Assignment#2 - Unsupervised Learning

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1. 테스트 환경

운영체제	Windows 11 23H Education edition
개발환경	Juypter Notebook (local)
Kernel	Python 3 (ipykernel)

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2. 실험 준비

a. Dataset

- Fashion MNIST data (테스트 dataset 만 사용)
  - **images**

shape	(10000, 784)
data type	float32 array

  - **labels**

shape	(10000,)
data type	array

- images, labels 전처리
  - images to pandas.DataFrame (변수 이름 : X_)

	0	1	2	3	4	5	6
0	0.0	0.0	0.0	0.000000	0.0	0.000000
1	0.0	0.0	0.0	0.000000	0.0	0.000000
2	0.0	0.0	0.0	0.000000	0.0	0.000000
3	0.0	0.0	0.0	0.000000	0.0	0.000000
4	0.0	0.0	0.0	0.007843	0.0	0.003922
...	...	...	...	...	...	...
9995	0.0	0.0	0.0	0.000000	0.0	0.000000
9996	0.0	0.0	0.0	0.000000	0.0	0.000000
9997	0.0	0.0	0.0	0.000000	0.0	0.000000
9998	0.0	0.0	0.0	0.000000	0.0	0.000000
9999	0.0	0.0	0.0	0.000000	0.0	0.000000

10000 rows × 784 columns

- labels to pandas.Series (변수 이름 : y_)

```
0       9
1       2
2       1
3       1
4       6
       ..
9995    9
9996    1
9997    8
9998    1
9999    5
Length: 10000, dtype: int64
```

b. Experimental Setup

Clustering Algorithm

- **KMeans**
  - **구현체** : sklearn.cluster.KMeans 클래스
  - **Hyper Paramter** :
    - n_cluster : 10
    - 나머지 파라미터는 모두 default parameter set 사용

- **DBSCAN**
  - **구현체** : sklearn.cluster.DBSCAN
  - **Hyper Parameter** :
    - eps (epsilon) : parameter search 알고리즘을 통해 구한 최적 epsilon 값
    - min_samples : parameter search 알고리즘을 통해 구한 최적 min_samples 값
    - n_jobs : -1 (all processor)
    - 나머지 파라미터는 모두 default parameter set 사용

c. Pseudo code of Algorithms

아래에 이번 과제에서 사용한 알고리즘들을 pseudo code형태로 나열.

- **Helper algorithms for main learning process**

[KMeans clustering algorithm]

Algorithm 1 K-means Clustering

```
procedure KMeansClustering(X, n_clusters = 10)
    kmeans ← KMeans(n_clusters = n_clusters).fit(X)
    labels ← kmeans.labels_
    return labels
end procedure
```

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[DBSCAN clustering algorithm]

Algorithm 2 DBSCAN Clustering

```
procedure DBSCANClustering(X, eps, min_samples)
    dbscan ← DBSCAN(eps = eps, min_samples = min_samples, n_jobs = −1).fit(X)
    return dbscan.labels_
end procedure
```

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[Transform from PCA dataset to t-SNE]

Algorithm 3 Transform PCA to t-SNE

```
procedure Transform PCA to t-SNE(X)
    n_components ← 2
    learning_rate ← 300
    perplexity ← 30
    early_exaggeration ← 12
    init ←' random'
    tSNE ← Create tSNE with parameters
    test_PCA ← DataFrame(X)
    X_test_tSNE ← tSNE.fit and transform PCA to tSNE
    X_test_tSNE ← convert tSNE to Dataframe(X_test_tSNE)
    return X_test_tSNE
end procedure
```

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[Hyper parameter search for DBSCAN]

