Experiment-11

AIM: Apply EM algorithm to cluster a set of data stored in a. CSV file. Use the same data set for clustering using k-Means algorithm. Compare the results of the set two algorithms and comment on the quality of clustering. You can add Java/Python ML library classes/API in the program.

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import make blobs
from sklearn.mixture import GaussianMixture
from matplotlib.patches import Ellipse
X, y true = make blobs(n samples=100, centers=4, cluster std=0.60,
random_state=0)
X = X[:, ::-1]
gmm = GaussianMixture(n components=4).fit(X)
labels = gmm.predict(X)
plt.scatter(X[:, 0], X[:, 1], c=labels, s=40, cmap='viridis')
probs = gmm.predict_proba(X)
print(probs[:5].round(3))
size = 50 * probs.max(1) ** 2
plt.scatter(X[:, 0], X[:, 1], c=labels, cmap='viridis', s=size)
def draw ellipse(position, covariance, ax=None, **kwargs):
  ax = ax or plt.gca()
  if covariance.shape == (2, 2):
    U, s, Vt = np.linalg.svd(covariance)
```

```
angle = np.degrees(np.arctan2(U[1, 0], U[0, 0]))
    width, height = 2 * np.sqrt(s)
  else:
    angle = 0
    width, height = 2 * np.sqrt(covariance)
  for nsig in range(1, 4):
    ax.add patch(Ellipse(position, nsig * width, nsig * height, angle, **kwargs))
def plot_gmm(gmm, X, label=True, ax=None):
  ax = ax or plt.gca()
  if label:
    labels = gmm.fit(X).predict(X)
    ax.scatter(X[:, 0], X[:, 1], c=labels, s=40, cmap='viridis', zorder=2)
  else:
    ax.scatter(X[:, 0], X[:, 1], s=40, zorder=2)
  ax.axis('equal')
  w_factor = 0.2 / gmm.weights_.max()
  for pos, covar, w in zip(gmm.means_, gmm.covariances_, gmm.weights_):
    draw ellipse(pos, covar, alpha=w * w factor)
gmm = GaussianMixture(n components=4, random state=42)
plot_gmm(gmm, X)
gmm = GaussianMixture(n_components=4, covariance_type='full', random_state=42)
plot_gmm(gmm, X)
plt.show()
```

Output:

[[0. 0. 1. 0.] [0. 1. 0. 0.] [0. 0. 1. 0.] [0. 0. 1. 0.]

