## Problem 1

• I used four different initial seeds as means to do the K-Means clustering, as shown in Figure 1 and 2. And I used square, triangle and sphere to represent data points from the 3 Gaussian distributions, respectively. We can see that the distribution of clustering results are similar in all four situations. The change of J(r, u) is shown in Figure 3. We can see that J(r, u) of different initial seeds all converged to the same value, except for a minor difference in the converging rate.

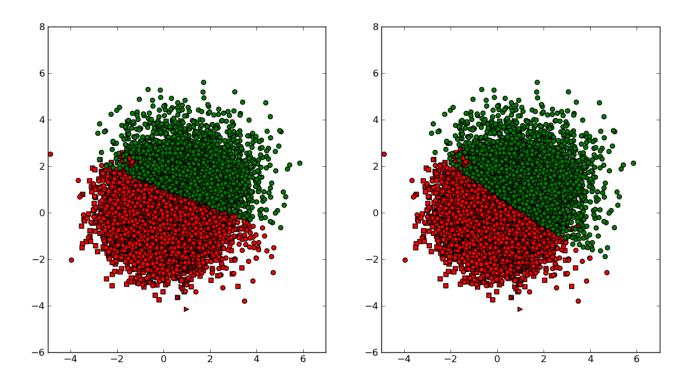


Figure 1: Visualization of K-Means Clustering Using Two Different Initial Seeds (K=2)

• I keep the initial seed the same, but change the value of k to be 2, 3, ..., 6. The visualization of clustering and change of J(r, u) are shown in Figures 4, 5 and 6. The value of J(r, u) (Figure 6) gets lower as Iteration Number increases; the minimum values of J(r, u) also decreases as K increases: for K = 2, 3, 4, 5, 6 J(r, u) are 236,190,150,117, and 107 (rounded).

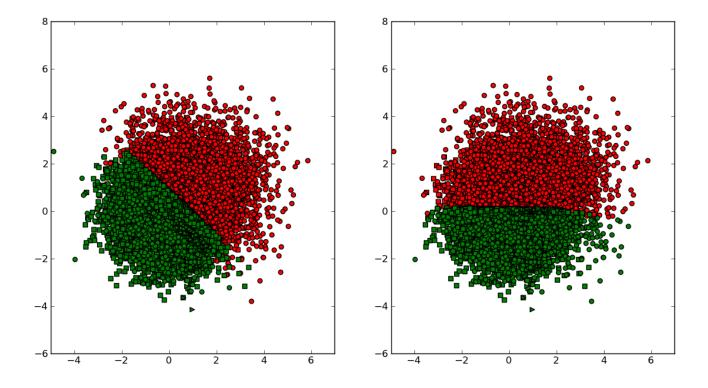


Figure 2: Visualization of K-Means Clustering Using Two Other Different Initial Seeds (K=2)

So when K=6 the value of J(r,u) gets the lowest. The distributions of clustering also differ a lot with different K values. To figure out the best K value, I also calculated the purity in each cluster, which is the maximum percentage of points from the same Gaussian distribution. The larger the purity, the more similar the points are in the cluster, so the better clustering it is. Looking through K=2,...,6, we want a clustering that has relatively high purity in each cluster and also has a relatively low J(r,u). Balancing both characteristics, I think K=4 does the best job. It has a J(r,u)=150, for away from the maximum J(r,u)=236, and not far away from the minimum J(r,u)=107. And the purity when K=4 is [0.91176470588235292, 0.7857142857142857, 0.8888888888884, 0.55172413793103448], which is relatively high for all clusters, comparing to other clustering methods.

## K=2,3,4,5,6!!

K: 2 Seed: 30

Iteration: 1 J(r,u): 349.389115369
Iteration: 2 J(r,u): 262.695090655
Iteration: 3 J(r,u): 250.745103117
Iteration: 4 J(r,u): 247.601074241

