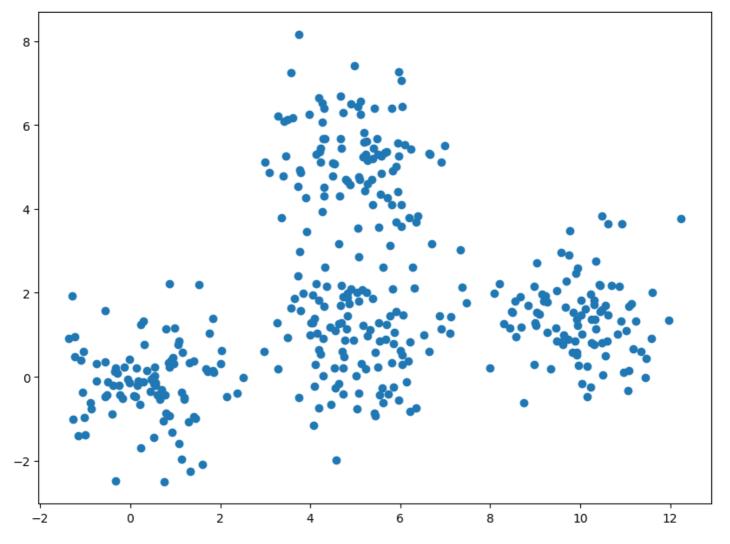
In [39]:

```
import numpy as np
import matplotlib.pyplot as plt

## Data generation
# write your code here
a=np.random.multivariate_normal([0.5,0], [[1, 0],[0, 1]], 100)
b=np.random.multivariate_normal([5,5], [[1, 0],[0, 1]], 100)
c=np.random.multivariate_normal([5,1], [[1, 0],[0, 1]], 100)
d=np.random.multivariate_normal([10,1.5], [[1, 0],[0, 1]], 100)
X=np.concatenate((a,b,c,d), axis=0)
plt.scatter(X[:,0],X[:,1])

n=X.shape[1]
n_iter=10
```



In [40]:

```
K=4
import random

# creating an empty centroid array
centroids=np.array([]).reshape(n,0)

# creating 5 random centroids
for k in range(K):
    centroids=np.c_[centroids,X[random.randint(0,m-1)]]

print(centroids)
```

In [41]: output={} # creating an empty array euclid=np.array([]).reshape(m,0) # finding distance between for each centroid for k in range(K): dist=np.sum((X-centroids[:,k])**2,axis=1) euclid=np.c_[euclid,dist] # storing the minimum value we have computed minimum=np.argmin(euclid,axis=1)+1

In [42]:

```
# computing the mean of separated clusters
cent={}
for k in range(K):
    cent[k+1]=np.array([]).reshape(2,0)

# assigning of clusters to points
for k in range(m):
    cent[minimum[k]]=np.c_[cent[minimum[k]],X[k]]
for k in range(K):
    cent[k+1]=cent[k+1].T

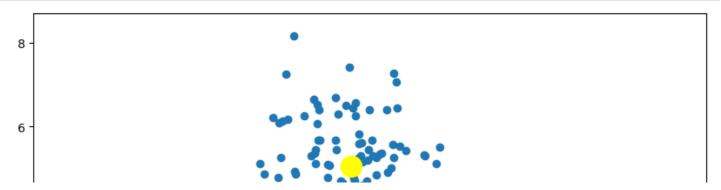
# computing mean and updating it
for k in range(K):
    centroids[:,k]=np.mean(cent[k+1],axis=0)
```

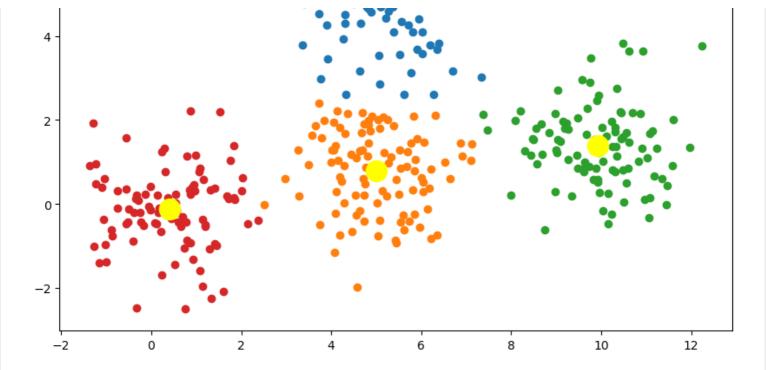
In [43]:

```
# repeating the above steps again and again
for i in range(n iter):
   euclid=np.array([]).reshape(m,0)
   for k in range(K):
        dist=np.sum((X-centroids[:,k])**2,axis=1)
       euclid=np.c [euclid, dist]
   C=np.argmin(euclid,axis=1)+1
   cent={}
   for k in range(K):
       cent[k+1]=np.array([]).reshape(2,0)
   for k in range(m):
       cent[C[k]]=np.c_[cent[C[k]],X[k]]
   for k in range(K):
       cent[k+1] = cent[k+1].T
   for k in range(K):
       centroids[:,k]=np.mean(cent[k+1],axis=0)
   final=cent
```

In [44]:

```
for k in range(K):
    plt.scatter(final[k+1][:,0],final[k+1][:,1])
plt.scatter(centroids[0,:],centroids[1,:],s=300,c='yellow')
plt.rcParams.update({'figure.figsize':(10,7.5), 'figure.dpi':100})
plt.show()
```



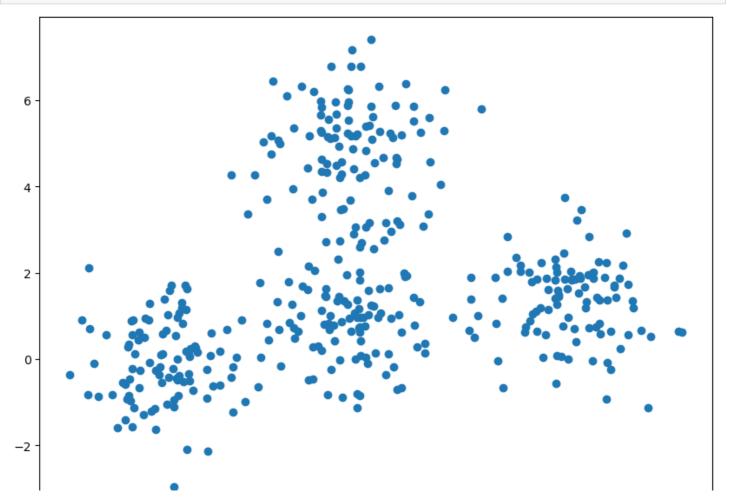


In [45]:

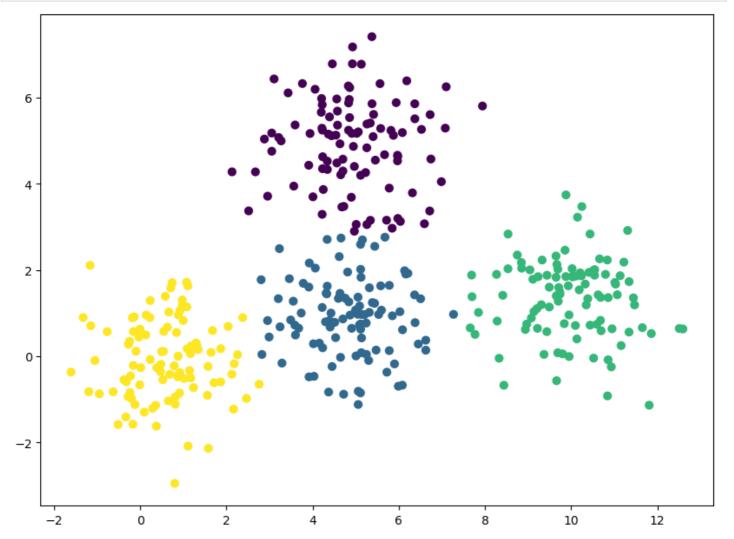
```
import numpy as np
import matplotlib.pyplot as plt

## Data generation
# write your code here
a=np.random.multivariate_normal([0.5,0], [[1, 0],[0, 1]], 100)
b=np.random.multivariate_normal([5,5], [[1, 0],[0, 1]], 100)
c=np.random.multivariate_normal([5,1], [[1, 0],[0, 1]], 100)
d=np.random.multivariate_normal([10,1.5], [[1, 0],[0, 1]], 100)
X=np.concatenate((a,b,c,d), axis=0)
plt.scatter(X[:,0],X[:,1])

n=X.shape[1]
n_iter=10
```



In [47]:



In [48]:

```
from sklearn.cluster import KMeans
def calculate mean covariance(X, prediction):
    d = X.shape[1]
    labels = np.unique(prediction)
    initial means = np.zeros((C, d))
    initial_cov = np.zeros((C, d, d))
    initial pi = np.zeros(C)
    counter=0
    for label in sorted(labels):
        ids = np.where(prediction == label) # returns indices
        initial pi[counter] = len(ids[0]) / X.shape[0]
        initial means[counter,:] = np.mean(X[ids], axis = 0)
        de meaned = X[ids] - initial means[counter,:]
        Nk = X[ids].shape[0]
        initial_cov[counter,:, :] = np.dot(initial_pi[counter] * de_meaned.T, de_meaned)
/ Nk
        counter+=1
    assert np.sum(initial_pi) == 1
    return (initial_means, initial_cov, initial_pi)
```

```
n_clusters = 3
kmeans = KMeans(n_clusters= n_clusters, max_iter=500, algorithm = 'auto')
fitted = kmeans.fit(X)
prediction = kmeans.predict(X)

m, c, pi = calculate_mean_covariance(X, prediction)
```

In [49]:

```
from scipy.stats import multivariate normal as mvn
class GMM:
    """ Gaussian Mixture Model
   Parameters
    _____
       k: int , number of gaussian distributions
       seed: int, will be randomly set if None
       max iter: int, number of iterations to run algorithm, default: 200
   Attributes
      centroids: array, k, number features
      cluster labels: label for each data point
   def
        __init___(self, C, n_runs):
       self.C = C # number of Guassians/clusters
       self.n runs = n runs
   def get params(self):
       return (self.mu, self.pi, self.sigma)
   def calculate mean covariance(self, X, prediction):
        """Calculate means and covariance of different
           clusters from k-means prediction
       Parameters:
       prediction: cluster labels from k-means
       X: N*d numpy array data points
       Returns:
       intial means: for E-step of EM algorithm
       intial cov: for E-step of EM algorithm
       d = X.shape[1]
       labels = np.unique(prediction)
       self.initial_means = np.zeros((self.C, d))
       self.initial cov = np.zeros((self.C, d, d))
       self.initial pi = np.zeros(self.C)
       counter=0
       for label in labels:
            ids = np.where(prediction == label) # returns indices
            self.initial pi[counter] = len(ids[0]) / X.shape[0]
            self.initial_means[counter,:] = np.mean(X[ids], axis = 0)
            de meaned = X[ids] - self.initial means[counter,:]
            Nk = X[ids].shape[0] # number of data points in current gaussian
            self.initial_cov[counter,:, :] = np.dot(self.initial_pi[counter] * de_meaned
.T, de meaned) / Nk
```

```
counter+=1
        assert np.sum(self.initial pi) == 1
        return (self.initial means, self.initial cov, self.initial pi)
    def initialise parameters(self, X):
        """Implement k-means to find starting
            parameter values.
            https://datascience.stackexchange.com/questions/11487/how-do-i-obtain-the-wei
ght-and-variance-of-a-k-means-cluster
        Parameters:
        X: numpy array of data points
        Returns:
        tuple containing initial means and covariance
       initial means: numpy array: (C*d)
       initial cov: numpy array: (C, d*d)
        n clusters = self.C
        kmeans = KMeans(n clusters= n clusters, init="k-means++", max iter=500, algorith
m = 'auto')
        fitted = kmeans.fit(X)
        prediction = kmeans.predict(X)
        self. initial means, self. initial cov, self. initial pi = self.calculate mean c
ovariance(X, prediction)
        return (self. initial means, self. initial cov, self. initial pi)
    def _e_step(self, X, pi, mu, sigma):
        """Performs E-step on GMM model
        Parameters:
       X: (N x d), data points, m: no of features
       pi: (C), weights of mixture components
       mu: (C x d), mixture component means
        sigma: (C \times d \times d), mixture component covariance matrices
        Returns:
        gamma: (N x C), probabilities of clusters for objects
        N = X.shape[0]
        self.gamma = np.zeros((N, self.C))
        const c = np.zeros(self.C)
        self.mu = self.mu if self. initial means is None else self. initial means
        self.pi = self.pi if self. initial pi is None else self. initial pi
        self.sigma = self.sigma if self. initial cov is None else self. initial cov
        for c in range(self.C):
            # Posterior Distribution using Bayes Rule
            self.gamma[:,c] = self.pi[c] * mvn.pdf(X, self.mu[c,:], self.sigma[c])
        # normalize across columns to make a valid probability
        gamma norm = np.sum(self.gamma, axis=1)[:,np.newaxis]
        self.gamma /= gamma norm
```

```
return self.gamma
   def _m_step(self, X, gamma):
        """Performs M-step of the GMM
       We need to update our priors, our means
        and our covariance matrix.
       Parameters:
       X: (N \times d), data
        gamma: (N x C), posterior distribution of lower bound
       Returns:
        _____
       pi: (C)
       mu: (C \times d)
        sigma: (C x d x d)
       N = X.shape[0] # number of objects
       C = self.gamma.shape[1] # number of clusters
       d = X.shape[1] # dimension of each object
        # responsibilities for each gaussian
        self.pi = np.mean(self.gamma, axis = 0)
        self.mu = np.dot(self.gamma.T, X) / np.sum(self.gamma, axis = 0)[:,np.newaxis]
       for c in range(C):
            x = X - self.mu[c, :] # (N x d)
            gamma diag = np.diag(self.gamma[:,c])
            x mu = np.matrix(x)
            gamma diag = np.matrix(gamma diag)
            sigma c = x.T * gamma diag * x
            self.sigma[c,:,:]=(sigma c) / np.sum(self.gamma, axis = 0)[:,np.newaxis][c]
       return self.pi, self.mu, self.sigma
   def compute loss function(self, X, pi, mu, sigma):
        """Computes lower bound loss function
       Parameters:
       X: (N \times d), data
       Returns:
        pi: (C)
       mu: (C \times d)
        sigma: (C x d x d)
       N = X.shape[0]
       C = self.gamma.shape[1]
       self.loss = np.zeros((N, C))
       for c in range(C):
            dist = mvn(self.mu[c], self.sigma[c],allow_singular=True)
            self.loss[:,c] = self.gamma[:,c] * (np.log(self.pi[c]+0.00001)+dist.logpdf(X
)-np.log(self.gamma[:,c]+0.000001))
       self.loss = np.sum(self.loss)
       return self.loss
   def fit(self, X):
        """Compute the E-step and M-step and
            Calculates the lowerbound
       Parameters:
```

```
X: (N \times d), data
   Returns:
    instance of GMM
   d = X.shape[1]
   self.mu, self.sigma, self.pi = self._initialise_parameters(X)
        for run in range(self.n runs):
            self.gamma = self. e step(X, self.mu, self.pi, self.sigma)
            self.pi, self.mu, self.sigma = self. m step(X, self.gamma)
            loss = self._compute_loss_function(X, self.pi, self.mu, self.sigma)
            if run % 10 == 0:
                print("Iteration: %d Loss: %0.6f" %(run, loss))
   except Exception as e:
       print(e)
   return self
def predict(self, X):
    """Returns predicted labels using Bayes Rule to
    Calculate the posterior distribution
   Parameters:
   X: ?*d numpy array
   Returns:
    labels: predicted cluster based on
   highest responsibility gamma.
   labels = np.zeros((X.shape[0], self.C))
   for c in range(self.C):
       labels [:,c] = self.pi[c] * mvn.pdf(X, self.mu[c,:], self.sigma[c])
   labels = labels .argmax(1)
   return labels
def predict proba(self, X):
    """Returns predicted labels
    Parameters:
   X: N*d numpy array
   Returns:
    _____
    labels: predicted cluster based on
   highest responsibility gamma.
   post proba = np.zeros((X.shape[0], self.C))
   for c in range(self.C):
        # Posterior Distribution using Bayes Rule, try and vectorise
        post_proba[:,c] = self.pi[c] * mvn.pdf(X, self.mu[c,:], self.sigma[c])
   return post proba
```

In [50]:

```
model = GMM(4, n_runs = 100)

fitted_values = model.fit(X)

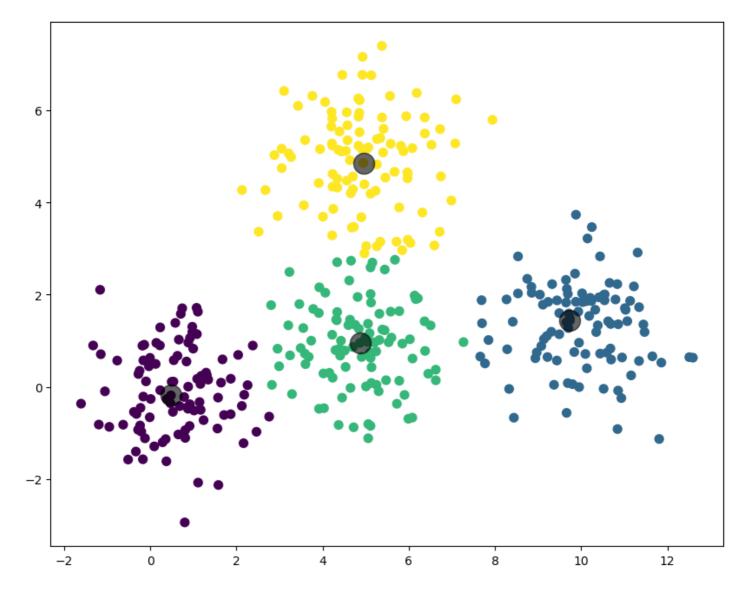
predicted_values = model.predict(X)

# compute centers as point of highest density of distribution
centers = np.zeros((4,2))
for i in range(model.C):
    density = mvn(cov=model.sigma[i], mean=model.mu[i]).logpdf(X)
    centers[i, :] = X[np.argmax(density)]

plt.figure(figsize = (10,8))
plt.scatter(X[:, 0], X[:, 1], c=predicted_values , s=50, cmap='viridis')

plt.scatter(centers[:, 0], centers[:, 1], c='black', s=300, alpha=0.6);
```

Iteration: 0 Loss: -1660.501398
Iteration: 10 Loss: -1653.847525
Iteration: 20 Loss: -1653.847330
Iteration: 30 Loss: -1653.847329
Iteration: 40 Loss: -1653.847329
Iteration: 50 Loss: -1653.847329
Iteration: 60 Loss: -1653.847329
Iteration: 70 Loss: -1653.847329
Iteration: 80 Loss: -1653.847329
Iteration: 90 Loss: -1653.847329



In [51]:

class FCM:

```
def _init_(self, n_clusters=10, max_iter=150, m=2, error=1e-5, random_state=42):
   assert m > 1
   self.u, self.centers = None, None
   self.n clusters = n clusters
   self.max iter = max iter
   self.m = m
   self.error = error
   self.random state = random state
def fit(self, X):
   self.n samples = X.shape[0]
   r = np.random.RandomState(self.random state)
   u = r.rand(self.n samples, self.n clusters)
   u = u / np.tile(u.sum(axis=1)[np.newaxis].T, self.n clusters)
    r = np.random.RandomState(self.random state)
    self.u = r.rand(self.n samples, self.n_clusters)
    self.u = self.u / np.tile(self.u.sum(axis=1)[np.newaxis].T, self.n clusters)
   for iteration in range(self.max_iter):
        u old = self.u.copy()
        self.centers = self.next centers(X)
        self.u = self. predict(X)
        selfClusterOut = self.predict(X)
        centers = self.centers
        for i in range(nPoints):
            plt.scatter(x[i], y[i], c = plotColor[selfClusterOut[i]], s = 10)
        for i in range(self.n clusters):
            plt.scatter(centers[i][0], centers[i][1], c = 'black', marker = 'X')
        plt.show()
        # Stopping rule
        if norm(self.u - u_old) < self.error:</pre>
           break
def next_centers(self, X):
   um = self.u ** self.m
   return (X.T @ um / np.sum(um, axis=0)).T
def predict(self, X):
   power = float(2 / (self.m - 1))
   temp = cdist(X, self.centers) ** power
   denominator = temp.reshape((X.shape[0], 1, -1)).repeat(temp.shape[-1], axis=1)
   denominator = temp[:, :, np.newaxis] / denominator
   return 1 / denominator .sum(2)
def predict(self, X):
    if len(X.shape) == 1:
        X = np.expand dims(X, axis=0)
   u = self. predict(X)
   return np.argmax(u, axis=-1)
```

In [53]:

```
from google.colab import drive
drive.mount('/content/drive')
```

```
X, y = make_circles(n_samples=750, factor=0.3, noise=0.1)
X = StandardScaler().fit_transform(X)
y_pred = DBSCAN(eps=0.3, min_samples=10).fit_predict(X)
y_kmeans = KMeans(2).fit_predict(X)
plt.scatter(X[:,0], X[:,1], c=y pred)
plt.show()
plt.scatter(X[:,0], X[:,1], c=y kmeans)
plt.show()
print('Number of clusters: {}'.format(len(set(y pred[np.where(y pred != -1)]))))
print('Homogeneity: {}'.format(metrics.homogeneity score(y, y pred)))
print('Completeness: {}'.format(metrics.completeness score(y, y pred)))
______
NameError
                                       Traceback (most recent call last)
<ipython-input-54-c6e26e7f8c07> in <module>
----> 1 X, y = make circles(n samples=750, factor=0.3, noise=0.1)
     2 X = StandardScaler().fit_transform(X)
     3 y pred = DBSCAN(eps=0.3, min samples=10).fit predict(X)
     4 y_kmeans = KMeans(2).fit_predict(X)
NameError: name 'make_circles' is not defined
In [ ]:
```