# IF4091 Pembelajaran Mesin

# Tugas Kecil I: Eksplorasi scikit-learn untuk Clustering pada Jupyter Notebook

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Kelas: K01

# **Import Statements**

```
In [37]:
```

```
from matplotlib.colors import LogNorm
from mst_clustering import MSTClustering
from pyclustering.cluster.bang import bang, bang_visualizer
from pyclustering.cluster.kmedoids import kmedoids
from sklearn import datasets
from sklearn.decomposition import PCA
from sklearn.externals import joblib
from sklearn.cluster import AgglomerativeClustering, DBSCAN, KMeans
from sklearn.metrics.pairwise import pairwise_distances
from sklearn.mixture import GaussianMixture
from sklearn.preprocessing import LabelEncoder

import matplotlib.pyplot as plt
import numpy as np
import pandas as pd
import pylab as pl
```

# **Load Datasets**

Pada tahap load, datasets diload kemudian labelnya dihapus karena ingin melakukan clustering (unsupervised learning)

```
In [38]:
```

# **Preprocessing**

Pada tahap preprocessing, data tennis akan diproses sehingga nilai setiap atribut diubah menjadi atribut integer dengan menggunakan LabelEncoder.

Setelah itu, data iris dan tennis akan ditampilkan dengan menggunakan PCA sehingga data yang memiliki dimensi lebih dari 2 dapat direpresentasikan dengan plot 2 dimensi.

```
In [39]:
```

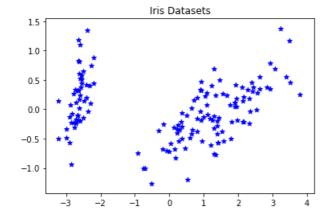
```
# change string/object values to integer
le = LabelEncoder()
tennis_df = pd.DataFrame()
for column in tennis_df_raw:
```

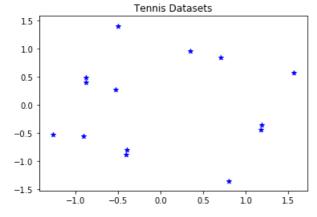
```
tennis_df[column] = le.fit_transform(tennis_df_raw[column])
```

```
In [40]:
```

```
def plot_initial(df, title):
    pca = PCA(n_components=2).fit(df)
    pca_2d = pca.transform(df)
    for i in range(0, pca_2d.shape[0]):
        c = pl.scatter(pca_2d[i,0],pca_2d[i,1],c='b',marker='*')
    pl.title(title)
    pl.show()

plot_initial(iris_df, "Iris Datasets")
plot_initial(tennis_df, "Tennis Datasets")
```