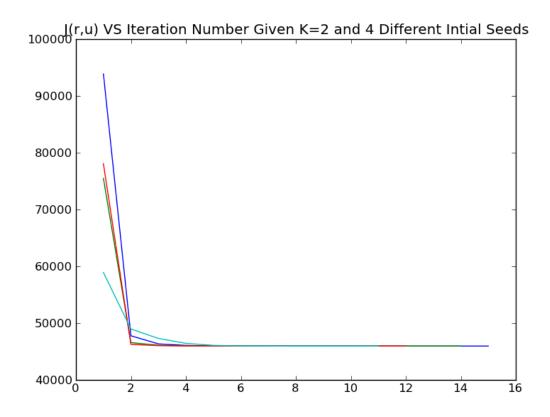


Figure 3: J(r, u) v.s. Iteration Numbers

Iteration: 5 J(r,u): 240.7570529
Iteration: 6 J(r,u): 236.19315226

Purities in 2 classes:

[0.77966101694915257, 0.58536585365853655]

K: 3 Seed: 30

Iteration: 1 J(r,u): 255.748828223
Iteration: 2 J(r,u): 191.034795635
Iteration: 3 J(r,u): 190.611716202
Iteration: 4 J(r,u): 190.238116299

Purities in 3 classes:

[0.5, 0.86956521739130432, 0.65625]

K: 4 Seed: 30

Iteration: 1 J(r,u): 220.464413702
Iteration: 2 J(r,u): 154.392972069
Iteration: 3 J(r,u): 150.385259237

Purities in 4 classes:

K: 5 Seed: 30

Iteration: 1 J(r,u): 196.781801235
Iteration: 2 J(r,u): 129.736789092
Iteration: 3 J(r,u): 119.810794795
Iteration: 4 J(r,u): 116.849929349

Purities in 5 classes:

K: 6 Seed: 30

Iteration: 1 J(r,u): 213.616799381
Iteration: 2 J(r,u): 115.70384947
Iteration: 3 J(r,u): 107.418080073

Purities in 6 classes:

[1.0, 0.8125, 1.0, 0.5714285714285714, 0.83870967741935487, 0.47826086956521741]

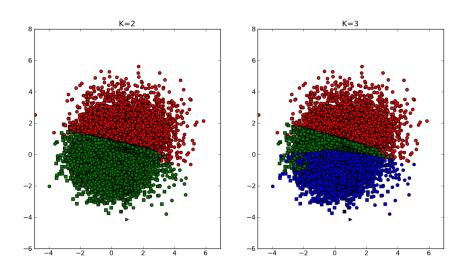


Figure 4: Visualization of K-Means Clustering Using K = 2, 3

- I kept K = 3 and used 10 different random initial seeds to test on the distribution of clustering. The visualization of clustering is shown in Figure 7, 8, 9, 10, and 11. The change of J(r, u) is shown in Figure 12. We can easily see that the 10 different random seeds have similar results in both the clustering distribution, and the minimization of distances (J(r, u)).
- I kept K = 3 and the same initial seed, but changed the mean values of the 3 Gaussian distributions from [0,0], [1,0], [1,1] to [-1,0], [1,0], [0,1] (Spread-out) and [0,0], [0,0.5], [0,1] (Close-up). The purity of each clustering are shown below, we learn that the more close-up the points are, the lower purity the clustering is. So the Spread-out points have the highest purity and

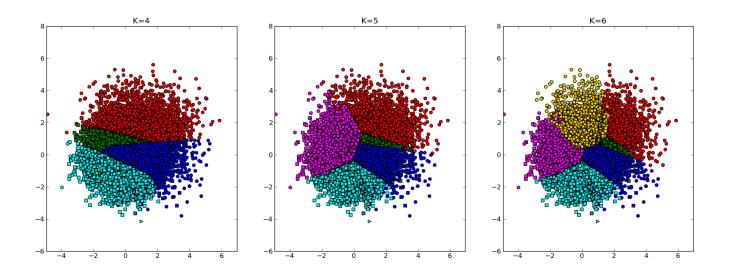


Figure 5: Visualization of K-Means Clustering Using K = 4, 5, 6

thus best in separation. The clustering visualization is shown in Figure 13. When looking at the change of J(r,u) in Figure 14, we find that the more close-up the points are, the easier to find a convergence of J(r,u), and the value of J(r,u) is also low. That tells us that we can not only determine the quality of clustering based on the value and convergence of J(r,u) but we need to also look at the purity.

```
3 Different Mean Values Given K=3!!
```

```
Original Means
Mean1: [0, 0] Mean2: [1, 0] Mean3: [0, 1]
K: 3 Seed: 30
...
Iteration: 34   J(r,u): 35512.468781
Purities in 3 classes:
[0.49061196105702365, 0.9263759828448892, 0.584287317620651]
Spread-out Means
Mean1: [-1, 0] Mean2: [1, 0] Mean3: [0, 1]
K: 3 Seed: 30
...
Iteration: 35   J(r,u): 36632.0234786
Purities in 3 classes:
[0.45540075990591639, 0.966066229985444, 0.41855788566503876]
Close-up Means
Mean1: [0, 0] Mean2: [0, 0.5] Mean3: [0, 1]
```

