

hw2

October 15, 2022

```
[1]: import scipy.io as sio
import numpy as np
import numpy.linalg as la
import matplotlib.pyplot as plt
```

1 1.

1.1 a.

At most there are three linearly independent columns, the set could be made up of any 3 of the four columns.

1.2 b.

The matrix's rank is 3, we know this because it has three linearly independent columns.

1.3 c.

$$\begin{bmatrix} 1 & 0 & 0 & 1 & 0 \\ 1 & 1 & 0 & 0 & 0 \\ 0 & 1 & 1 & 0 & 0 \\ 0 & 0 & 1 & 1 & 0 \end{bmatrix} \begin{bmatrix} 1 & 1 & 0 & 0 \\ 0 & 1 & 1 & 0 \\ 0 & 0 & 1 & 1 \\ 1 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 \end{bmatrix} = \begin{bmatrix} 2 & 1 & 0 & 1 \\ 1 & 2 & 1 & 0 \\ 0 & 1 & 2 & 1 \\ 1 & 0 & 1 & 2 \end{bmatrix}$$

The rank of this matrix is 3 as well (it also has 3 linearly independent columns)

2 2.

2.1 a.

Yes, there are two linearly independent columns. If the columns are x_1, x_2 there is no a and b such that $ax_1 + bx_2 = 0$ except if both a and b are 0.

2.2 b.

Yes, there are three linearly independent columns. If the columns are x_1, x_2, x_3 there is no a, b , and c such that $ax_1 + bx_2 + cx_3 = 0$ except if a, b , and c are 0.

2.3 c.

No, there are two linearly independent columns. If the columns are x_1, x_2, x_3 then $3x_2 - x_1 = x_3$ or in other words $-x_1 + 3x_2 - x_3 = 0$

2.4 d.

The rank of the matrix is 2 since there are two linearly independent columns. Since there are only 2 columns the rank can be at most 2.

3 3.

3.1 a.

$$w^T(3x) = \begin{bmatrix} 3x_1w_1 \\ 3x_2w_2 \\ \dots \\ 3x_iw_i \end{bmatrix}$$

and df/dw of that matrix would be:

$$\begin{bmatrix} 3x_1 \\ 3x_2 \\ \dots \\ 3x_i \end{bmatrix}$$

So the gradient would be

$$3x$$

3.2 b.

$$(w - x)^T(w - x) =$$

$$w^T w - x^T w + x^T x - w^T x$$

$$df/dw = 2w - x - x$$

$$= 2w - 2x$$

3.3 c.

$$x^T \begin{bmatrix} 1 & 2 \\ 3 & 4 \end{bmatrix} w =$$

$$\begin{bmatrix} x_1 + 3x_2 & 2x_1 + 4x_2 \end{bmatrix} \begin{bmatrix} w_1 \\ w_2 \end{bmatrix} =$$

$$\begin{bmatrix} w_1(x_1 + 3x_2) \\ w_2(2x_1 + 4x_2) \end{bmatrix}$$

df/dw of that matrix would simply be $\begin{bmatrix} x_1 + 3x_2 \\ 2x_1 + 4x_2 \end{bmatrix}$

3.4 d.

$$w^T \begin{bmatrix} 1 & 1 \\ -1 & 1 \end{bmatrix} w$$

$$df/dw = \begin{bmatrix} 1 & 1 \\ -1 & 1 \end{bmatrix} w + \begin{bmatrix} 1 & -1 \\ 1 & 1 \end{bmatrix} w$$

$$= \begin{bmatrix} 2 & 0 \\ 0 & 2 \end{bmatrix} w$$

$$= 2w$$

3.5 e.

$$w^T \begin{bmatrix} 1 & 3 \\ 5 & 9 \end{bmatrix} w$$

$$df/dw = \begin{bmatrix} 1 & 3 \\ 3 & 9 \end{bmatrix} w + \begin{bmatrix} 1 & 3 \\ 3 & 9 \end{bmatrix} w$$

$$= \begin{bmatrix} 2 & 6 \\ 6 & 18 \end{bmatrix} w$$

4 4.

4.1 a.

```
[4]: matlab_data_file = sio.loadmat('face_emotion_data.mat')
     X = matlab_data_file['X']
     y = matlab_data_file['y']
     n,p = np.shape(X)
```

```
[5]: w = la.inv(X.T@X)@X.T@y
```

```
[6]: print(w)
```

```
[[ 0.94366942]
 [ 0.21373778]
 [ 0.26641775]
 [-0.39221373]
 [-0.00538552]
 [-0.01764687]
 [-0.16632809]
 [-0.0822838 ]
 [-0.16644364]]
```

4.2 b.

If you were given a new face with a new vector to describe its features, lets call it x you could simply use the dot product of x and w to find an estimate of y ie $\tilde{y} = x^T w$. Since 1 indicates

smiling and -1 indicates not smiling. If \tilde{y} is above 0 then the prediction is smiling ie $\hat{y} = 1$, if \tilde{y} is 0 or below then the prediction is not smiling ie $\hat{y} = -1$

4.3 c.

Assuming each feature is normalized then features 1, 3, and 4 would seem to be the most important since those have the largest absolute values. This means that as long as the features are normalized a 1 point change in any of those features would cause a larger change in the prediction of y . If you added 1 to feature 1 then we'd expect our prediction for y to change by .944 vs if you added 1 to feature 9 we'd expect the prediction to decrease by only .166.

4.4 d.