# **Model Training**

```
In [41]:
```

```
k_iris = 3
k_tennis = 2
```

#### In [42]:

```
def plot_cluster(df, model_labels, title):
    pca = PCA(n_components=2).fit(df)
    pca_2d = pca.transform(df)
    unique_labels = list(set(model_labels))
    colors = ['r', 'g', 'b']
    markers = ['+', 'o', '*']
    clusters = [None] * len(unique_labels)
    for i in range(0, pca_2d.shape[0]):
        idx = unique_labels.index(model_labels[i])
        clusters[idx] = pl.scatter(pca_2d[i,0], pca_2d[i,1], c=colors[idx], marker=markers[idx])
    legends = []
    for i, cluster in enumerate(clusters):
        if cluster != None:
            legends.append('Cluster ' + str(i))
    pl.legend(clusters, legends)
    pl.title(title)
```

```
pl.show()
def plot cluster DBSCAN(df, model labels, title):
   pca = PCA(n components=2).fit(df)
    pca 2d = pca.transform(df)
    unique_labels = list(set(model_labels))
    colors = ['r', 'g', 'b', 'k']
markers = ['+', 'o', '*', 'v']
    clusters = [None] * len(unique labels)
    is_core = [True] * len(unique labels)
    for i in range(0, pca_2d.shape[0]):
        idx = unique_labels.index(model_labels[i])
        is_core[idx] = model_labels[i] != -1
        clusters[idx] = pl.scatter(pca_2d[i,0], pca_2d[i,1], c=colors[idx], marker=markers[idx])
    legends = []
    for i, cluster in enumerate(clusters):
        if cluster != None:
            if is core[i]:
                legends.append('Cluster ' + str(i))
                legends.append('Outlier')
    pl.legend(clusters, legends)
    pl.title(title)
    pl.show()
def plot cluster GMM(df, k, title, cov='full', threshold=0.001):
    pca = PCA(n_components=2).fit(df)
    pca 2d = pca.transform(df)
    model = GaussianMixture(n components=k, covariance type=cov, tol=threshold).fit(pca 2d)
    x = np.linspace(-20., 30.)
    y = np.linspace(-20., 40.)
    X, Y = np.meshgrid(x, y)
    XX = np.array([X.ravel(), Y.ravel()]).T
    Z = -model.score samples(XX)
    Z = Z.reshape(X.shape)
    CS = plt.contour(X, Y, Z, norm=LogNorm(vmin=1.0, vmax=1000.0),
                     levels=np.logspace(0, 3, 10))
    CB = plt.colorbar(CS, shrink=0.8, extend='both')
    plt.scatter(pca_2d[:, 0], pca_2d[:, 1], .8)
    plt.title(title)
    plt.axis('tight')
    plt.show()
```

# 1. Agglomerative Clustering

Parameter Agglomerative Clustering:

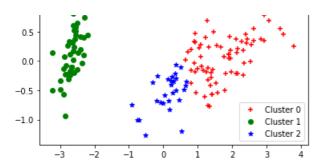
- 1. n\_clusters: jumlah cluster (default=2)
- 2. affinity: metrik yang digunakan untuk menghitung jarak linkage (default=euclidean)
- 3. connectivity: matriks ketetanggan node, untuk structured hierarchical algorithm (default=None)
- 4. compute\_full\_tree: apakah proses clustering dihentikan setelah n\_cluster iterasi atau tidak (default=auto)
- 5. linkage: kriteria linkage yang digunakan, misalnya 'ward', 'complete', 'average' (default=ward)

#### a. Eksperimen Agglomerative Clustering dengan parameter linkage complete

```
In [43]:
```

```
iris_agglo1 = AgglomerativeClustering(n_clusters=k_iris, linkage='complete').fit(iris_df)
tennis_agglo1 = AgglomerativeClustering(n_clusters=k_tennis, linkage='complete').fit(tennis_df)

plot_cluster(iris_df, iris_agglo1.labels_, "Iris Agglomerative Clustering Clusters - Complete Linkage")
plot_cluster(tennis_df, tennis_agglo1.labels_, "Tennis Agglomerative Clustering Clusters -
Complete Linkage")
```



# 

0.0

# b. Eksperimen Agglomerative Clustering dengan parameter linkage ward

0.5

1.0

1.5

#### In [44]:

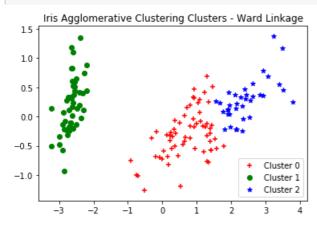
-1.5

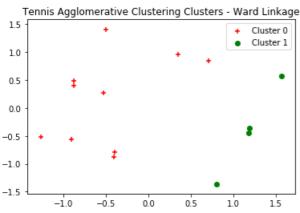
-1.0

-0.5

iris\_agglo = AgglomerativeClustering(n\_clusters=k\_iris, affinity='euclidean', linkage='ward').fit(i
ris\_df)
tennis\_agglo = AgglomerativeClustering(n\_clusters=k\_tennis, affinity='euclidean', linkage='ward').f
it(tennis\_df)

plot\_cluster(iris\_df, iris\_agglo.labels\_, "Iris Agglomerative Clustering Clusters - Ward Linkage")
plot\_cluster(tennis\_df, tennis\_agglo.labels\_, "Tennis Agglomerative Clustering Clusters - Ward Linkage")





## c. Simpan model

#### In [45]:

```
joblib.dump(iris_agglo, 'iris_agglo.pkl')
joblib.dump(tennis_agglo, 'tennis_agglo.pkl')
print("Agglomerative Clustering models saved")
```

Agglomerative Clustering models saved

# 2. DBSCAN

Beberapa parameter DBSCAN:

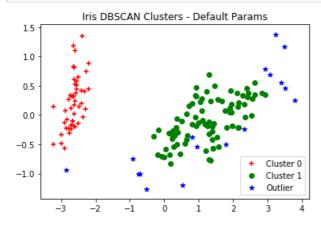
- 1. eps: jarak maksimum diantara 2 data yang dianggap saling bertetangga (default=0.5)
- 2. min\_samples: jumlah tetangga data minimal untuk dianggap sebagai core point (default=5)
- 3. metric: metrik yang digunakan untuk menghitung jarak antar data (default-euclidean
- 4. algorithm: algoritma NN, misalnya 'auto', 'ball\_tree', 'kd\_tree', 'brute' (default=auto)

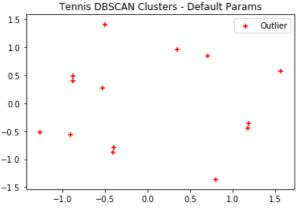
#### a. Eksperimen DBSCAN dengan parameter default

## In [46]:

```
iris_dbscan1 = DBSCAN().fit(iris_df)
tennis_dbscan1 = DBSCAN().fit(tennis_df)

plot_cluster_DBSCAN(iris_df, iris_dbscan1.labels_, "Iris_DBSCAN Clusters - Default Params")
plot_cluster_DBSCAN(tennis_df, tennis_dbscan1.labels_, "Tennis_DBSCAN Clusters - Default Params")
```



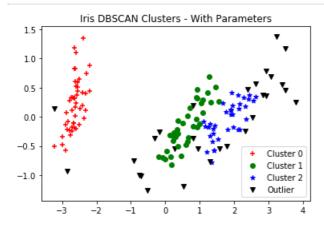


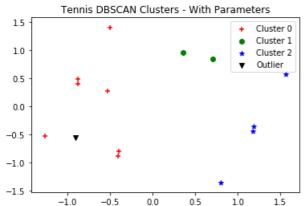
# b. Eksperimen DBSCAN dengan parameter epsilon, min samples, dan algoritma DBSCAN

# In [47]:

```
iris_dbscan = DBSCAN(eps=0.42, min_samples=5, algorithm='kd_tree').fit(iris_df)
tennis_dbscan = DBSCAN(eps=1, min_samples=2, algorithm='kd_tree').fit(tennis_df)

plot_cluster_DBSCAN(iris_df, iris_dbscan.labels_, "Iris_DBSCAN Clusters - With Parameters")
plot_cluster_DBSCAN(tennis_df, tennis_dbscan.labels_, "Tennis_DBSCAN Clusters - With Parameters")
```





## c. Simpan model

#### In [48]:

```
joblib.dump(iris_dbscan, 'iris_dbscan.pkl')
joblib.dump(tennis_dbscan, 'tennis_dbscan.pkl')
print("DBSCAN models saved")
```

DBSCAN models saved

# 3. K-Means

Beberapa parameter K-Means:

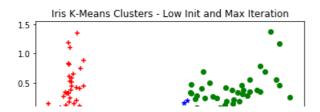
- 1. n\_clusters: jumlah cluster yang dihasilkan (default=8)
- 2. init: metode inisialisasi, misalnya 'k-means++', 'random', atau ndarray (default=k-means++)
- 3. n\_init: jumlah algoritma k-means akan dijalankan pada seed/centroid yang berbeda untuk menemukan seed terbaik
- 4. max\_iter: jumlah iterasi maksimum (default=300)