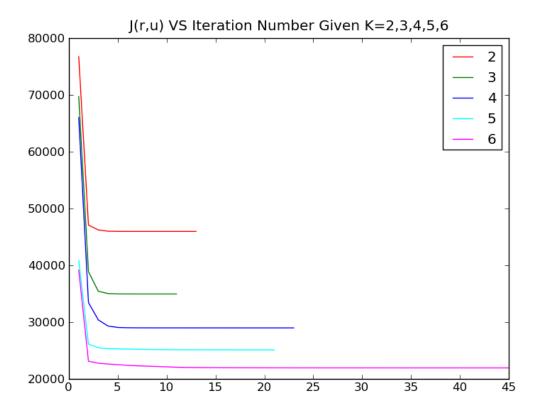


Figure 6: J(r, u) v.s. Iteration Numbers

```
K: 3 Seed: 30
...
Iteration: 22  J(r,u): 35143.2752124
Purities in 3 classes:
[0.42643391521197005, 0.91406617531754741, 0.73117742539642905]
```

• I kept K=3 and the same initial seed, but changed the variance values of the 3 Gaussian distributions from

$$[[1,0],[0,1]],[[0.1,0],[0,1]],[[2,0],[0,2]]$$
 to (Small Variance) 
$$[[0.1,0],[0,0.1]],[[0.01,0],[0,0.1]],[[0.2,0],[0,0.2]]$$
 and (Large Variance) 
$$[[3,0],[0,3]],[[0.3,0],[0,3]],[[6,0],[0,6]]$$

. Obviously, when the variance is small, the points are close-up to the center of each Gaussian distribution, thus K-Means clustering has a better separation in the points, so the purity for

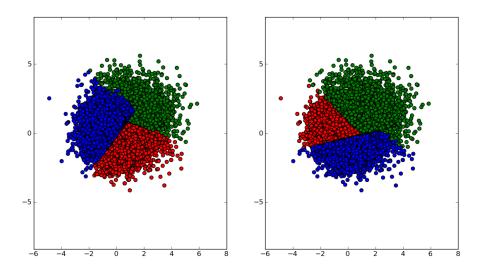


Figure 7: Visualization of K-Means Clustering Using Two Different Initial Seeds (K=3)

small variance is the best. We can also look at the Small Variance subplot in Figure 16, which shows that most of the points are clustered into the center shown in blue, and that agrees with the original model that 70% of points comes from the center (mean is [0,0]) Gaussian distribution. A small variance also helps in the convergence in J(r,u) and has a lower J(r,u) value as shown in Figure 16.

```
3 Different Variances Given K=3!!
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```
Var1: [[1, 0], [0, 1]] Var2: [[0.1, 0], [0, 1]] Var3: [[2, 0], [0, 2]]

K: 3 Seed: 30
...
Iteration: 34   J(r,u): 35512.468781
Purities in 3 classes:
[0.49061196105702365, 0.9263759828448892, 0.584287317620651]

Var1: [[0.1, 0], [0, 0.1]] Var2: [[0.01, 0], [0, 0.1]] Var3: [[0.2, 0], [0, 0.2]]

K: 3 Seed: 30
...
Iteration: 9   J(r,u): 4624.61488682
Purities in 3 classes:
[0.74108527131782942, 0.99985211475894709, 0.65734865374721319]

Var1: [[3, 0], [0, 3]] Var2: [[0.3, 0], [0, 3]] Var3: [[6, 0], [0, 6]]
```

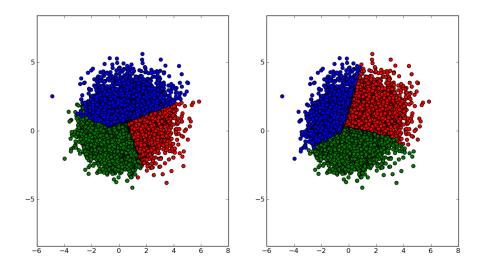


Figure 8: Visualization of K-Means Clustering Using Two Other Different Initial Seeds (K=3)

K: 3 Seed: 30

. . .

Iteration: 48 J(r,u): 80763.949289

Purities in 3 classes:

[0.55916386711459498, 0.74062278839348905, 0.7665476963011032]

• I wrote a program for the Gaussian Mixture Model and the results of clustering assignments are shown in Figures 17; The distribution of each cluster in X and Y dimensions are shown in 18, respectively. For visualization I set the points to be clustered into the Gaussian distribution with highest  $\gamma$  value. I set the maximum iteration to be 10000, and the tolerance of difference in  $\gamma$  as 0.001. The value of difference finally converged in 17 iterations.

diff 1.0

diff 0.247866014954

diff 0.0311447554769

diff 0.0239822131299

diff 0.0209311785987

diff 0.0174222991875

diff 0.0140759698151

diff 0.0111796108491

diff 0.00878255676385

diff 0.00684495360007

diff 0.00530996310052

diff 0.0041120766978

diff 0.00318604679511

diff 0.00247052366001

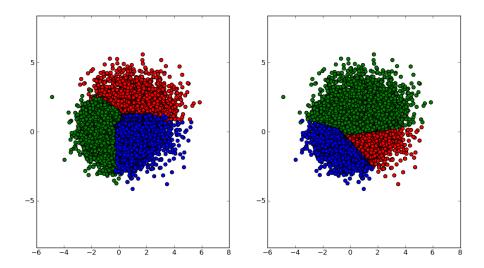


Figure 9: Visualization of K-Means Clustering Using Two Other Different Initial Seeds (K=3)

diff 0.00192263148965

diff 0.00150724858789

diff 0.00119614961777

converged in 17 iterations

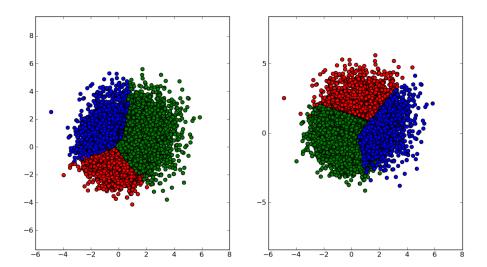


Figure 10: Visualization of K-Means Clustering Using Two Other Different Initial Seeds (K=3)

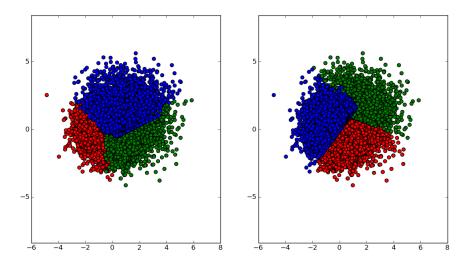


Figure 11: Visualization of K-Means Clustering Using Two Other Different Initial Seeds (K=3)

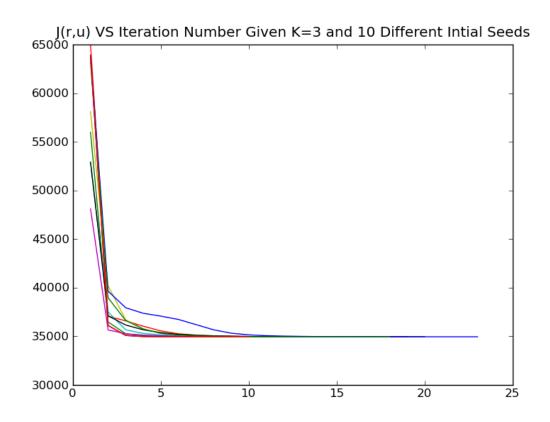


Figure 12: J(r, u) v.s. Iteration Numbers

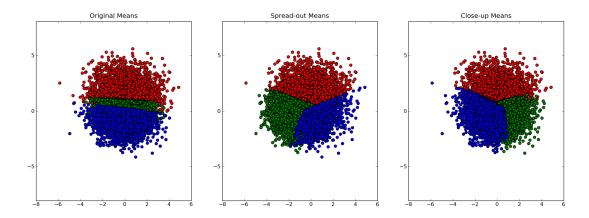


Figure 13: Visualization of K-Means Clustering Using Three Different Mean Values (K=3)

