# Clustering\_excercise

May 29, 2019

### 1 Clustering

The use of watermark is optional. You can install this IPython extension via "pip install watermark". For more information, please see: https://github.com/rasbt/watermark.

### 2 Grouping objects by similarity using k-means

### 2.1 K-means clustering using scikit-learn

```
In [ ]: from sklearn.datasets import make_blobs
        X, y = make_blobs(n_samples=150,
                          n_features=2,
                          centers=3,
                          cluster_std=0.5,
                          shuffle=True,
                          random_state=0)
In [ ]: print(X,y)
In [ ]: import matplotlib.pyplot as plt
        plt.scatter(X[:, 0], X[:, 1],
                    c='white', marker='o', edgecolor='black', s=50)
        plt.grid()
        plt.tight_layout()
        #plt.savefig('images/11_01.png', dpi=300)
        plt.show()
In [ ]: from sklearn.cluster import KMeans
```

```
km = KMeans(n_clusters=3,
                    init='random',
                    max_iter = 300,
                    random_state=0)
        y_km = km.fit_predict(X)
In [ ]: print(y_km)
In [ ]: X[y_km == 0, 0]
In [ ]: X[y_km == 1, 0]
In []: print(km.cluster_centers_)
In []: plt.scatter(X[y_km == 0, 0],
                    X[y_{km} == 0, 1],
                    s=50, c='lightgreen',
                    marker='s', edgecolor='black',
                    label='cluster 1')
        plt.scatter(X[y_km == 1, 0],
                    X[y_{km} == 1, 1],
                    s=50, c='orange',
                    marker='o', edgecolor='black',
                    label='cluster 2')
        plt.scatter(X[y_km == 2, 0],
                    X[y_{km} == 2, 1],
                    s=50, c='lightblue',
                    marker='v', edgecolor='black',
                    label='cluster 3')
        plt.scatter(km.cluster_centers_[:, 0],
                    km.cluster_centers_[:, 1],
                    s=250, marker='*',
                    c='red', edgecolor='black',
                    label='centroids')
        plt.legend(scatterpoints=1)
        plt.grid()
        plt.tight_layout()
        #plt.savefig('images/11_02.png', dpi=300)
        plt.show()
```

### 2.2 Using the elbow method to find the optimal number of clusters

```
for i in range(1, 11):
            km = KMeans(n_clusters=i,
                        init='random',
                        random_state=0)
            km.fit(X)
            distortions.append(km.inertia_)
        plt.plot(range(1, 11), distortions, marker='o')
        plt.xlabel('Number of clusters')
        plt.ylabel('Distortion')
        plt.tight_layout()
        #plt.savefig('images/11_03.png', dpi=300)
        plt.show()
In [ ]: from sklearn.cluster import KMeans
        km = KMeans(n_clusters=3,
                    init='k-means++',
                    \max iter = 300,
                    random_state=0)
        y_km = km.fit_predict(X)
In []: plt.scatter(X[y_km == 0, 0],
                    X[y_{km} == 0, 1],
                    s=50, c='lightgreen',
                    marker='s', edgecolor='black',
                    label='cluster 1')
        plt.scatter(X[y_km == 1, 0],
                    X[y_{km} == 1, 1],
                    s=50, c='orange',
                    marker='o', edgecolor='black',
                    label='cluster 2')
        plt.scatter(X[y_km == 2, 0],
                    X[y_{km} == 2, 1],
                    s=50, c='lightblue',
                    marker='v', edgecolor='black',
                    label='cluster 3')
        plt.scatter(km.cluster_centers_[:, 0],
                    km.cluster_centers_[:, 1],
                    s=250, marker='*',
                    c='red', edgecolor='black',
                    label='centroids')
        plt.legend(scatterpoints=1)
        plt.grid()
        plt.tight_layout()
        #plt.savefig('images/11_02.png', dpi=300)
       plt.show()
In []: # kmeans++ - , kmeans
```

```
# inertia_ : SSE
        distortions = []
        for i in range(1, 11):
            km = KMeans(n_clusters=i,
                        init='k-means++',
                        random state=0)
            km.fit(X)
            distortions.append(km.inertia_)
        plt.plot(range(1, 11), distortions, marker='o')
        plt.xlabel('Number of clusters')
        plt.ylabel('Distortion')
        plt.tight_layout()
        #plt.savefig('images/11_03.png', dpi=300)
        plt.show()
2.3 silhouette
  • a -
  • b -
  • silhouette = (b-a)/max(b,a)
In []: import numpy as np
        from matplotlib import cm
        from sklearn.metrics import silhouette_samples
        km = KMeans(n_clusters=3,
                    init='k-means++',
                    random state=0)
        y_km = km.fit_predict(X)
        print(X[:5])
        print(y_km[:5])
        cluster_labels = np.unique(y_km)
        print(cluster_labels)
        n_clusters = cluster_labels.shape[0]
        print(n_clusters)
        silhouette_vals = silhouette_samples(X, y_km, metric='euclidean')
        print(silhouette_vals[:5])
        print(len(silhouette_vals))
        y_ax_lower, y_ax_upper = 0, 0
        yticks = []
In [ ]: np.mean(silhouette_samples(X, y_km, metric='euclidean'))
In [ ]: plt.plot(range(len(silhouette_vals)), sorted(silhouette_vals))
```

```
In []: clist = ['g','b','y']
        for i, c in enumerate(cluster_labels):
            c_silhouette_vals = silhouette_vals[y_km == c]
            c_silhouette_vals.sort()
            y ax upper += len(c silhouette vals)
              color = cm.jet(float(i) / n_clusters)
            plt.barh(range(y_ax_lower, y_ax_upper), c_silhouette_vals, height=1.0,
                     edgecolor='none', color=clist[i])
            yticks.append((y_ax_lower + y_ax_upper) / 2.)
            y_ax_lower += len(c_silhouette_vals)
        silhouette_avg = np.mean(silhouette_vals)
        plt.axvline(silhouette_avg, color="red", linestyle="--")
        plt.yticks(yticks, cluster_labels + 1)
        plt.ylabel('Cluster')
        plt.xlabel('Silhouette coefficient')
        plt.tight layout()
        #plt.savefig('images/11_04.png', dpi=300)
        plt.show()
  Comparison to "bad" clustering:
In [ ]: km = KMeans(n_clusters=2,
                    init='k-means++',
                    n init=10,
                    max_iter=300,
                    tol=1e-04.
                    random_state=0)
        y_km = km.fit_predict(X)
        plt.scatter(X[y_km == 0, 0],
                    X[y_{km} == 0, 1],
                    s = 50,
                    c='lightgreen',
                    edgecolor='black',
                    marker='s',
                    label='cluster 1')
        plt.scatter(X[y_km == 1, 0],
                    X[y_{km} == 1, 1],
                    s = 50,
                    c='orange',
                    edgecolor='black',
                    marker='o',
                    label='cluster 2')
```

```
plt.scatter(km.cluster_centers_[:, 0], km.cluster_centers_[:, 1],
                    s=250, marker='*', c='red')#, label='centroids')
        plt.legend()
        plt.grid()
        plt.tight_layout()
        #plt.savefig('images/11_05.png', dpi=300)
        plt.show()
In []: cluster_labels = np.unique(y_km)
        n_clusters = cluster_labels.shape[0]
        silhouette_vals = silhouette_samples(X, y_km, metric='euclidean')
        y_ax_lower, y_ax_upper = 0, 0
        yticks = []
        colorlist=['g','b']
        for i, c in enumerate(cluster_labels):
            c_silhouette_vals = silhouette_vals[y_km == c]
            c_silhouette_vals.sort()
            y_ax_upper += len(c_silhouette_vals)
              color = cm.jet(float(i) / n_clusters)b
            plt.barh(range(y_ax_lower, y_ax_upper), c_silhouette_vals, height=1.0,
                     edgecolor='none', color=colorlist[i])
            yticks.append((y_ax_lower + y_ax_upper) / 2.)
            y_ax_lower += len(c_silhouette_vals)
        silhouette_avg = np.mean(silhouette_vals)
        plt.axvline(silhouette_avg, color="red", linestyle="--")
        plt.yticks(yticks, cluster_labels + 1)
        plt.ylabel('Cluster')
        plt.xlabel('Silhouette coefficient')
        plt.tight_layout()
        #plt.savefig('images/11_06.png', dpi=300)
        plt.show()
In [ ]: \# \iff silhouette .
In [ ]:
```

## 3 Hierachical Clustering - Agglomerative Clustering

1. 2. 3. 4. . 5.

• ward - .

```
average -
  • complete -
In []: import mglearn
In [ ]: import matplotlib.font_manager as fm
        path = 'C:\\Windows\\Fonts\\ALGER.TTF'
        fontprop = fm.FontProperties(fname=path, size=18)
        mglearn.plots.plot_agglomerative_algorithm()
In [ ]: from sklearn.cluster import AgglomerativeClustering
        from sklearn.datasets import make blobs
        import matplotlib.pyplot as plt
        %matplotlib inline
        # import matplotlib.font_manager as fm
        \# path = 'C:\\Windows\\Fonts\\ALGER.TTF'
        # fontprop = fm.FontProperties(fname=path, size=18)
        X, y = make_blobs(random_state=1)
        agg = AgglomerativeClustering(n_clusters=3)
        assignment = agg.fit_predict(X)
        print(assignment[:20])
        mglearn.discrete_scatter(X[:, 0],X[:,1], assignment)
       plt.legend([' 0', ' 1', ' 2'], loc='best')
       plt.xlabel(' 0')
       plt.ylabel(' 1')
In []: mglearn.plots.plot_agglomerative()
In []: from scipy.cluster.hierarchy import dendrogram, ward
        linkage_array = ward(X) #
        row_dendr = dendrogram(linkage_array, # dendrogram
                               labels=y,
       plt.tight_layout()
       plt.ylabel('Euclidean distance')
       plt.show()
In []:
In []:
```

#### 3.1 Grouping clusters in bottom-up fashion

#### 3.2 Applying agglomerative clustering via scikit-learn

## 4 Locating regions of high density via DBSCAN

#### 4.1

```
• - e
• (MinPts) e
• -e (MinPts) , e
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• ,e ()
• - , ,
• -()
• random , ,

In []: Image(filename='images/11_13.png', width=500)
```

```
X, y = make_moons(n_samples=200, noise=0.05, random_state=0)
plt.scatter(X[:, 0], X[:, 1])
plt.tight_layout()
#plt.savefig('images/11_14.png', dpi=300)
plt.show()

In []: # <> clustering algorithm , .

In []:
K-means and hierarchical clustering:

In []:
Density-based clustering:

5 Summary
...

Readers may ignore the next cell.

In []: ! python ../.convert_notebook_to_script.py --input ch11.ipynb --output ch11.py
```

In [ ]: from sklearn.datasets import make\_moons