We could just use the 3 most important features removing all features except for 1, 3 and 4 so that X would be $n \times 3$ and our W would be a vector of length 3.

4.5 e.

```
[7]: Xs = np.split(X, 8)
     ys = np.split(y, 8)
```

```
[8]: total_error_9 = 0
     total_error_3 = 0
     for idx in range(len(Xs)):
         training_X = np.concatenate(Xs[:idx] + Xs[idx+1:])
         training_y = np.concatenate(ys[:idx] + ys[idx+1:])
         test_X = Xs[idx]
         test_y = ys[idx]
         w_hat = la.inv(training_X.T@training_X)@training_X.T@training_y
         y_tilde = test_X@w_hat
         y_hat = np.where(y_tilde > 0, 1, -1)
         errors = ((y_hat - test_y)!=0).sum()
         total_error_9 = total_error_9 + errors/16
         w_hat = la.inv(training_X[:, [0,2,3]].T@training_X[:, [0,2,3]])@training_X[:,
         ↪, [0,2,3]].T@training_y
         y_tilde = test_X[:, [0,2,3]]@w_hat
         y_hat = np.where(y_tilde > 0, 1, -1)
         errors = ((y_hat - test_y)!=0).sum()
         total_error_3 = total_error_3 + errors/16

     avg_error_9 = total_error_9/8
     avg_error_3 = total_error_3/8
```

4.6 f.

```
[9]: print(f"The average error using 9 features was {avg_error_9}")
      print(f"The average error using 3 features was {avg_error_3}")
```

The average error using 9 features was 0.046875

The average error using 3 features was 0.078125

5 5.

```
[10]: # load x and y vectors
d = sio.loadmat('polydata.mat')
z = d['x']
y = d['y']
# n = number of data points
# N = number of points to use for interpolation
# z = points where interpolant is evaluated
# p = array to store the values of the interpolated polynomials
n = z.size
N = 100
z_test = np.linspace(np.min(z), np.max(z), N)
p = np.zeros((3 , N) )
for d in [1 , 2 , 3]:
    X = np.zeros((n,d + 1))
    for i in range(n):
        for j in range(d + 1):
            X[i, j] = z[i]**j

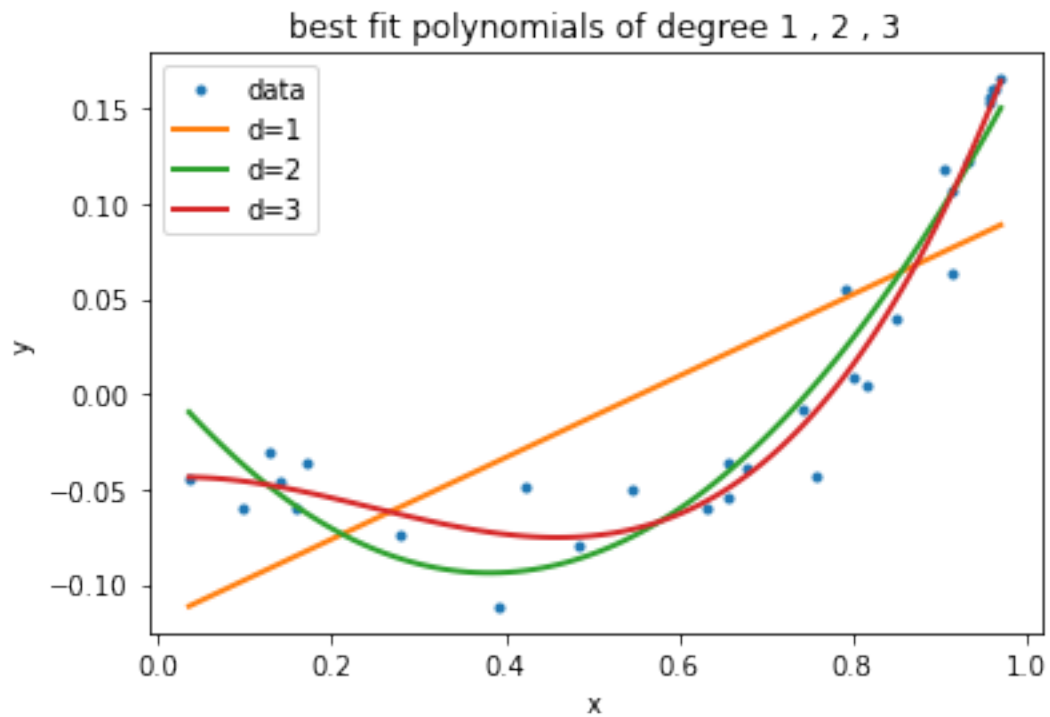
    w = la.inv(X.T@X)@X.T@y

    X_test = np.zeros((N, d+1))
    for i in range(N):
        for j in range(d + 1):
            X_test[i, j] = z_test[i]**j

    p[d-1] = (X_test@w).T

# generate X- matrix for this choice of d
# solve least - squares problem . w is the list
# of polynomial coefficients
# evaluate best -fit polynomial at all points z_test ,
# and store the result in p
# NOTE ( optional ): this can be done in one line
# with the polyval command !
# plot the datapoints and the best -fit polynomials
plt.plot(z, y, '.', z_test, p[0,:], z_test, p[1,:], z_test, p[2,:], linewidth=2)
plt.legend(['data', 'd=1', 'd=2', 'd=3'], loc='upper left')
```

```
plt.title('best fit polynomials of degree 1 , 2 , 3')
plt.xlabel('x')
plt.ylabel('y')
plt.show()
```



[]: