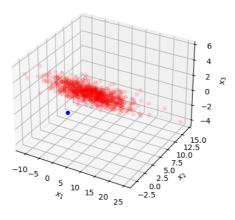
2024/3/22 晚上11:23 Assign6Starter

Figure



2a) Does the data appear to lie in a low-dimensional subspace? Why or why not? Remember the definition of a subspace.

No, it does not lie in a low-dimensional subspace, because the line does not go through the origin.

2b) What could you do to the data so that it lies (approximately) in a low-dimensional subspace?

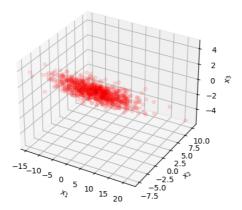
I couldsubsract the data by the meann to mak a low-dimensional line goes through the origin, making it a subspacee

```
In [3]: # Subtract mean
X_m = X - np.mean(X, 0)
In [4]: # display zero mean scatter plot
fig = plt.figure()
    ax = fig.add_subplot(111, projection='3d')
    ax.scatter(X_m[:,0], X_m[:,1], X_m[:,2], c='r', marker='o', alpha=0.1)

ax.scatter(0,0,0,c='b', marker='o')
    ax.set_xlabel('$x_1$')
    ax.set_ylabel('$x_2$')
    ax.set_zlabel('$x_3$')

plt.show()
```

Figure

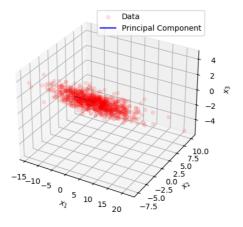


2c) The third section of the code removes the mean (average) value of the 1000 data points. Use the rotate tool to inspect the scatterplot of the data with the mean removed. Does the mean-removed data appear to lie in a low-dimensional sub-space

Yes, the mean-removed data appears to lie in a low-dimensional sub-space, becauseit goes through the origin.?

```
In [5]: # Use SVD to find first principal component
        U,s,VT = np.linalg.svd(X_m,full_matrices=False)
        # complete the next line of code to assign the first principal component to a
        # a = VT[:1,:] # a matrix
In [6]: a.shape
Out[6]: (3,)
In [7]: # display zero mean scatter plot and first principal component
        fig = plt.figure()
        ax = fig.add_subplot(111, projection='3d')
        #scale length of line by root mean square of data for display
        ss = s[0]/np.sqrt(np.shape(X_m)[0])
        ax.scatter(X_m[:,0],\ X_m[:,1],\ X_m[:,2],\ c='r',\ marker='o',\ label='Data',\ alpha=0.1)
        ax.plot([0,ss*a[0]],[0,ss*a[1]],[0,ss*a[2]], c='b',label='Principal Component')
        ax.set_xlabel('$x_1$')
        ax.set_ylabel('$x_2$')
ax.set_zlabel('$x_3$')
        ax.legend()
```

Figure



2d) comment on how well a one-dimensional subspace captures the data

It captures the most important distribution direction of the dat.a

2e) Let xzi, i = 1, 2, . . . , 1000 be the individual mean-removed data points and a the unit-norm vector representing the best one-dimensional subspace for the data. Thus, xzi ≈ awi. Find wi in terms of the SVD matrices U, S, and V.

2024/3/22 晚上11:23 Assign6Starter

```
In [8]: # X_hat = U[:,:1] @ S_matrix[:1,:1] @VT[:1,:]
# a = vt1 --> wi = u1 @ S_matrix[:1,:1]
S_matrix = np.zeros_like(X_m)
np.fill_diagonal(S_matrix, s)
wi = U[:,:1] @ S_matrix[:1,:1]
wi.shape
Out[8]: (1000, 1)
```

2f) Now write the original data xi, i = 1, 2, . . . , 1000 as xi ≈ awi + b. What is b?

b is the mean of X.

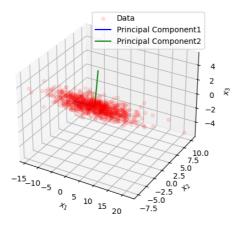
2g) Let E be the difference between X and the rank-one approximation. Find a mathematical expression for ||E||F2 in terms of the singular values of the mean-removed data Xz.

```
In [9]: X_hat = U[:,:1] @ S_matrix[:1,:1] @ VT[:1,:]
E = X_m - X_hat
np.linalg.norm(E,ord='fro')**(2)
```

Out[9]: 626.6899203862777

2h) Now try a rank-two approximation. Use the SVD to find an orthonormal basis for the best plane containing the mean-removed data. Display the mean-removed data and the bases for the plane..

Figure



2i) Your rank-two approximation for the original data is $xi \approx a1w1i + a2w2i + b$, i = 1, 2, ..., 1000. Express w2i, i = 1, 2, ..., 1000 in terms of the SVD of the mean-removed data matrix Xz. Display a scatter plot of the original data (in red) and the rank-two approximations in blue. Does the rank-two approximation lie in a plane? Does that plane capture the dominant components of the data?

Indeed, the rank-2 approximation exists within a plane, and it's evident from the 3D chart that this plane effectively captures the most significant components of the data.

```
In [11]:
    w2i = U[:,1:2] @ S_matrix[1:2,1:2]
    X_hat = U[:,:2] @ S_matrix[2:,:2] @ VT[:2,:]

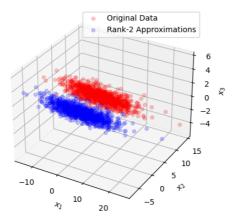
    fig = plt.figure()
    ax = fig.add_subplot(111, projection='3d')

    ax.scatter(X[:,0], X[:,1], X[:,2], c='r', marker='o', label='Original Data', alpha=0.2)
    ax.scatter(X_hat[:,0], X_hat[:,1], X_hat[:,2], c='b', marker='o', label='Rank-2 Approximations', alpha=0.2)

    ax.set_xlabel('$x_1$')
    ax.set_ylabel('$x_2$')
    ax.set_zlabel('$x_2$')
    ax.set_zlabel('$x_3$')

    ax.legend()
    plt.show()
```

Figure



2j) Let E be the difference between X and the rank-two approximation. Find a mathematical expression for ||E||F2 in terms of the singular values of the mean-removed data Xz.

