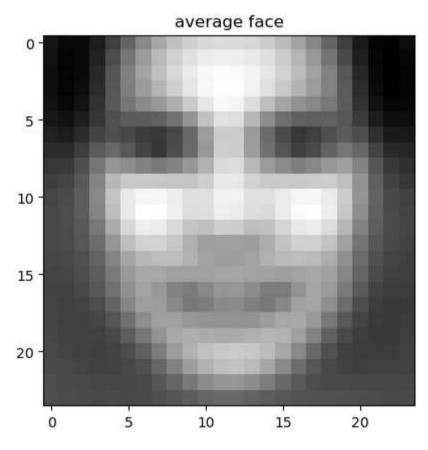
Out[6]: <matplotlib.image.AxesImage at 0x14923f0fb20>



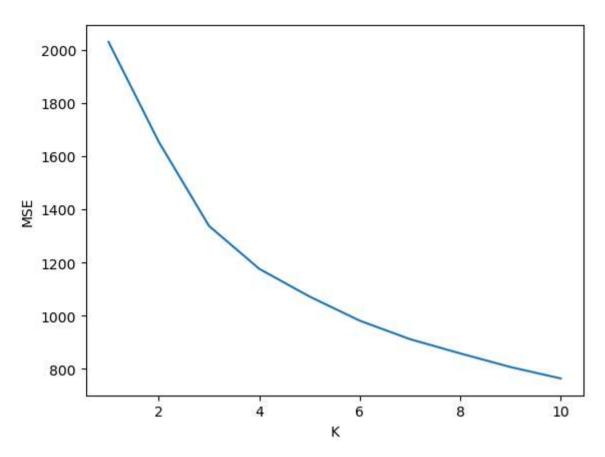


```
In [7]: img = np.reshape(mu,(24,24))
    plt.imshow( img.T , cmap="gray")
    plt.title("average face")
```

Out[7]: Text(0.5, 1.0, 'average face')



```
In [8]: from scipy.linalg import svd
         U, s, V = svd(X0, full_matrices=False)
         W = np.dot(U, np.diag(s))
         print ('Shape of W = (%d, %d)' % W.shape, 'Shape of V = (%d, %d)' % V.shape)
        Shape of W = (4916, 576) Shape of V = (576, 576)
In [24]: def cal_mse_of_k(k):
             mse = np.mean((X0-X0k)**2)
             return mse
In [25]: k_list = []
         Mse = []
         for i in range(1,11):
             k_list.append(i)
             Mse.append(cal_mse_of_k(i))
In [29]: plt.plot(k_list,Mse)
         plt.xlabel("K")
         plt.ylabel("MSE")
Out[29]: Text(0, 0.5, 'MSE')
```



```
In [51]: for j in range(3):
    a = 2 * np.median(np.abs(W[:, j])) # The scalar

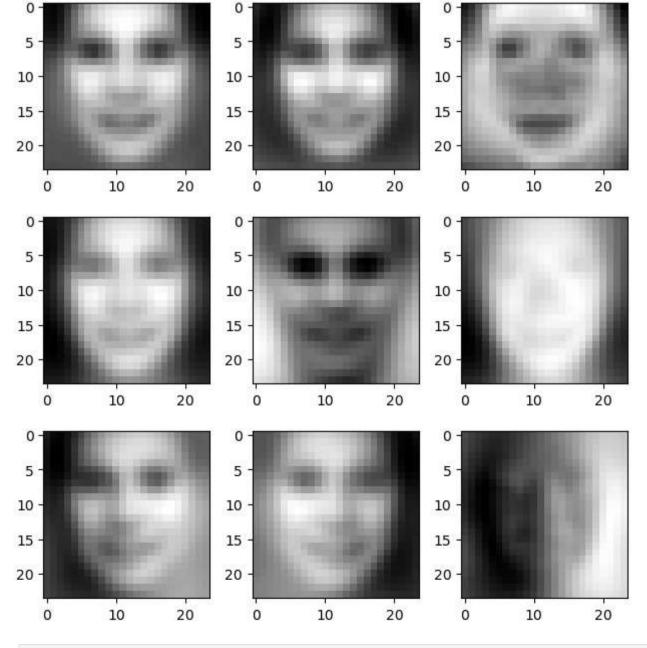
# Compting the "direction" as a function of the mean.
p1 = mu - a * V[j,:]
p2 = mu + a * V[j,:]
fig, (ax1, ax2,ax3) = plt.subplots(1, 3)

img = np.reshape(p1,(24,24))
    ax1.imshow(img.T, cmap="gray")

img2 = np.reshape(p2,(24,24))
ax2.imshow(img2.T, cmap="gray")

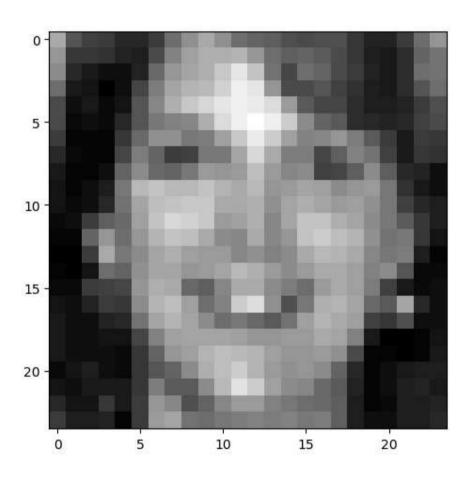
img3 = np.reshape(p1-p2,(24,24))
ax3.imshow(img3.T, cmap="gray")
plt.tight_layout()

plt.show()
```

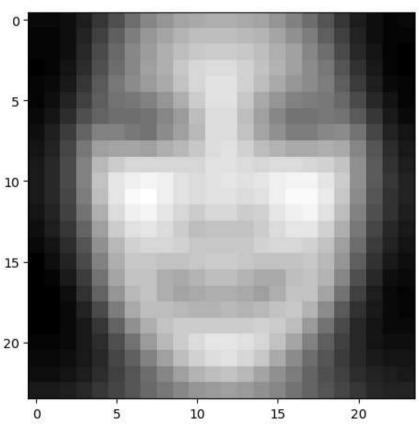


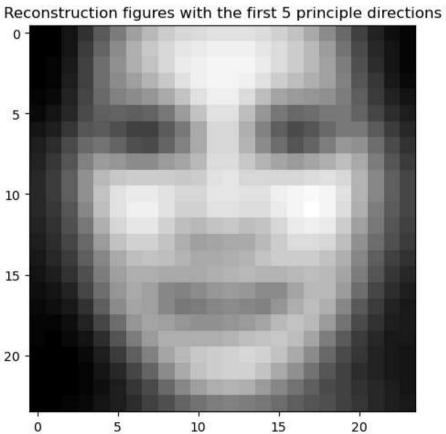
```
In [73]: idx = np.random.choice(X.shape[0], 2, replace=False)
   idx = [88, 14]
```

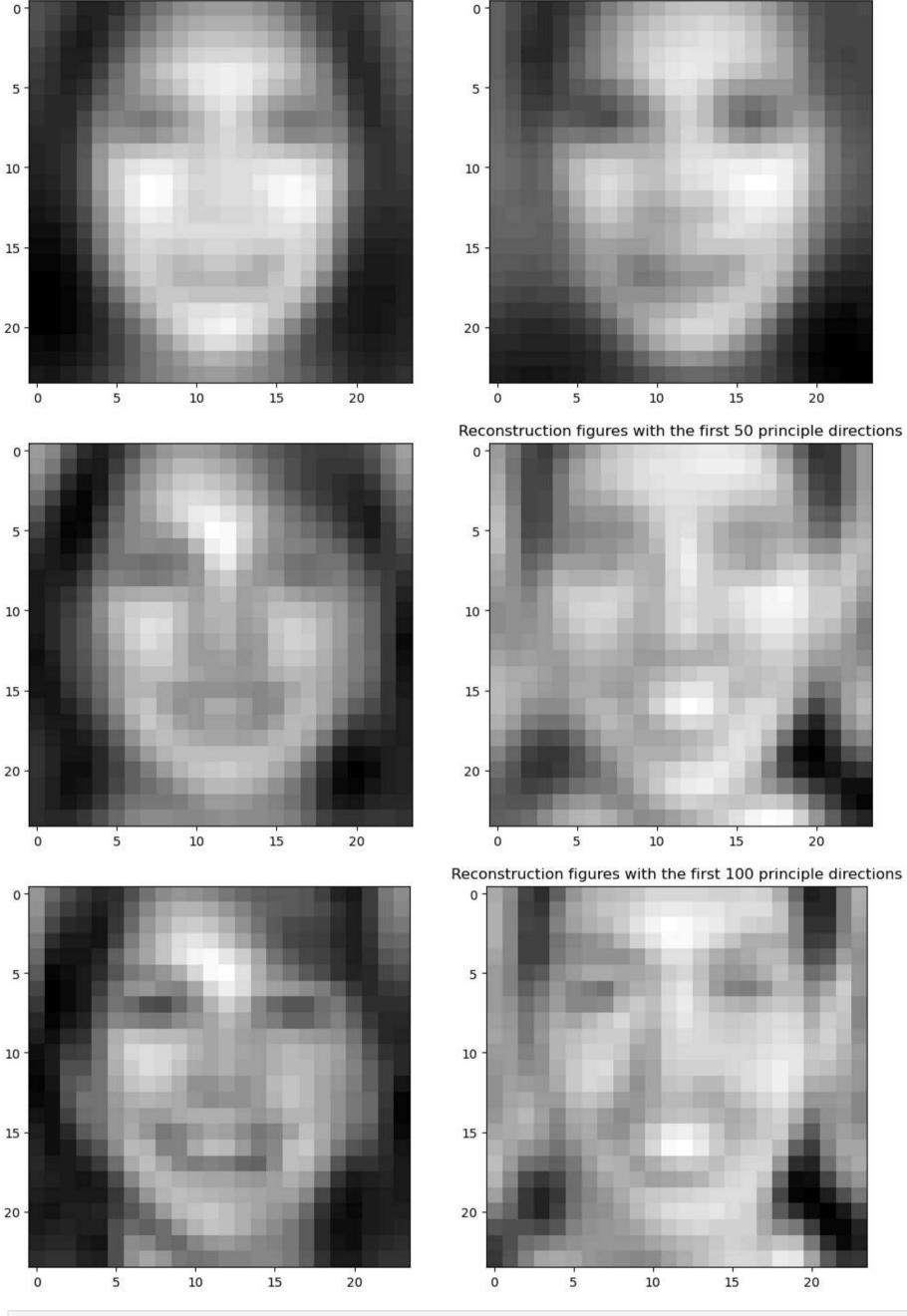
```
k_{list} = [5,10,50,100]
f, ax = plt.subplots(1, 2, figsize=(12, 15))
plt.title("Original Figures")
for j in range(len(idx)):
       i = idx[j]
       img = X[i] # DON'T FORGET TO ADD THE MU
       img = np.reshape(img,(24,24)) # reshape flattened data into a 24*24 patch
       # We've seen the imshow method in the previous discussion :)
       ax[j].imshow( img.T , cmap="gray")
for k in k_list:
    f, ax = plt.subplots(1, 2, figsize=(12, 15))
    plt.title("Reconstruction figures with the first {} principle directions".format(k))
    for j in range(len(idx)):
       i = idx[j]
       img = np.dot(W[i, :k], V[:k]) + mu # DON'T FORGET TO ADD THE MU
       img = np.reshape(img,(24,24)) # reshape flattened data into a 24*24 patch
       # We've seen the imshow method in the previous discussion :)
        ax[j].imshow( img.T , cmap="gray")
```











Reconstruction figures with the first 10 principle directions

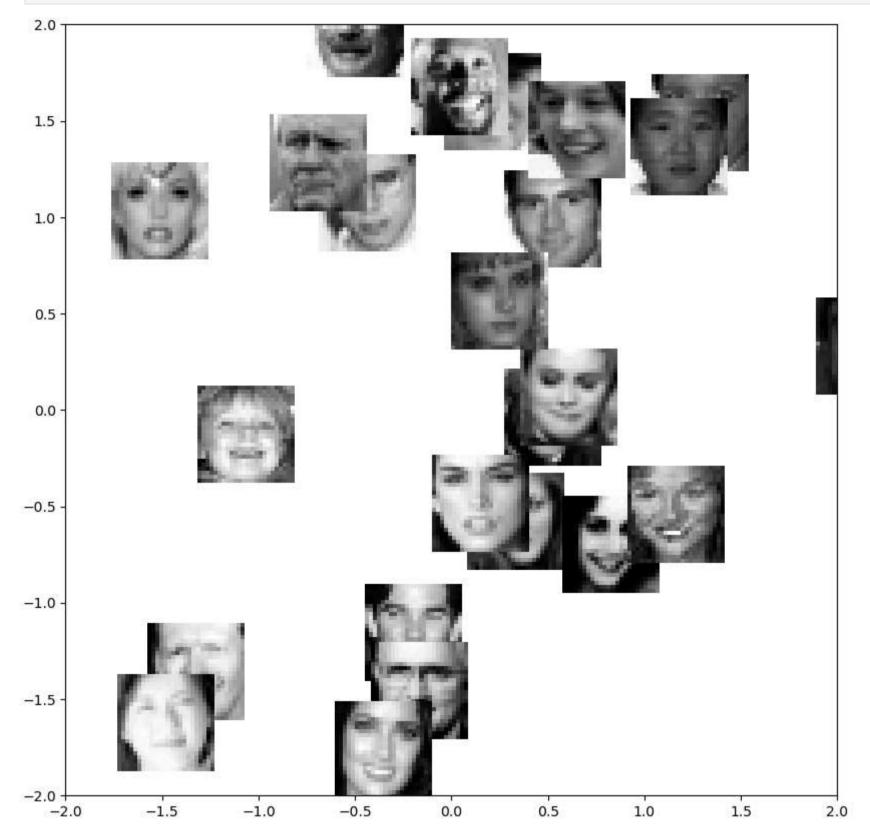
```
In [76]: import mltools as ml
f, ax = plt.subplots(1, 1, figsize=(15, 10))

idx = np.random.choice(X.shape[0], 25, replace=False)

coord, params = ml.transforms.rescale(W[:,0:2])
for i in idx:
    loc = (coord[i, 0], coord[i, 0] + 0.5, coord[i, 1], coord[i, 1] + 0.5)
```

```
img = np.reshape(X[i,:], (24,24))
    ax.imshow(img.T , cmap="gray", extent=loc)

ax.axis([-2, 2, -2, 2])
plt.show()
```



Statement of Collaboration
I finished my homework by myself.

In []: