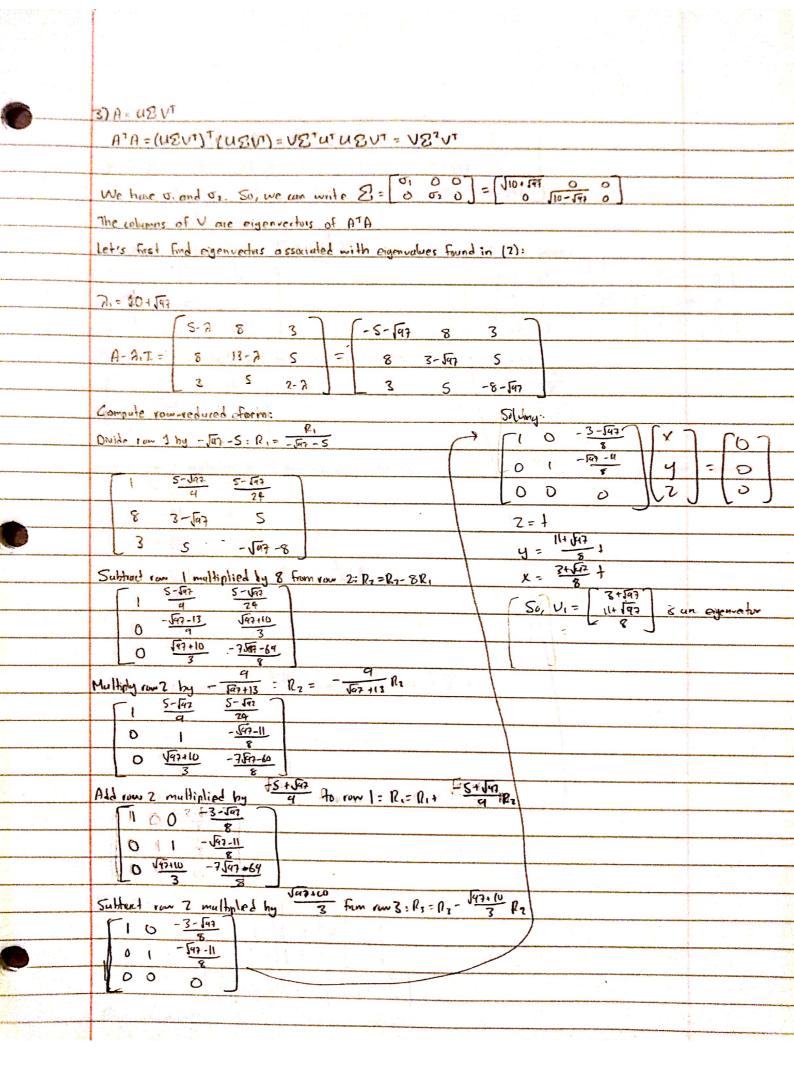


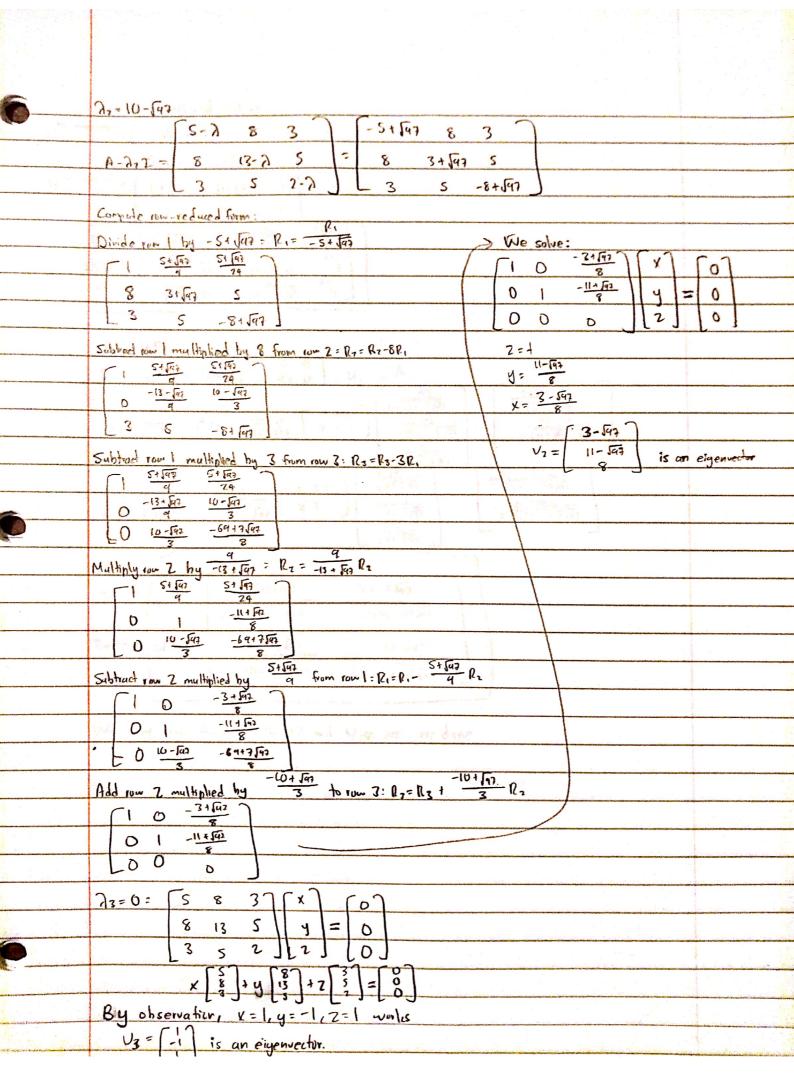
Problem 2) 1) A proof that something is "not" true by a counter-example is a valid analytical proof Consider k. 2 and detect made of 4 points in R: x112=1, x112=2, x113=3, x11=4. Inhabite k-movins with controids M = 2, M = 4 and assume if x 111 is equally distant to multiple controls per. The point will be assigned to the centroid whose order is smallest, i.e k with the smallest value for k cargmin 11 x (1)-Mill? Running k mouns with this initialization: Cluster Assignment: C'= 1 since you = 1 cluser to pri= 2 than pri= 4 C2- 1 since x(2)= 2 closer to pr= 2 than pr= 4 c3=1 since x(3)=3 change distance to me and probat select me since index convention co= 2 snc x(4)= 4 cluser to µ = 9 thun pr= 2 Move controlds: H = 1+7+3 2 = 2 , H2 = 4/=4 Cluster controls have not changed, so algorithm terminates The distortion is: J= m & 11x 11- Mem 1/2 = $\frac{1}{4} \left[(1-2)^{2} \cdot (2-2)^{2} + (3-2)^{2} \cdot (4-4)^{2} \right]$ = 4 (1+0+1+0) = 4 To show this is not globally optimal solution, we will start with a different inhalization and show this converges to a smaller distortion $x^{(1)}=1$ $x^{(2)}=2$ $x^{(3)}=3$ $x^{(4)}=9$ Cluster Assignment: [1]=1, (12)=1, (13)=2, (14)=2 (hused on minimum distance) Move centrale: M1 = 1+2 = 1.5 M2 = 2+4 = 3.5 The distribut in this cuse is: J2 = 1 2 11x10 - M. 11/2 = 1/4 ((x111-MLIN) 2+ - -+ (x10)-MLIN)2) = 4((4)249) = 4

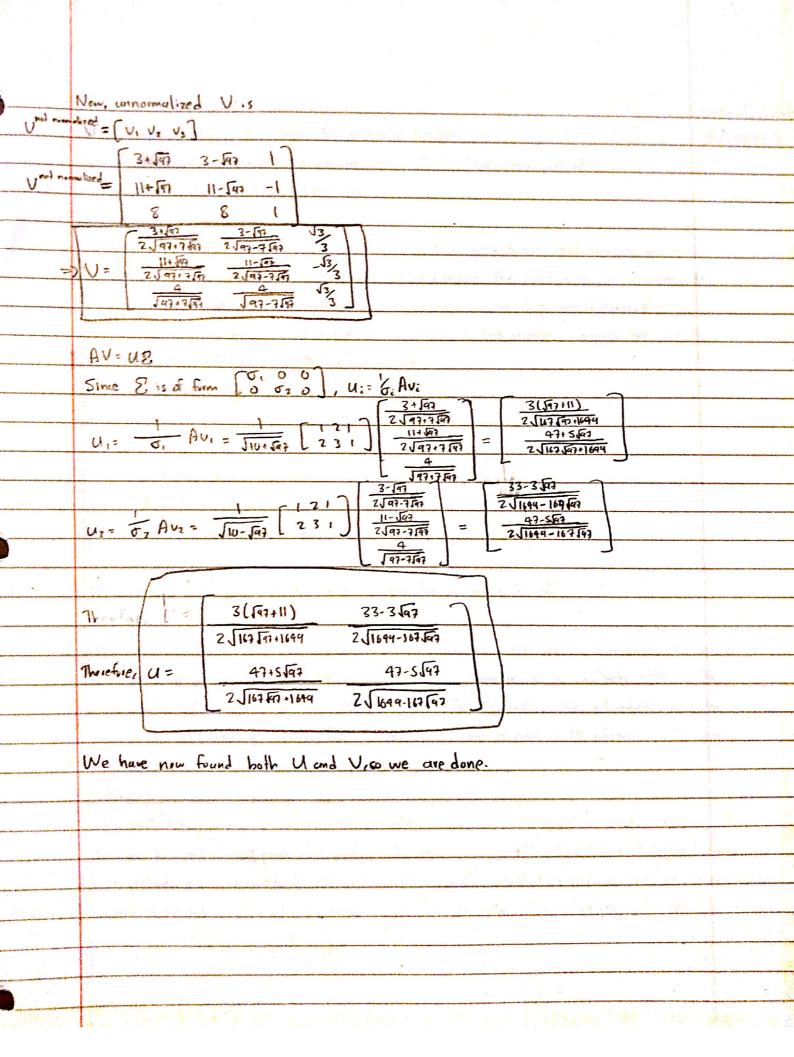
Since J= 4 < 2 = J., the cluster centroids that we obtain in the first cuse (after the algorithm ends)