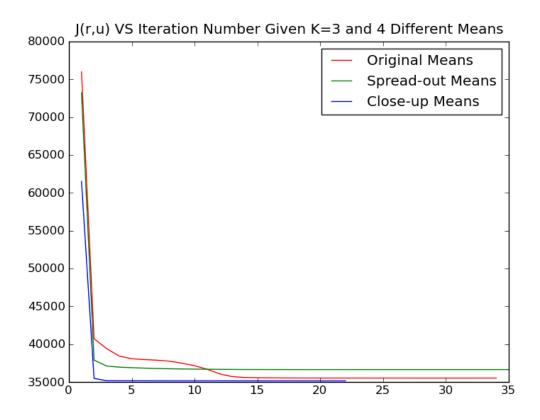


Figure 14: J(r, u) v.s. Iteration Numbers

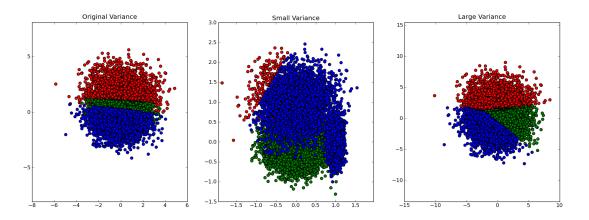


Figure 15: Visualization of K-Means Clustering Using Three Different Variance Values (K=3)

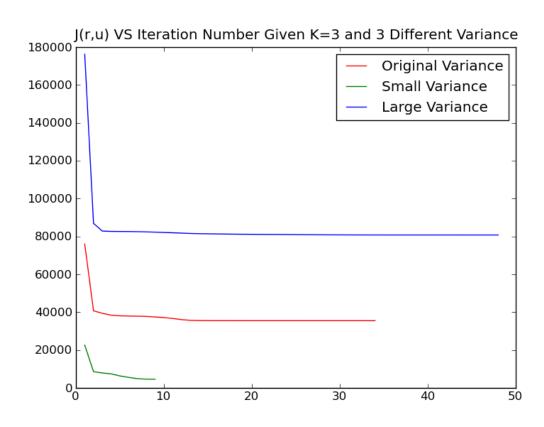


Figure 16: J(r, u) v.s. Iteration Numbers

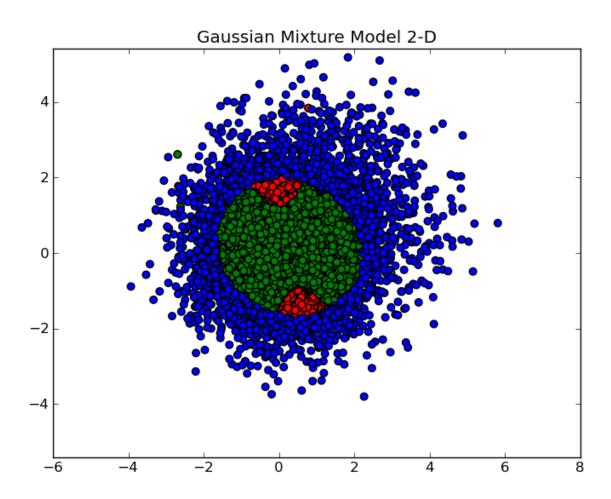


Figure 17: Visualization of Gaussian Mixture Model

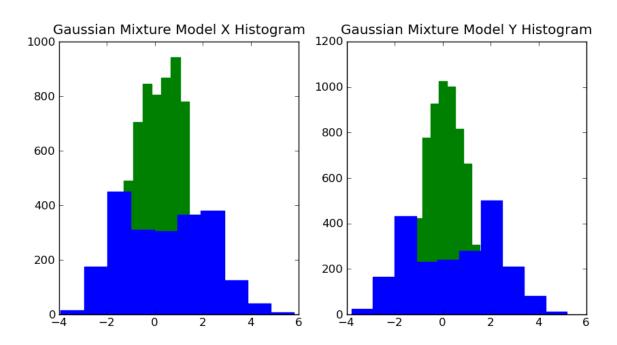


Figure 18: Histogram of Gaussian Mixture Model