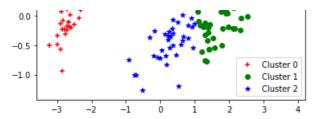
#### a. Eksperimen K-Means dengan parameter n\_init dan jumlah maksimum iterasi yang kecil

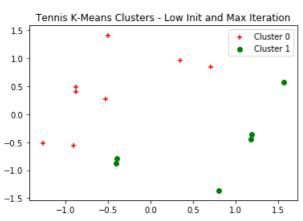
#### In [75]:

```
iris_kmeans1 = KMeans(n_clusters=k_iris, n_init=1, max_iter=2).fit(iris_df)
tennis_kmeans1 = KMeans(n_clusters=k_tennis, n_init=1, max_iter=2).fit(tennis_df)

plot_cluster(iris_df, iris_kmeans1.labels_, "Iris K-Means Clusters - Low Init and Max Iteration")
plot_cluster(tennis_df, tennis_kmeans1.labels_, "Tennis K-Means Clusters - Low Init and Max Iteration")
```





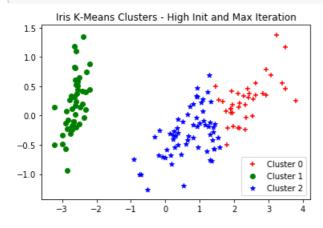


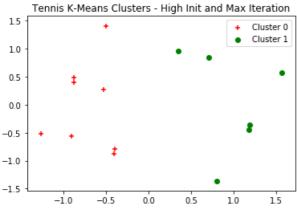
#### b. Eksperimen K-Means dengan parameter n\_init dan jumlah maksimum iterasi besar

#### In [50]:

```
iris_kmeans = KMeans(n_clusters=k_iris, n_init=20, max_iter=100).fit(iris_df)
tennis_kmeans = KMeans(n_clusters=k_tennis, n_init=20, max_iter=100).fit(iris_df)

plot_cluster(iris_df, iris_kmeans.labels_, "Iris K-Means Clusters - High Init and Max Iteration")
plot_cluster(tennis_df, tennis_kmeans.labels_, "Tennis K-Means Clusters - High Init and Max Iteration")
```





#### c. Simpan Model

#### In [51]:

```
joblib.dump(iris_kmeans, 'iris_kmeans.pkl')
joblib.dump(tennis_kmeans, 'tennis_kmeans.pkl')
```

```
print("K-Means models saved")
```

K-Means models saved

# 4. Gaussian Mixture

Beberapa parameter GMM:

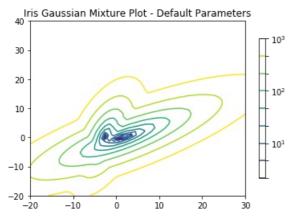
- 1. n\_components: jumlah komponen mixture (default=1)
- 2. covariance\_type: tipe kovariansi, misalnya 'full', 'tied', 'diag', 'spherical' (default=full)
- 3. tol: threshold konvergensi (default=1e-3)

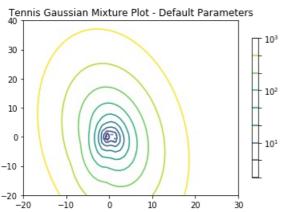
# a. Eksperimen GMM dengan parameter default

#### In [52]:

```
iris_gauss1 = GaussianMixture(n_components=k_iris).fit(iris_df)
tennis_gauss1 = GaussianMixture(n_components=k_tennis).fit(tennis_df)

plot_cluster_GMM(iris_df, k_iris, "Iris Gaussian Mixture Plot - Default Parameters")
plot_cluster_GMM(tennis_df, k_tennis, "Tennis Gaussian Mixture Plot - Default Parameters")
```



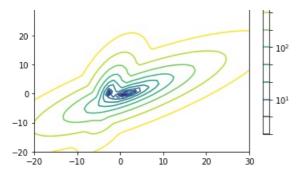


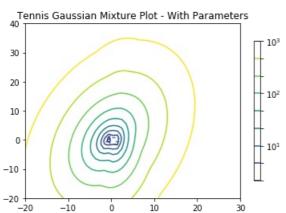
## b. Eksperimen GMM dengan parameter komponen, tipe kovarian, dan threshold

# In [78]:

```
iris_gauss = GaussianMixture(n_components=k_iris, covariance_type='spherical', tol=0.0001).fit(iris_df)
tennis_gauss = GaussianMixture(n_components=k_tennis, covariance_type='spherical', tol=0.0001).fit(tennis_df)

plot_cluster_GMM(iris_df, k_iris, "Iris Gaussian Mixture Plot - With Parameters")
plot_cluster_GMM(tennis_df, k_tennis, "Tennis Gaussian Mixture Plot - With Parameters")
```





#### c. Simpan model

#### In [54]:

```
joblib.dump(iris_gauss, 'iris_gauss.pkl')
joblib.dump(tennis_gauss, 'tennis_gauss.pkl')
print("Gaussian Mixture models saved")
```

Gaussian Mixture models saved

# 5. K-Medoids

Beberapa parameter K-Medoids:

- 1. data: data yang ingin diproses
- 2. initial\_index\_medoids: titik-titik centroid awal
- 3. tolerance: threshold untuk konvergensi (default=0.001)
- 4. ccore: true apabila ingin menggunakan library C dalam proses K-Medoids (default=True)

## In [55]:

```
def data_clusters_to_cluster(cluster_idxs):
    size = 0
    for i in range(0, len(cluster_idxs)):
        size += len(cluster_idxs[i])
    labels = [0] * size
    for i in range(0, len(cluster_idxs)):
        for j in range(0, len(cluster_idxs[i])):
            labels[cluster_idxs[i][j]] = i
    return labels
```

# a. Eksperimen K-Medoids dengan parameter initial medoid random dan toleransi tinggi

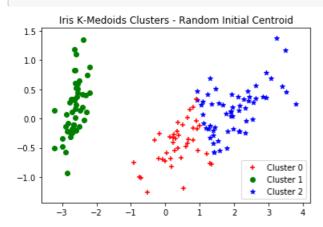
## In [91]:

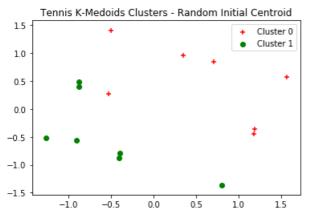
```
iris_init_medoids1 = np.random.randint(low=0, high=iris_df.shape[0], size=k_iris)
iris_kmedoids1 = kmedoids(data=iris_df.get_values(), initial_index_medoids=iris_init_medoids1,
tolerance=0.5)
iris_kmedoids1.process()

tennis_init_medoids1 = np.random.randint(low=0, high=tennis_df.shape[0], size=k_tennis)
```

```
tennis_kmedoids1 = kmedoids(data=tennis_df.get_values(),
initial_index_medoids=tennis_init_medoids1, tolerance=0.5)
tennis_kmedoids1.process()

plot_cluster(iris_df, data_clusters_to_cluster(
    iris_kmedoids1.get_clusters()), "Iris K-Medoids Clusters - Random Initial Centroid")
plot_cluster(tennis_df, data_clusters_to_cluster(
    tennis_kmedoids1.get_clusters()), "Tennis K-Medoids Clusters - Random Initial Centroid")
```





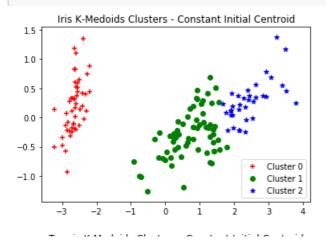
#### b. Eksperimen K-Medoids dengan initial centroid tetap dan toleransi rendah

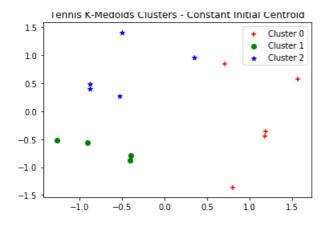
#### In [57]:

```
iris_init_medoids = [25, 75, 125]
iris_kmedoids = kmedoids(data=iris_df.get_values(), initial_index_medoids=iris_init_medoids)
iris_kmedoids.process()

tennis_init_medoids = [4, 8, 12]
tennis_kmedoids = kmedoids(data=tennis_df.get_values(), initial_index_medoids=tennis_init_medoids)
tennis_kmedoids.process()

plot_cluster(iris_df, data_clusters_to_cluster(
    iris_kmedoids.get_clusters()), "Iris K-Medoids Clusters - Constant Initial Centroid")
plot_cluster(tennis_df, data_clusters_to_cluster(
    tennis_kmedoids.get_clusters()), "Tennis K-Medoids Clusters - Constant Initial Centroid")
```





# c. Simpan model

```
In [58]:
```

```
joblib.dump(iris_kmedoids.get_cluster_encoding(), 'iris_kmedoids.pkl')
joblib.dump(tennis_kmedoids.get_cluster_encoding(), 'tennis_kmedoids.pkl')
print("K-Medoids models saved")
```

K-Medoids models saved

# 6. MST Clustering

Beberapa parameter MST:

- 1. cutoff: jumlah edge yang dihapus
- 2. cutoff\_scale: hapus edge dengan jarak melebihi skala
- 3. min\_cluster\_size: jumlah minimum data per cluster, bila lebih kecil maka data akan diassign pada background (default=1)
- 4. metric: metrik untuk menghitung jarak antar data (default=euclidean)

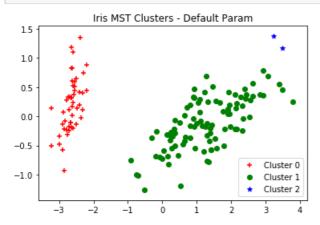
## a. Eksperimen MST dengan parameter default

```
In [59]:
```

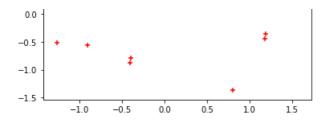
```
iris_mst1 = MSTClustering(cutoff=k_iris-1).fit(iris_df)
tennis_mst1 = MSTClustering(cutoff=k_tennis-1).fit(tennis_df)

plot_cluster(iris_df, iris_mst1.fit_predict(iris_df).tolist(), "Iris MST Clusters - Default Param")

plot_cluster(tennis_df, tennis_mst1.fit_predict(iris_df).tolist(), "Tennis MST Clusters - Default Param")
```





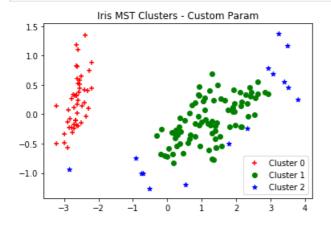


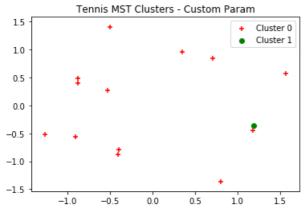
#### b. Eksperimen MST dengan parameter jumlah cluster dan metrik

#### In [60]:

```
iris_mst = MSTClustering(cutoff=k_iris-1, cutoff_scale=0.8, min_cluster_size=10, metric='manhattan'
).fit(iris_df)
tennis_mst = MSTClustering(cutoff=k_tennis-1, cutoff_scale=0.5, min_cluster_size=1, metric='manhatt
an').fit(tennis_df)

plot_cluster(iris_df, iris_mst.fit_predict(iris_df).tolist(), "Iris MST Clusters - Custom Param")
plot_cluster(tennis_df, tennis_mst.fit_predict(iris_df).tolist(), "Tennis MST Clusters - Custom Param")
"iris: Cluster"
```





## c. Simpan model

## In [99]:

```
joblib.dump(iris_mst, 'iris_mst.pkl')
joblib.dump(tennis_mst1, 'tennis_mst.pkl')
print("MST models saved")
```

MST models saved

# 7. Grid Clustering

Beberapa parameter Grid:

- 1. data: data yang ingin dicluster
- 2. levels: jumlah pemisahan 1 blok pada grid

## a. Eksperimen Grid Clustering

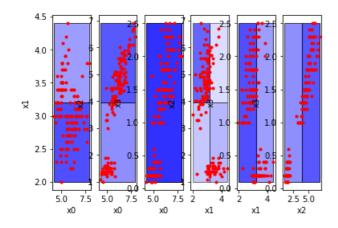
#### In [62]:

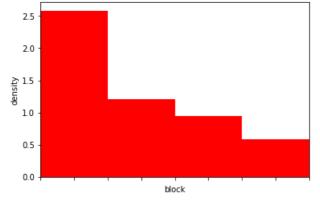
```
iris_grid = bang(data=iris_df.values.tolist(), levels=k_iris)
iris_grid.process()

tennis_grid = bang(data=tennis_df.values.tolist(), levels=k_tennis)
tennis_grid.process()

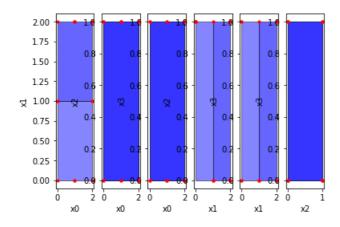
print("Iris Grid Clustering Blocks and Dendogram")
bang_visualizer.show_blocks(iris_grid.get_directory())
bang_visualizer.show_dendrogram(iris_grid.get_dendrogram())
print()
print("Tennis Grid Clustering Blocks and Dendogram")
bang_visualizer.show_blocks(tennis_grid.get_directory())
bang_visualizer.show_dendrogram(tennis_grid.get_dendrogram())
```

Iris Grid Clustering Blocks and Dendogram

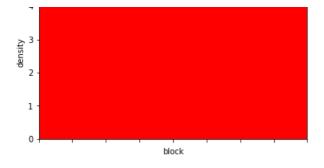




Tennis Grid Clustering Blocks and Dendogram







#### b. Simpan model

```
In [63]:
```

```
joblib.dump(iris_grid, 'iris_grid.pkl')
joblib.dump(tennis_grid, 'tennis_grid.pkl')
print("Grid Clustering models saved")
```

Grid Clustering models saved

# **Load Models**

#### In [102]:

```
iris_agglo_loaded = joblib.load('iris_agglo.pkl')
tennis_agglo_loaded = joblib.load('tennis_agglo.pkl')
iris_dbscan_loaded = joblib.load('iris_dbscan.pkl')
tennis_dbscan_loaded = joblib.load('tennis_dbscan.pkl')
iris_kmeans_loaded = joblib.load('iris_kmeans.pkl')
tennis_kmeans_loaded = joblib.load('tennis_kmeans.pkl')
iris_gauss_loaded = joblib.load('tennis_gauss.pkl')
tennis_gauss_loaded = joblib.load('tennis_gauss.pkl')
iris_kmedoids_loaded = joblib.load('iris_kmedoids.pkl')
tennis_kmedoids_loaded = joblib.load('tennis_kmedoids.pkl')
iris_mst_loaded = joblib.load('tennis_mst.pkl')
tennis_mst_loaded = joblib.load('tennis_mst.pkl')
iris_grid_loaded = joblib.load('iris_grid.pkl')
tennis_grid_loaded = joblib.load('tennis_grid.pkl')
```

# **New Instances**

Didefinisikan sebuah instans baru untuk iris dan tennis

```
In [65]:
```

```
iris_new = {
    'sepal length (cm)': [5.4],
    'sepal width (cm)': [3.7],
    'petal length (cm)': [1.5],
    'petal width (cm)': [0.2]
}
tennis_new = {'outlook': [0], 'temp': [1], 'humidity': [1], 'windy': [0]}
iris_new_df = pd.DataFrame(data=iris_new)
tennis_new_df = pd.DataFrame(data=tennis_new)

iris_new_df_c = pd.concat([iris_df, iris_new_df])
tennis_new_df_c = pd.concat([tennis_df, tennis_new_df])
```

