

Clustering by Kmeans and GMM

September 28, 2023

```
[159]: import pandas as pd
import numpy as np
from sklearn import datasets

np.random.seed(254)

X, y = datasets.make_classification(1000, n_features=2, n_informative=2,
    n_redundant=0, n_repeated=0, n_classes=3,
    n_clusters_per_class=1)

df= pd.DataFrame({"feature_1": X[:,0], "feature_2": X[:,1], "category":y},
    columns=["feature_1", "feature_2", "category"])

df.shape
```

```
[159]: (1000, 3)
```

```
[160]: df.head()
```

```
[160]:   feature_1  feature_2  category
0    1.316455    1.906756         0
1    0.499191   -1.668219         2
2    0.937946    0.834301         0
3    0.930180    0.648558         0
4    0.309252   -1.365109         2
```

```
[161]: df.describe()
```

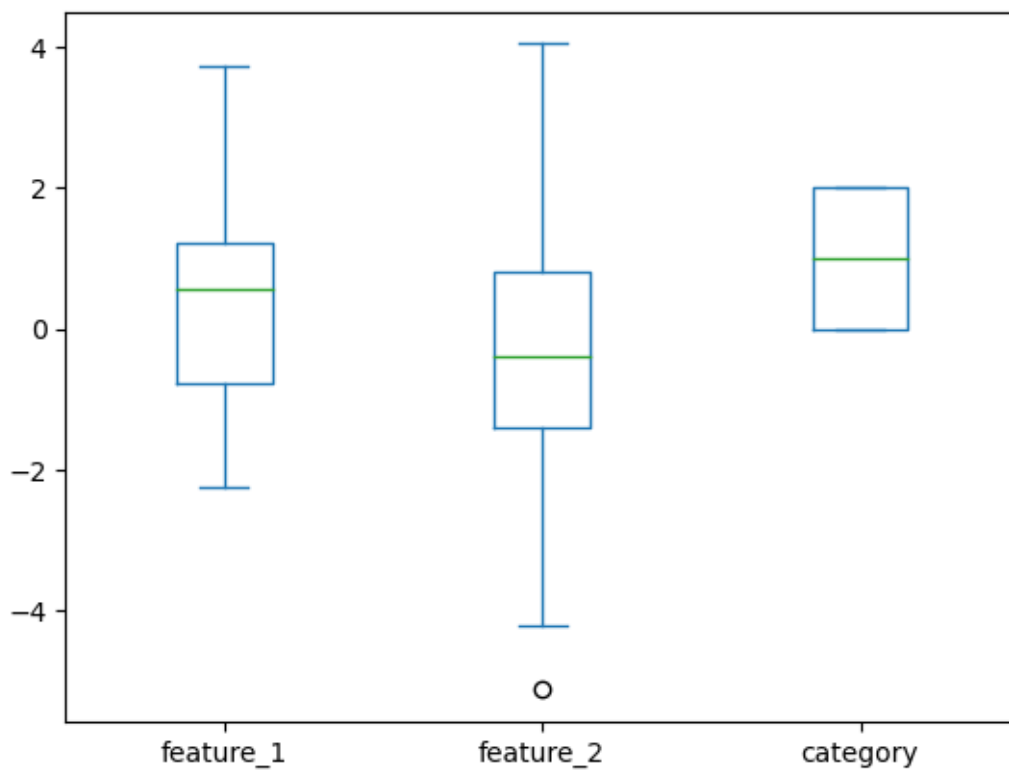
```
[161]:
```

	feature_1	feature_2	category
count	1000.000000	1000.000000	1000.000000
mean	0.332238	-0.299943	0.998000
std	1.142013	1.516528	0.818942
min	-2.234235	-5.113364	0.000000
25%	-0.779555	-1.399917	0.000000
50%	0.568081	-0.391351	1.000000
75%	1.221679	0.795395	2.000000
max	3.724172	4.034715	2.000000

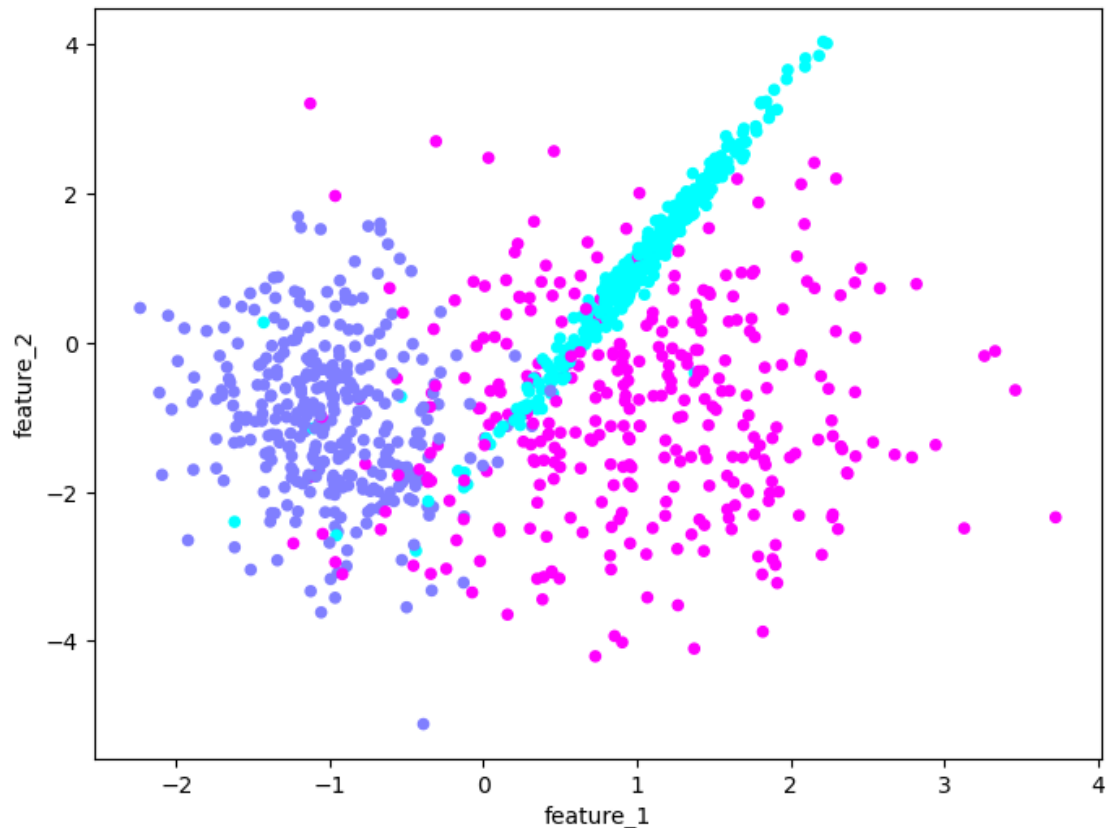
```
[162]: df.category.value_counts()
```

```
[162]: 0    336  
      2    334  
      1    330  
      Name: category, dtype: int64
```

```
[163]: import matplotlib.pyplot as plt  
      import matplotlib as pylab  
  
      df.plot.box()  
      plt.show()
```



```
[164]: df.plot.scatter("feature_1", "feature_2", c="category", cmap=pylab.cm.cool,  
      ↪ figsize=(8,6), colorbar=False)  
      plt.show()
```



```
[165]: from sklearn import preprocessing
```

```
[166]: scaler = preprocessing.StandardScaler()
```

```
[167]: scaler.fit(df[["feature_1", "feature_2"]])
```

