2. The SVD.

$$X_{1} = \begin{bmatrix} 3 \\ 0 \end{bmatrix}, \quad X_{2} = \begin{bmatrix} 1 \\ 2 \end{bmatrix}, \quad X_{3} = \begin{bmatrix} 0 \\ 16 \end{bmatrix}$$
 $X = [x_{1} \ x_{2} \ x_{3}] \in \mathbb{R}^{2 \times 3}$
 $(\alpha_{1}) - \frac{16}{6} \times 1 + \frac{16}{2} \begin{bmatrix} 1 \\ 2 \end{bmatrix} = x_{3}$
 $U_{1} = \frac{x_{1}}{\|x_{1}\|_{2}} = \begin{bmatrix} 3 \\ 0 \end{bmatrix} / 3 = \begin{bmatrix} 1 \\ 0 \end{bmatrix}$
 $X_{2} = [x_{1}] - [x_{2}] = [x_{2}] - [x_{3}] = [x_{3}]$

$$\underline{X}_{3}' = \underline{X}_{3} - \underline{U}_{1}(\underline{U}_{1}^{T} \underline{X}_{2}) = \begin{bmatrix} 1 \\ 2 \end{bmatrix} - \begin{bmatrix} 1 \\ 0 \end{bmatrix} \begin{bmatrix} 1 \\ 0 \end{bmatrix} \begin{bmatrix} 1 \\ 1 \end{bmatrix} \\
= \begin{bmatrix} 1 \\ 2 \end{bmatrix} - \begin{bmatrix} 1 \\ 0 \end{bmatrix} \\
\underline{U}_{3} = \frac{\underline{X}_{3}'}{\|\underline{X}_{2}'\|_{2}} = \begin{bmatrix} 0 \\ 2 \end{bmatrix} / 2 = \begin{bmatrix} 0 \\ 1 \end{bmatrix} \\
\Rightarrow U = \begin{bmatrix} 1 \\ 0 \end{bmatrix}$$

$$0 = \begin{bmatrix} 0 & 1 \end{bmatrix}$$
(b.)
(i) GIR^2

$$P_{\nu} = \nu (\nu^{\mathsf{T}} \nu)^{\mathsf{T}} \nu^{\mathsf{T}}$$

(ii)
$$d_{i}^{2} = \|x_{i} - P_{i}x_{i}\|_{2}^{2} = (x_{i} - P_{i}x_{i})^{T}(x_{i} - P_{i}x_{i})$$

 $= x_{i}^{T}x_{i} - x_{i}^{T}P_{i}^{T}x_{i} - x_{i}^{T}P_{i}x_{i} + x_{i}^{T}P_{i}x_{i}$
 $= \|x_{i}\|_{2}^{2} - 2x_{i}^{T}P_{i}x_{i} + x_{i}^{T}P_{i}x_{i}$
 $= \|x_{i}\|_{2}^{2} - 2x_{i}^{T}P_{i}x_{i} + x_{i}^{T}P_{i}x_{i}$
 $= \|x_{i}\|_{2}^{2} - x_{i}^{T}P_{i}x_{i}$

(iii)

$$d_i^2 = ||x_i||_2^2 - x_i^T P_i x_i$$

$$V = [x], \quad x_i = [x]$$

$$\Rightarrow d_i^2 = \chi_i^T \chi_i - \frac{(\kappa \chi_{i1} + \kappa \chi_{i2})^2}{\kappa^2 + \kappa^2}$$

$$X = \begin{bmatrix} 3 & 1 & 0 \\ 0 & 2 & \sqrt{6} \end{bmatrix}$$

$$\frac{3}{24} (U_1 X_{11} + V_2 X_{12})^2 = (3V_1)^2 + (V_1 + 2V_2)^2 + (76V_2)^2$$

$$= 9V_1^2 + V_1^2 + 4V_1 V_2 + 4V_3^2 + 6V_3^2$$

シルニル ヨンパニーョルニルニ 元ョルが

$$= /oV_1^2 + 4UV_2 + /oV_3^2$$

$$= /o (V_1^2 + V_2^2) + 4V_1 V_2$$
$$= /o + 4V_1 V_2$$

$$v = [x], x_i = [x_i]$$

V. J. T. TA

diz. So vis not unighe.

$$VV^{\mathsf{T}} = V'V'^{\mathsf{T}}$$
.

$$\begin{aligned}
\sigma_{1} &= \| \times^{T} \mathcal{L}_{1} \|_{2} \\
&= \| \begin{bmatrix} 3 & 0 \\ 1 & 2 \\ 0 & 76 \end{bmatrix} \begin{bmatrix} \frac{1}{12} \\ \frac{1}{12} \end{bmatrix} \|_{2} &= \| \begin{bmatrix} \frac{3}{12} \\ \frac{3}{12} \\ \frac{1}{12} \end{bmatrix} \|_{2} &= \sqrt{\frac{9}{2} + \frac{9}{2} + \frac{6}{2}} &= \sqrt{\frac{1}{12}} = 2\sqrt{\frac{3}{12}} \\
&= \sqrt{\frac{9}{2} + \frac{9}{2} + \frac{6}{2}} &= \sqrt{\frac{1}{12}} = 2\sqrt{\frac{3}{12}}
\end{aligned}$$

$$\mathcal{J}_{2} = \| \mathbf{x}^{T} \mathbf{y}_{2} \|_{2} \\
= \| \begin{bmatrix} \mathbf{3} & 0 \\ 1 & 2 \\ 0 & 16 \end{bmatrix} \begin{bmatrix} \mathbf{x}_{1} \\ -\mathbf{x}_{2} \end{bmatrix} \|_{2} = \| \begin{bmatrix} \mathbf{3}_{1} \\ -\mathbf{x}_{2} \\ -\mathbf{x}_{2} \end{bmatrix} \|_{2} = \mathbf{1} \mathbf{9} + \frac{1}{2} + \frac{1}{2} = \mathbf{1} \mathbf{8} = 2 \mathbf{1} \mathbf{2}$$

$$\Rightarrow \Sigma = \begin{bmatrix} 25 & 0 & 0 \\ 0 & 25 & 0 \end{bmatrix}$$

$$X = \sum_{i=1}^{r} \sigma_i u_i v_i^T$$

(b)
$$\chi_{k} = \sum_{i=1}^{k} \sigma_{i} u_{i} v_{i}^{T}$$

$$4. \quad X = \begin{bmatrix} 5 & 0 \\ 0 & 1 \end{bmatrix}$$

X is a diagonal matrix

$$\Rightarrow U = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}, \Sigma = \begin{bmatrix} 5 & 0 \\ 0 & 1 \end{bmatrix}, V = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}$$

X is a diagonal matrix

cmsc253 hw4

November 5, 2023

```
[]: import numpy as np import numpy.linalg as la # matrix rank, inverse import scipy.io import matplotlib.pyplot as plt # plots import scipy.io as sio import sys from matplotlib import colors

Q1 Gram-Schmidt. (a)

[]: def gram_schmidt(X):
    # X is an n-by-p matrix.
    # Returns U an orthonormal matrix.
    # eps is a threshold value to identify if a vector # is nearly a zerous vector.
```

```
[]: def gram_schmidt(X):
         eps = 1e-12
         n, p = X.shape
         U = np.zeros((n, 0))
         for j in range(p):
             # Get the j-th column of matrix X
             v = X[:, j]
             # Write your own code here: Perform the
             # orthogonalization by subtracting the projections on
             # all columns of U. And then check whether the vector
             # you get is nearly a zero vector.
             v = v - U@U.T@v
             v = np.reshape(v, (-1, 1))
             if np.linalg.norm(v) > eps:
                 \# Normalize vector v and append it to U
                 U = np.hstack((U, v / la.norm(v)))
         return U
     X = np.array([[1, 1/2, 1/3, 1/4, 1/5, 1/6, 1/7],
                   [1/2, 1/3, 1/4, 1/5, 1/6, 1/7, 1/8],
                   [1/3, 1/4, 1/5, 1/6, 1/7, 1/8, 1/9],
                   [1/4, 1/5, 1/6, 1/7, 1/8, 1/9, 1/10],
                   [1/5, 1/6, 1/7, 1/8, 1/9, 1/10, 1/11],
                   [1/6, 1/7, 1/8, 1/9, 1/10, 1/11, 1/12],
```

```
gram schmidt(X)
[]: array([[8.13304645e-01, -5.43794290e-01, 1.99095308e-01,
             -5.51132525e-02, 1.19578493e-02, -1.96870730e-03,
             -2.10219152e-04],
            [ 4.06652323e-01, 3.03317262e-01, -6.88560548e-01,
              4.75965265e-01, -1.97392654e-01, 5.41151990e-02,
              3.01288446e-03],
            [ 2.71101548e-01, 3.93949644e-01, -2.07134626e-01,
             -4.90111167e-01, 6.10839674e-01, -3.26893986e-01,
              1.54577469e-02],
            [ 2.03326161e-01, 3.81744394e-01, 1.12389156e-01,
             -4.39598473e-01, -2.54179063e-01, 6.41045574e-01,
             -2.08187091e-01],
            [ 1.62660929e-01, 3.51412668e-01, 2.91518455e-01,
             -1.12276608e-01, -4.99206689e-01, -2.02224426e-01,
              6.15118567e-01],
            [ 1.35550774e-01, 3.20235052e-01, 3.89191824e-01,
              2.30886766e-01, -1.50594721e-01, -5.41817789e-01,
             -7.06123659e-01],
            [ 1.16186378e-01, 2.92095792e-01, 4.40744903e-01,
              5.20623748e-01, 5.01273338e-01, 3.80535581e-01,
              2.81830797e-01]])
     (b)
[]: def hilbert matrix(n):
         X = \text{np.array}([[1.0 / (i + j - 1) \text{ for } i \text{ in } )
                        range(1, n + 1)] for j in range(1, n + 1)])
         return X
     n = 7
     X = hilbert_matrix(n)
     U = gram_schmidt(X)
     L1 = np.identity((U.T@U).shape[0]) - U.T@U
     print(L1)
     err = la.norm(L1, ord=1)
     print(f'L1 matrix norm is {err}')
    [[ 0.00000000e+00 6.73072709e-16 -2.36616282e-15 2.80331314e-14
      -1.92849903e-12 1.25247715e-10 -9.71088800e-09]
     [ 6.73072709e-16  0.00000000e+00 -2.76723089e-14  3.71008779e-13
      -2.63988831e-12 -4.05814549e-11 5.49175827e-09]
     [-2.36616282e-15 -2.76723089e-14 1.11022302e-16 8.52865001e-12
      -2.31620473e-10 5.20528393e-09 -1.33284293e-07]
     [ 2.80331314e-14 3.71008779e-13 8.52865001e-12 -2.22044605e-16
      -9.04092928e-09 4.29613409e-07 -1.80981921e-05]
     [-1.92849903e-12 -2.63988831e-12 -2.31620473e-10 -9.04092928e-09
```

[1/7, 1/8, 1/9, 1/10, 1/11, 1/12, 1/13]])

```
[ 1.25247715e-10 -4.05814549e-11 5.20528393e-09 4.29613409e-07
       2.66371580e-05 0.00000000e+00 -2.27098006e-01]
     [-9.71088800e-09 5.49175827e-09 -1.33284293e-07 -1.80981921e-05
      -2.30322707e-03 -2.27098006e-01 1.11022302e-16]]
    L1 matrix norm is 0.22941947980869065
[]: def modified gram schmidt(X):
         \# Define a threshold value to identify if a vector \# is nearly a zero
      ⇒vector.
        eps = 1e-12
        n, p = X.shape
        U = np.zeros((n, 0))
        for j in range(p):
             # Get the j-th column of matrix X
            v = X[:, j]
            for i in range(j):
                 # Compute and subtract the projection of
                 # vector v onto the i-th column of U
                 v = v - np.dot(U[:, i], v) * U[:, i]
            v = np.reshape(v, (-1, 1))
             # Check whether the vector we get is nearly # a zero vector
             if np.linalg.norm(v) > eps:
                 # Normalize vector v and append it to U
                 U = np.hstack((U, v / la.norm(v)))
        return U
     n = 7
     X = hilbert_matrix(n)
     U_modified = modified_gram_schmidt(X)
     L1_modified = np.identity((U_modified.T@U_modified).shape[0]) - \
        U_modified.T@U_modified
     L1 modified
     print(L1 modified)
     err_modified = la.norm(L1_modified, ord=1)
     print(f'Modified L1 matrix norm is {err_modified}')
    [[ 0.00000000e+00 1.54737334e-15 -2.16146545e-14 3.62931907e-13
      -7.97459321e-12 3.16995839e-10 -1.90386089e-08]
     [ 1.54737334e-15  0.00000000e+00  1.02695630e-15 -2.63955524e-14
       9.00696184e-13 -2.74882339e-11 1.03917761e-09]
     [-2.16146545e-14 1.02695630e-15 -2.22044605e-16 3.05311332e-15
      -2.71893619e-13 1.51737511e-11 -6.93317848e-10]
     [ 3.62931907e-13 -2.63955524e-14 3.05311332e-15 0.00000000e+00
      -1.66533454e-15 -2.34368080e-13 4.83647289e-11]
     [-7.97459321e-12 9.00696184e-13 -2.71893619e-13 -1.66533454e-15
```

0.00000000e+00 2.66371580e-05 -2.30322707e-03]

```
0.00000000e+00 2.01227923e-14 -2.16494878e-12]
[ 3.16995839e-10 -2.74882339e-11 1.51737511e-11 -2.34368080e-13 2.01227923e-14 2.22044605e-16 -1.46549439e-14]
[-1.90386089e-08 1.03917761e-09 -6.93317848e-10 4.83647289e-11 -2.16494878e-12 -1.46549439e-14 -2.22044605e-16]]
Modified L1 matrix norm is 2.0821648873819987e-08
```

In the traditional approach to Gram-Schmidt, each vector in the sequence is made orthogonal to all previously processed vectors by subtracting their projections. The modified Gram-Schmidt re-orthogonalizes each vector against all previously orthogonalized vectors, one at a time. This stepwise approach reduces the accumulation of errors and maintains better orthogonality. Therefore, the modified Gram-Schmidtis process is generally superior to the original process.

Q6 (a)

```
[]: d = np.load('face_emotion_data.npz')
     X = d['X']
     y = d['y']
     n, p = np.shape(X)
     # error rate for truncated SVD
     error_SVD = np.zeros((8, 7))
     # SVD parameters to test
     k_vals = np.arange(9) + 1
     param_err_SVD = np.zeros(len(k_vals))
     subset_count = 8
     subsets = np.array_split(np.arange(n), 8)
     for i in range(subset count):
         X_hold_out = X[subsets[i]]
         y_hold_out = y[subsets[i]]
         for idx, j in enumerate([j for j in range(subset_count) if j != i]):
             X_train = np.concatenate([X[subsets[p]] for p in range(8) if p != i
                                       and p != j])
             y_train = np.concatenate([y[subsets[p]] for p in range(8) if p != i
                                       and p != j])
             U, Sigma, VT = la.svd(X_train, full_matrices=False)
             X test = X[subsets[j]]
             y_test = y[subsets[j]]
             w_hat_lst = list()
             for k_idx, k in enumerate(k_vals):
                 Sigma_plus = np.diag(1 / Sigma[:k])
```

```
VT_k = VT[:k, :]
U_k = U[:, :k]

w_hat = VT_k.T@Sigma_plus@U_k.T@y_train
w_hat_lst.append(w_hat)
y_hat = X_test@w_hat

param_err_SVD[k_idx] += np.sum(np.sign(y_hat) != y_test) / 16

# print(param_err_SVD)
k_best = np.argmin(param_err_SVD)
w_hat_best = w_hat_lst[k_best]

y_hat_hold_out = X_hold_out@w_hat_best
error_SVD[i][idx] = np.sum(np.sign(y_hat_hold_out) != y_hold_out) / 16
print(f"Error_estimate: {np.mean(error_SVD)}")
```

(b)

```
[]: d = np.load('face_emotion_data.npz')
     X = d['X']
     y = d['y']
     n, p = np.shape(X)
     # error rate for regularized least squares
     error RLS = np.zeros((8, 7))
     # RLS parameters to test
     lambda_vals = np.array([0, 0.5, 1, 2, 4, 8, 16])
     param_err_RLS = np.zeros(len(lambda_vals))
     subset_count = 8
     subsets = np.array_split(np.arange(n), 8)
     for i in range(subset_count):
         X_hold_out = X[subsets[i]]
         y_hold_out = y[subsets[i]]
         for idx, j in enumerate([j for j in range(subset_count) if j != i]):
             X_train = np.concatenate([X[subsets[p]] for p in range(8) if p != i
                                       and p != j])
             y_train = np.concatenate([y[subsets[p]] for p in range(8) if p != i
                                       and p != j]
             X_test = X[subsets[j]]
             y_test = y[subsets[j]]
```

(c)

```
[]: X = d['X']
     y = d['y']
     new_features = X@np.random.rand(9,3)
     X_new = np.concatenate((X, new_features), axis = 1)
     ### (a) ' ###
     n, p = np.shape(X_new)
     # error rate for regularized least squares
     error_SVD = np.zeros((8, 7))
     # SVD parameters to test
     k_vals = np.arange(9) + 1
     param_err_SVD = np.zeros(len(k_vals))
     subset_count = 8
     subsets = np.array_split(np.arange(n), 8)
     for i in range(subset_count):
         X_hold_out = X_new[subsets[i]]
         y_hold_out = y[subsets[i]]
         for idx, j in enumerate([j for j in range(subset_count) if j != i]):
             X_train = np.concatenate([X new[subsets[p]] for p in range(8) if p != i
                                       and p != j])
             y_train = np.concatenate([y[subsets[p]] for p in range(8) if p != i
```

```
and p != j])
        U, Sigma, VT = la.svd(X_train, full_matrices=False)
        X_test = X_new[subsets[j]]
        y_test = y[subsets[j]]
        w_hat_lst = list()
        for k_idx, k in enumerate(k_vals):
            Sigma_plus = np.diag(1 / Sigma[:k])
            VT_k = VT[:k, :]
            U_k = U[:, :k]
            w_hat = VT_k.T@Sigma_plus@U_k.T@y_train
            w_hat_lst.append(w_hat)
            y_hat = X_test@w_hat
            param_err_SVD[k_idx] += np.sum(np.sign(y_hat) != y_test) / 16
        # print(param_err_SVD)
        k_best = np.argmin(param_err_SVD)
        w_hat_best = w_hat_lst[k_best]
        y_hat_hold_out = X_hold_out@w_hat_best
        error_SVD[i][idx] = np.sum(np.sign(y_hat_hold_out) != y_hold_out) / 16
print(f"Error estimate: {np.mean(error_SVD)}")
```

```
[]: ### (b)' ###
# error rate for regularized least squares
error_RLS = np.zeros((8, 7))
# RLS parameters to test
lambda_vals = np.array([0, 0.5, 1, 2, 4, 8, 16])
param_err_RLS = np.zeros(len(lambda_vals))

subset_count = 8

subsets = np.array_split(np.arange(n), 8)

for i in range(subset_count):
    X_hold_out = X_new[subsets[i]]
    y_hold_out = y[subsets[i]]

for idx, j in enumerate([j for j in range(subset_count) if j != i]):
    X_train = np.concatenate([X_new[subsets[p]] for p in range(8) if p != i
```

```
and p != j])
        y_train = np.concatenate([y[subsets[p]] for p in range(8) if p != i
                                  and p != j])
        X_test = X_new[subsets[j]]
        y_test = y[subsets[j]]
        w_hat_lst = list()
        for lmd_idx, lmd in enumerate(lambda_vals):
            w_hat = la.inv(X_train.T@X_train + lmd*np.identity(p))@X_train.
 →T@y_train
            w_hat_lst.append(w_hat)
            y_hat = X_test@w_hat
            param_err_RLS[lmd_idx] += np.sum(np.sign(y_hat) != y_test) / 16
        # print(param_err_SVD)
        k_best = np.argmin(param_err_RLS)
        w_hat_best = w_hat_lst[k_best]
        y_hat_hold_out = X_hold_out@w_hat_best
        error_RLS[i][idx] = np.sum(np.sign(y_hat_hold_out) != y_hold_out) / 16
print(f"Error estimate: {np.mean(error_RLS)}")
```

The average error rate of Truncated SVD solution is 0.052455357142857144. The average error rate of Regularized LS is 0.0546875. With new features, Avg error rate of SVD_new is 0.049107142857142856 and Avg error rate of RLS_new is 0.05357142857142857. It seems that these new features are helpful for classification, but not significantly so. One possible explanation is that since the new features are generated as random linear combinations of the original features, they might not necessarily add any meaningful or informative variance to the model that would improve classification.