# **Assigment Cluster**

# 1. Agglomerative Clustering

# In [66]: iris\_predict\_agglo = iris\_agglo\_loaded.fit\_predict(iris\_new\_df\_c) [len(iris\_df)] print("iris: Cluster", iris\_predict\_agglo) tennis\_predict\_agglo = tennis\_agglo\_loaded.fit\_predict(tennis\_new\_df\_c) [len(tennis\_df)] print("tennis: Cluster", tennis\_predict\_agglo) iris: Cluster 1 tennis: Cluster 0

# 2. DBSCAN

```
In [67]:
```

```
iris_predict_dbscan = iris_dbscan_loaded.fit_predict(iris_new_df_c)[len(iris_df)]
print("iris: Cluster", iris_predict_dbscan)
tennis_predict_dbscan = tennis_dbscan_loaded.fit_predict(tennis_new_df_c)[len(tennis_df)]
print("tennis: Cluster", tennis_predict_dbscan)

iris: Cluster 0
tennis: Cluster 1
```

# 3. K-Means

```
In [68]:
```

```
iris_predict_kmeans = iris_kmeans_loaded.predict(iris_new_df)[0]
print("iris: Cluster", iris_predict_kmeans)
tennis_predict_kmeans = tennis_kmeans_loaded.predict(tennis_new_df)[0]
print("tennis: Cluster", tennis_predict_kmeans)

iris: Cluster 1
tennis: Cluster 1
```

# 4. Gaussian Mixture

```
In [69]:
```

```
iris_predict_gauss = iris_gauss_loaded.predict(iris_new_df)[0]
print("iris: Cluster", iris_predict_gauss)
tennis_predict_gauss = tennis_gauss_loaded.predict(tennis_new_df)[0]
print("tennis: Cluster", tennis_predict_gauss)

iris: Cluster 1
tennis: Cluster 1
```

# 5. K-Medoids

```
In [70]:
```

```
iris_predict_kmedoids = kmedoids(data=iris_new_df_c.get_values(),
initial_index_medoids=iris_init_medoids)
iris_predict_kmedoids.process()
print("iris: Cluster", data_clusters_to_cluster(iris_predict_kmedoids.get_clusters())[len(iris_df)])
tennis_predict_kmedoids = kmedoids(data=tennis_new_df_c.get_values(),
initial_index_medoids=tennis_init_medoids)
tennis_predict_kmedoids.process()
print("tennis: Cluster", data_clusters_to_cluster(tennis_predict_kmedoids.get_clusters())
[len(tennis_df)])
```

iris: Cluster 0
tennis: Cluster 2

# 6. MST Clustering

```
In [104]:
```

```
iris_predict_mst = iris_mst_loaded.fit_predict(iris_new_df_c)[len(iris_df)]
print("iris: Cluster", iris_predict_mst)
tennis_predict_mst = tennis_mst_loaded.fit_predict(tennis_new_df_c)[len(tennis_df)]
print("tennis: Cluster", tennis_predict_mst)

iris: Cluster 0
tennis: Cluster 1
```

# 7. Grid Clustering

```
In [72]:
```

```
iris_predict_grid = bang(data=iris_new_df_c.values.tolist(), levels=k_iris)
iris_predict_grid.process()
print("iris: Cluster", data_clusters_to_cluster(iris_predict_grid.get_clusters())[len(iris_df)])
tennis_predict_grid = bang(data=tennis_new_df_c.values.tolist(), levels=k_tennis)
tennis_predict_grid.process()
print("tennis: Cluster", data_clusters_to_cluster(tennis_predict_grid.get_clusters())
[len(tennis_df)])
```

iris: Cluster 0
tennis: Cluster 0