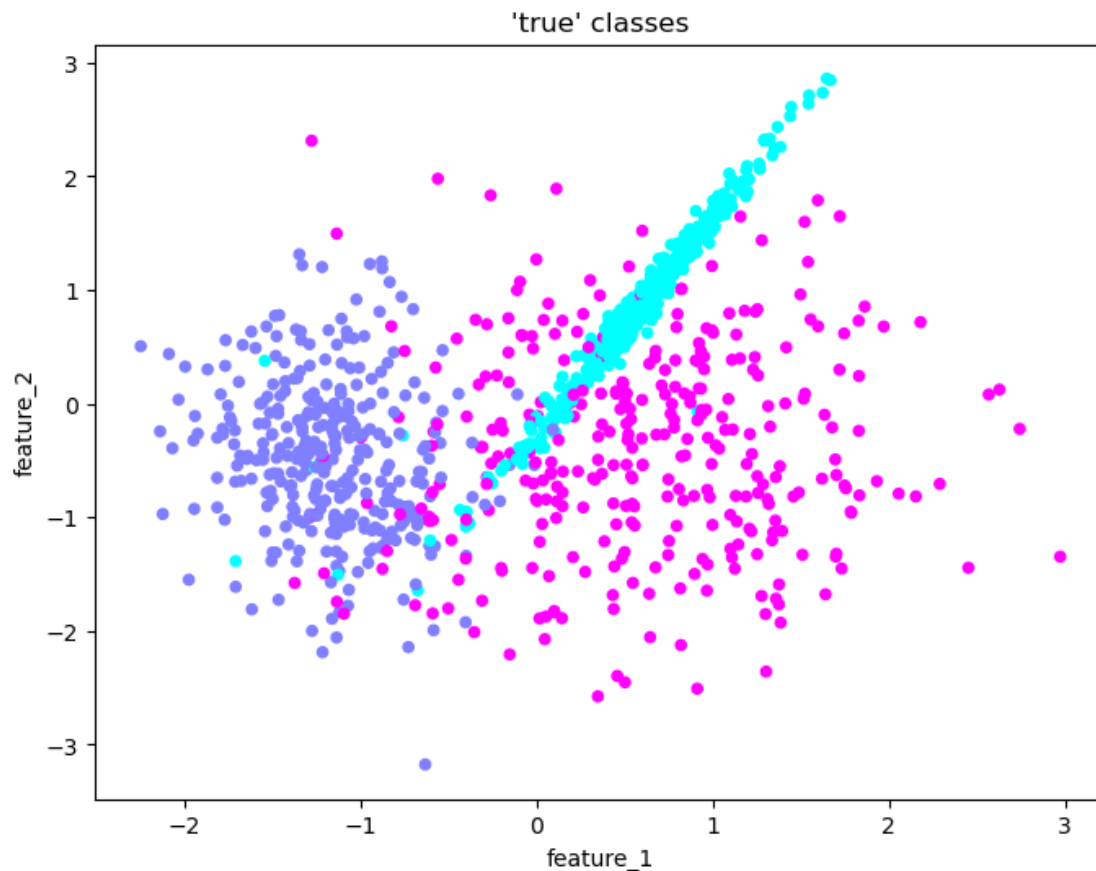
```
[167]: StandardScaler()
```

```
[168]: standardized = scaler.transform(df[["feature_1", "feature_2"]])  
standardized
```

```
[168]: array([[ 0.86225717,  1.45582779],  
          [ 0.14626453, -0.90269403],  
          [ 0.53065139,  0.74829594],  
          ...,  
          [-1.11216242, -1.02386509],  
          [ 1.82951322,  0.24260611],  
          [ 0.12922246,  0.15564846]])
```

```
[169]: df_scaled = df.copy()  
df_scaled.loc[:, ["feature_1", "feature_2"]] = standardized
```

```
[170]: df_scaled.plot.scatter("feature_1", "feature_2", c="category", cmap=pylab.cm.
      ↪ cool, title="'true' classes", figsize=(8,6), colorbar=False)
plt.show()
```