2k) Find and compare the numerical values for ||E||F2 using both the rank-1 and rank-2 approximation.

The approximation error for rank-1 is approximately 626.69, whereas for rank-2, it is about 152.95. This significant reduction in error with rank-2 demonstrates that it captures more essential features of the data compared to the rank-1 model.

3a) Truncated SVD. Use the pseudo-inverse V $\Sigma r-1UT$, where $\Sigma r-1$ is computed by inverting the r largest singular values. Hence the regularization parameter r takes values $r=1,2,\ldots,9$.

```
In [13]: data = loadmat('face_emotion_data.mat')
             X, y = data['X'], data['y']
              total_error = []
              for i in range(8):
                   for j in range(8):
    if i == j: continue
                          test_idx_1 = np.arange(i*16, (i+1)*16)
                          test_idx_2 = np.arange(j*16, (j+1)*16)
train_idx = np.setdiff1d(np.arange(128), test_idx_1)
train_idx = np.setdiff1d(train_idx, test_idx_2)
                          X_train, y_train = X[train_idx, :], y[train_idx, :]
X_test_1, y_test_1 = X[test_idx_1, :], y[test_idx_1, :]
X_test_2, y_test_2 = X[test_idx_2, :], y[test_idx_2, :]
                          param = 0
                          lowest = 1
                          \label{eq:continuity} \begin{array}{ll} \mbox{for r in range(1,10):} \\ \mbox{U, s, VT = np.linalg.svd}(\mbox{X\_train, full\_matrices=False}) \end{array}
                               S = np.diag(1/ s[:r])
approx = VT[:r, :].T @ S @ U[:,:r].T @ y_train
                                y pred = np.sign(X test 2 @ approx)
                                for idx, item in enumerate(y_pred):
    if item == y_test_2[idx]:
        error.append(0)
                                      else:
                                           error.append(1)
                               error_percentage = sum(error) / len(error)
# print(error_percentage)
                                if error_percentage < lowest:</pre>
                                     lowest = error_percentage
param = r
                          U, s, VT = np.linalg.svd(X_train, full_matrices=False)
                         S = np.diag(1/ s[:param])
w = VT[:param, :].T @ S @ U[:,:param].T @ y_train
                          y_pred_1 = np.sign(X_test_1 @ w)
                          for idx, item in enumerate(y_pred_1):
    if item == y_test_1[idx]:
                                error.append(0)
else:
                                      error.append(1)
                          error_percentage = sum(error) / len(error)
                          total_error.append(error_percentage)
              print("Error SVD: ", sum(total_error)/ 56)
            Error SVD: 0.11160714285714286
```

3b) Ridge Regression

2024/3/22 晚上11:23 Assign6Starter

```
lowest = 1
                               lambda_list = [0, 0.5, 1, 2, 4, 8, 16]
                                if i == j: continue
                               test_idx_1 = np.arange(i*16, (i+1)*16)
test_idx_2 = np.arange(j*16, (j+1)*16)
train_idx = np.setdiff1d(np.arange(128), test_idx_1)
train_idx = np.setdiff1d(train_idx, test_idx_2)
X_train, y_train = X[train_idx, :], y[train_idx, :]
X_test_1, y_test_1 = X[test_idx_1, :], y[test_idx_1, :]
X_test_2, y_test_2 = X[test_idx_2, :], y[test_idx_2, :]
                                for x in lambda_list:
                                     U, s, Yr = np.linalg.svd(X_train, full_matrices=False)
S = s / (s**2 +x)
S = np.diag(S)
                                       w = VT.T @ S @ U.T @y_train
                                        y_pred = np.sign(X_test_2 @ w)
                                        for idx, item in enumerate(y_pred):
    if item == y_test_2[idx]:
                                              error.append(0)
else:
                                       error.append(1)
error_percentage = sum(error) / len(error)
# print(error_percentage)
                                       if error_percentage < lowest:
    lowest = error_percentage</pre>
                               U, s, VT = np.linalg.svd(X_train, full_matrices=False)
S = s / (s**2 + param)
S = np.diag(S)
# print(S)
                               new_w = VT.T @ S @ U.T @y_train
y_pred_1 = np.sign(X_test_1 @ new_w)
                               error = []
for idx, item in enumerate(y_pred_1):
    if item == y_test_1[idx]:
        error.append(0)
                                       else:
error.append(1)
                                error_percentage = sum(error) / len(error)
total_error.append(error_percentage)
                 print("Error SVD - Ridge: ", sum(total_error)/ 56)
```

Error SVD - Ridge: 0.04799107142857143

△ Assignment 6

A: nxn, rank 1 mantrix: VE, k= 1211, h power method converges to vs (within 1%) # iterations

Initial vector bo = 1 []
A is symmetric

A = 61 V1 U1 = 61 U1 VI

for iteration =

b1= 61 VI UT bo 11 61 41 41/11/2

= GIMIPO.NI 1614760/11/11/12

Ulis orthonormal [1 /1 1/2

11 dall= 12/11/21/2

> M