drave higher distriction than this case, so are not a globally optimal solution

Publem 3) SVD 1) The rank of A is the number of linearly independent rows of A Since row 2 cannot be written as a scalar multiple of row 1, the rows of motive A are linearly independent. Alternatively, we can show the rows are linearly independent by showing: a. [] . a. [] = 0 implies d.= x.= 0. no do this, we have: Q+ 707 = 0 2 x1+3 d1=0 @ 3 d,+ a,= 0 0-0: (a2=0) (2-13: 0,+70,=0 =) a,+26)=0=) [a,=0] 50 the rows are independent. 2 linearly independent rows means runk (A) = 2 Eigenvalues of ATA: Set: det (ATA-AI) = 0 =0 (5-7) det ([13-7 5-7]) - 8 det ([8 5]) + 3 det ([8 13-7] (5-7)[(13-7)17-21-25]-8(8(7-2)-15)+3(40-3(13-21)=0 (5-7) (26-157+72-75) -8 (16-87-15) +3 (40-39+37)=0 (5-7)(72-157+1)-8(1-87)+3(1+37)=0 572-757+5-73+1572-7-8+697+3+97=0 -23+2072-37=0 73-2072+37=0 7(72-207+3)=0 $\gamma = 0$ or $\lambda = \frac{20 \pm \sqrt{20^2 - 4(1 \times 3)}}{2} = \frac{20 \pm \sqrt{400 - 12}}{2} = 10 \pm \sqrt{97}$ Singular values are square rout of nonzero eigenvalues: J= 110+197 Vz = \$ 10- 197







1) Implement kmeans function- this performs one trial of the k-means algorithm, starting with a random initialization of k data points

```
import numpy as np
In [3]:
         def kmeans(data, k):
             num examples, num features = data.shape
             old centroids = np.zeros((k, num features))
             random indices = np.random.choice(num examples, size=k, replace=False)
             cluster centroids = data[random indices, :]
             cluster assignments = [0]*num examples
             repeat = True
             while repeat:
                 # Cluster Assignment Step
                 clusteridx_to examples = dict()
                 for i in range(num examples):
                     # Find cluster point closest to data point
                     mindist, minidx = np.linalg.norm(data[i]-cluster centroids[0]), 0
                     for j in range(1, k):
                         if np.linalg.norm(data[i]-cluster centroids[j]) < mindist:</pre>
                             mindist, minidx = np.linalq.norm(data[i]-cluster centroids[j]), j
                     # Store data point in index to examples dictionary
                     cluster assignments[i] = minidx
                     if minidx in clusteridx to examples:
                         clusteridx to examples[minidx].append(i)
                     else:
                         clusteridx to examples[minidx] = [i]
                 # Move centroids step
                 for clusteridx, examples in cluster in clusteridx to examples.items():
                     cluster centroids[clusteridx] = np.mean(np.array([data[i] for i in examples in cluster]), axis=0)
                 # Repeat iff cluster centroids have changed
                 comparison = old centroids == cluster centroids
                 repeat = not comparison.all()
                 old centroids = np.copy(cluster centroids)
             return cluster centroids, cluster assignments
```

2) Run kmeans for breast cancer data for 2 <= k <= 8

```
from sklearn.datasets import load breast cancer
In [6]:
          dataset = load breast cancer()
          data = dataset.data
          num examples = data.shape[0]
          distortions = []
          for k in range(2,8):
              cluster centroids, cluster assignments = kmeans(data, k)
              distortion = 0
              for i in range(num examples):
                  distortion += np.linalg.norm(data[i] - cluster centroids[cluster assignments[i]])**2
              distortions.append(distortion/num examples)
          distortions
Out[6]: [136982.6008405956,
          88783.42629047789,
          51364.748070263326,
          36331.63689693994,
          30099.055519437814,
          23468.0867249605841
        3) Plot distortion vs k
In [10]:
          import matplotlib.pyplot as plt
          plt.scatter(list(range(2,8)), distortions)
          plt.title("Distortion vs k")
          plt.xlabel("k")
          plt.ylabel("Distortion")
Out[10]: Text(0, 0.5, 'Distortion')
```