Algorithm 1 Find Best Parameters for DBSCAN

```
procedure FindBestParamsDBSCAN(true_labels)
    eps_list ← arange(0.5, 10.0, 0.1)
    min_sample_list ← arange(5, 105, 1)
    best_params_per_dim ← {dim : [] for dim in [784, 100, 50, 10]}
    for dim, X_test_tSNE in zip(dimensions, X_transformed) do
        for eps in eps_list do
            for min_sample in min_sample_list do
                labels ←DBSCAN_Clustering(X_test_tSNE, eps, min_sample)
                cur_score ← Evaluate_Clustering(true_labels, labels)
                if cluster_count ≥ 9 then
                    Append new ARI score, eps and min sample
                    sort params by ARI in descending order
                    slice params at index 2
                end if
            end for
        end for
    end for
    return best_params_per_dim
end procedure
```

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3. 실험 진행

아래의 pseudo code를 따라,

- 각 차원 784, 100, 54, 10에 대해
  - PCA 적용
  - tSNE 변형
  - 변형된 데이터셋에 대해 Kmeans 와 dbscan 클러스터링 알고리즘으로 학습후, ARI score계산 및 시각 (dbscan은 parameter search로 얻은 최적 params중 임의로 선택하여 통일 적용)

a. learning process

Algorithm 1 Main Execution

```
procedure MainExecution(X_, y_)
    dimensions ← [784, 100, 50, 10]
    ari_kmeans ← []
    ari_dbscan ← []
    for dim in dimensions do
        if dim = 784 then
            X_reduced ← X_                          ▷ Original image dimensions
        else
            X_reduced ← ApplyPCA(X_, dim)
        end if
        running algorithms and visualize for both kmeans and dbscan
    end for
end procedure
```

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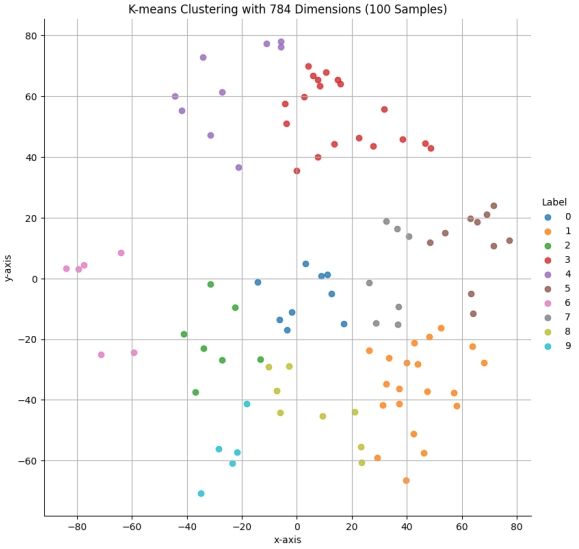
4. 실험 결과

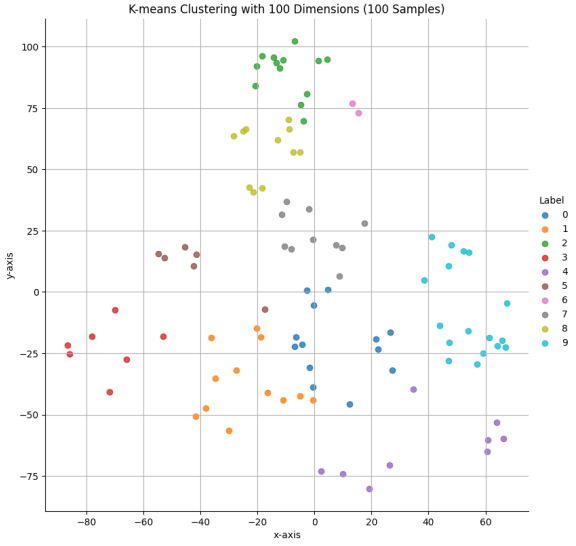
a. ARI scores for both of KMeans and DBSCAN

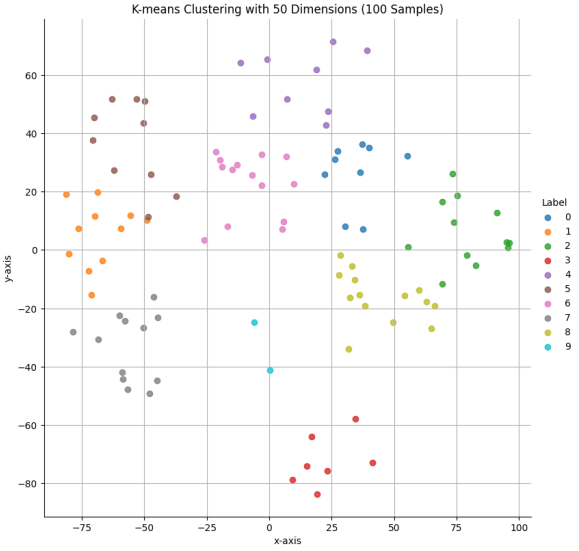
	Dimensions	K-means ARI	DBSCAN ARI
0	784	0.436577	0.440892
1	100	0.461160	0.424607
2	50	0.405413	0.438050
3	10	0.425742	0.400305

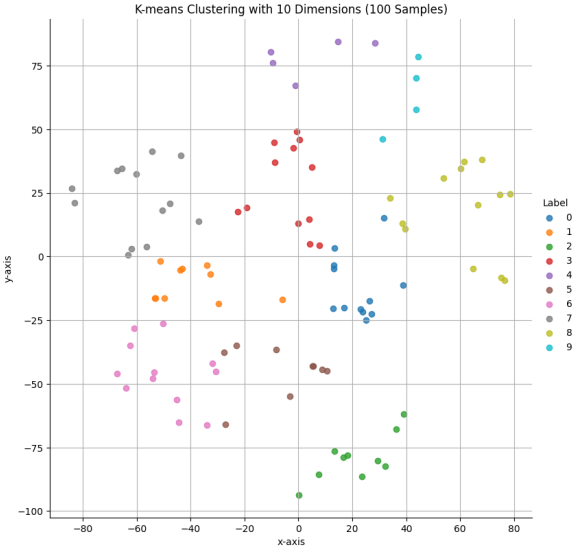
b. KMeans visualizatoin

Kmeans cluster 100 samples plotted

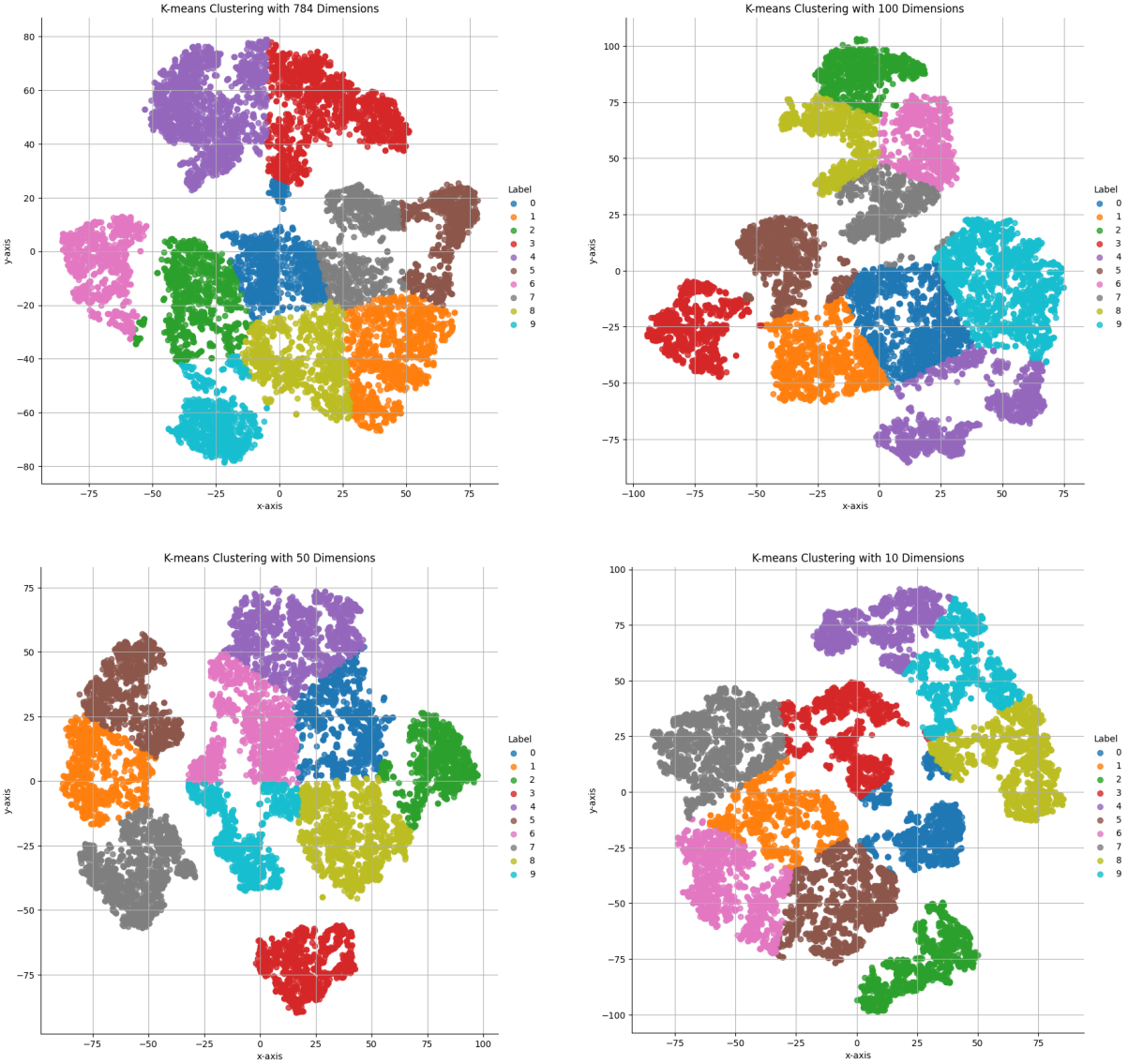






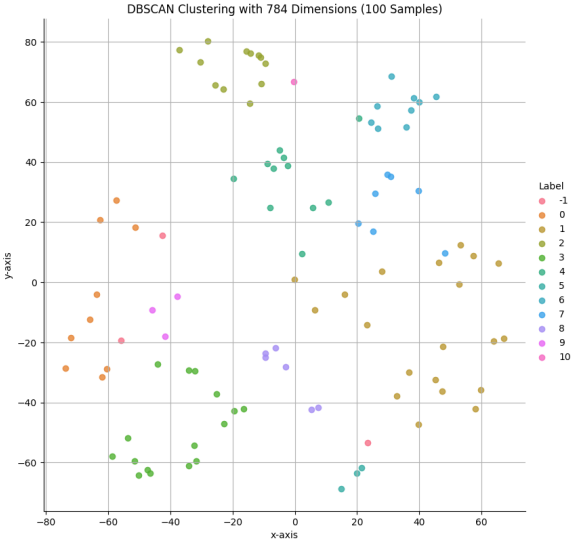


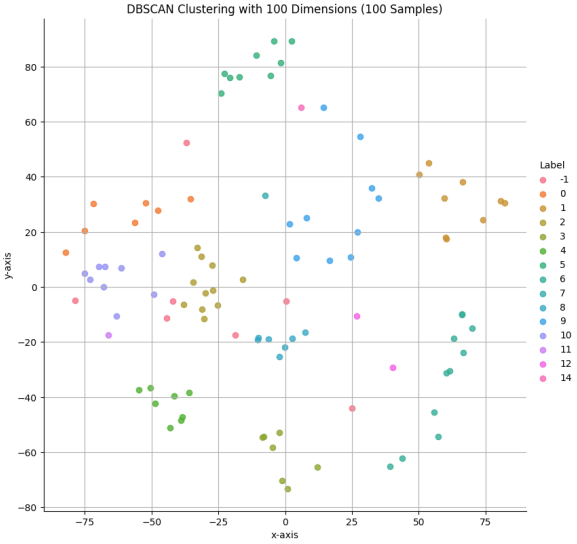
KMeans cluster Without sampling

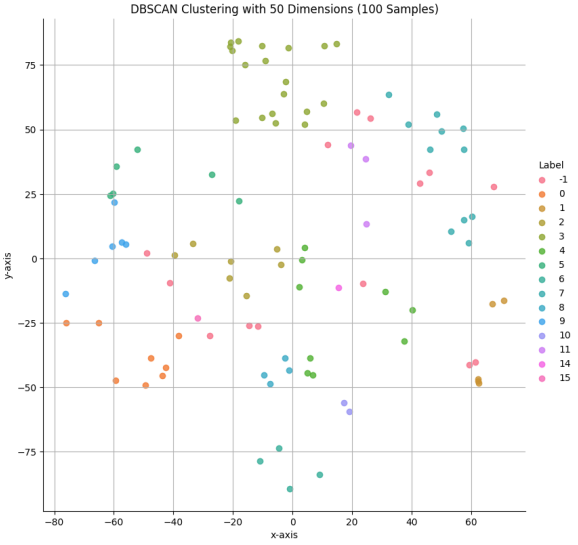


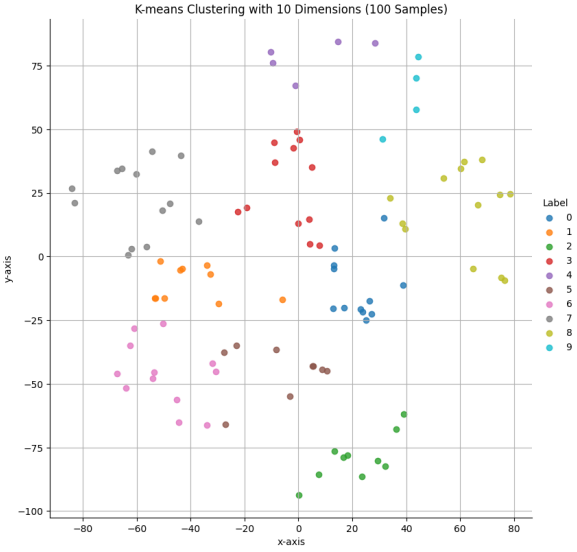
c. DBSCAN visualization (plotted 100 samples)

DBSCAN 100 samples plotted

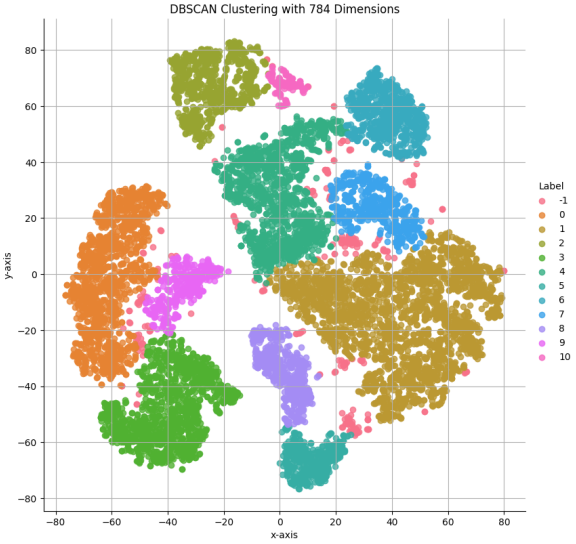


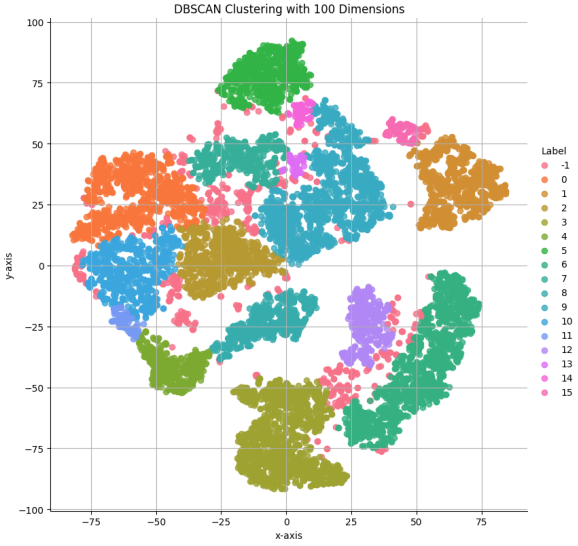


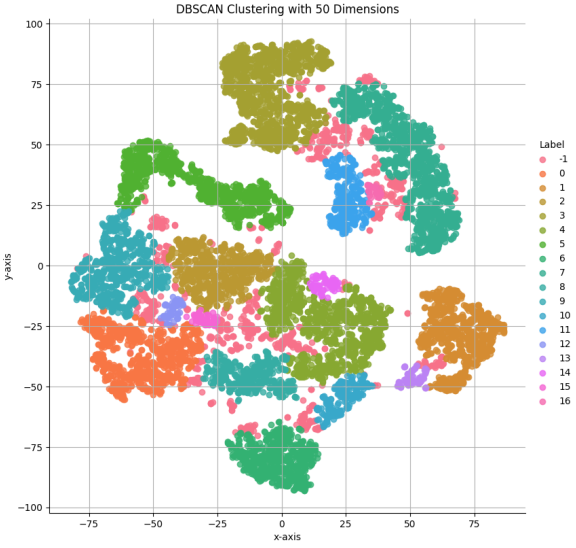


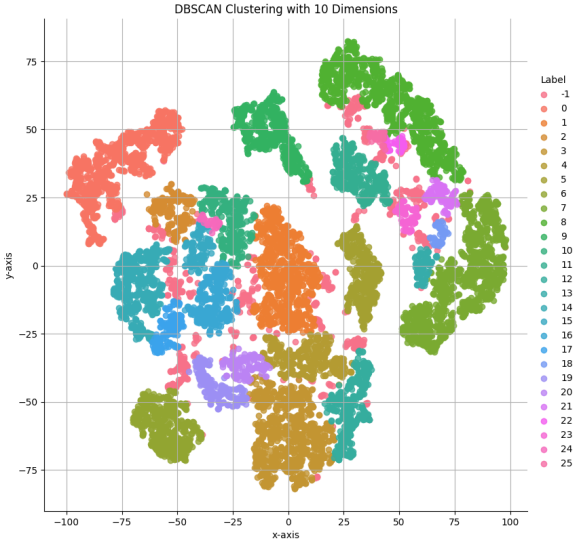


DBSCAN Without sampling


DBSCAN Clustering with 784 Dimensions
x-axis
y-axis
Label
-1
0
1
2
3
4
5
6
7
8
9
10


DBSCAN Clustering with 100 Dimensions
x-axis
y-axis
Label
-1
0
1
2
3
4
5
6
7
8
9
10
11
12
13
14
15


DBSCAN Clustering with 50 Dimensions
x-axis
y-axis
Label
-1
0
1
2
3
4
5
6
7
8
9
10
11
12
13
14
15
16


DBSCAN Clustering with 10 Dimensions
x-axis
y-axis
Label
-1
0
1
2
3
4
5
6
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