1 K-mean clustering

```
[171]: from sklearn.cluster import KMeans
```

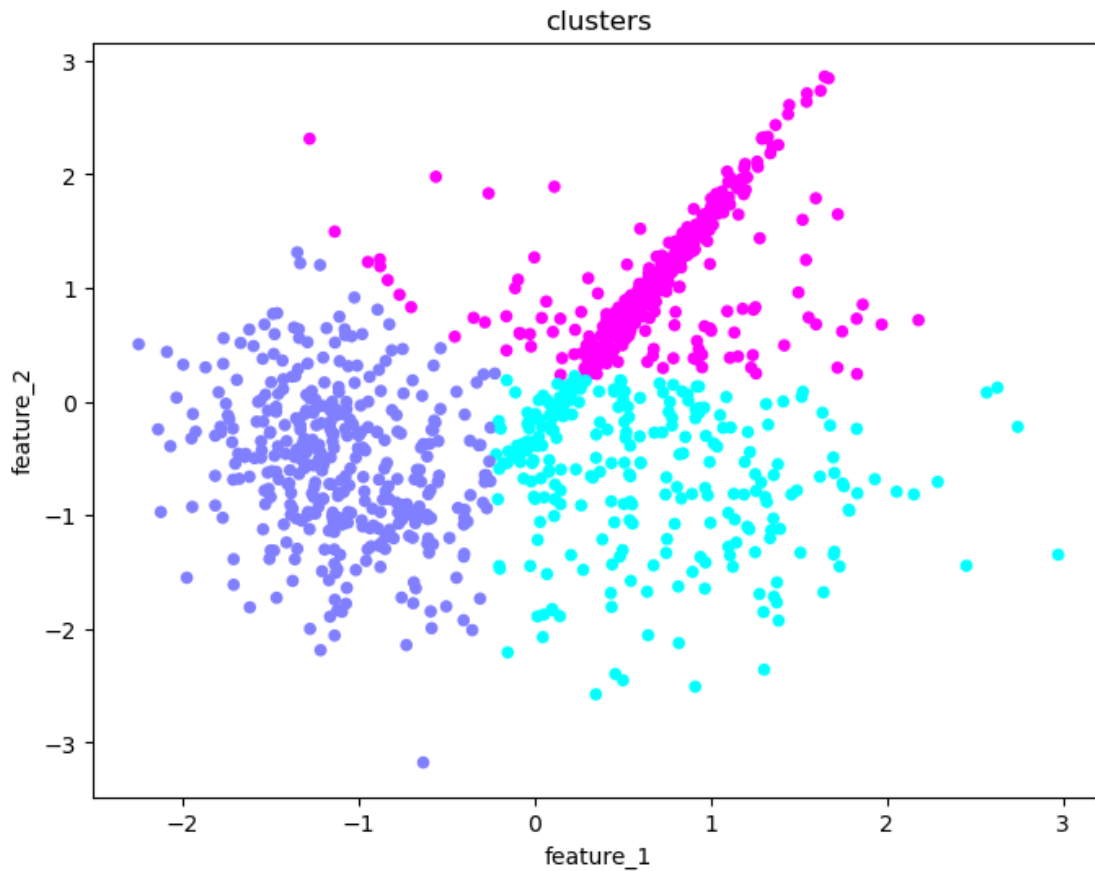
```
[172]: kmeans = KMeans(n_clusters=3, random_state=42)
```

```
[173]: kmeans.fit(df_scaled[["feature_1", "feature_2"]])
```

```
[173]: KMeans(n_clusters=3, random_state=42)
```

```
[174]: df_scaled["kmeans_cluster"] = kmeans.predict(df_scaled[["feature_1",
      ↪ "feature_2"]])
```

```
[175]: df_scaled.plot.scatter("feature_1", "feature_2", c="kmeans_cluster", cmap=pylab.
      ↪ cm.cool, figsize=(8,6), title="clusters", colorbar=False)
plt.show();
```



2 GMM Clustering

```
[176]: from sklearn.mixture import GaussianMixture
      from sklearn.decomposition import PCA
```

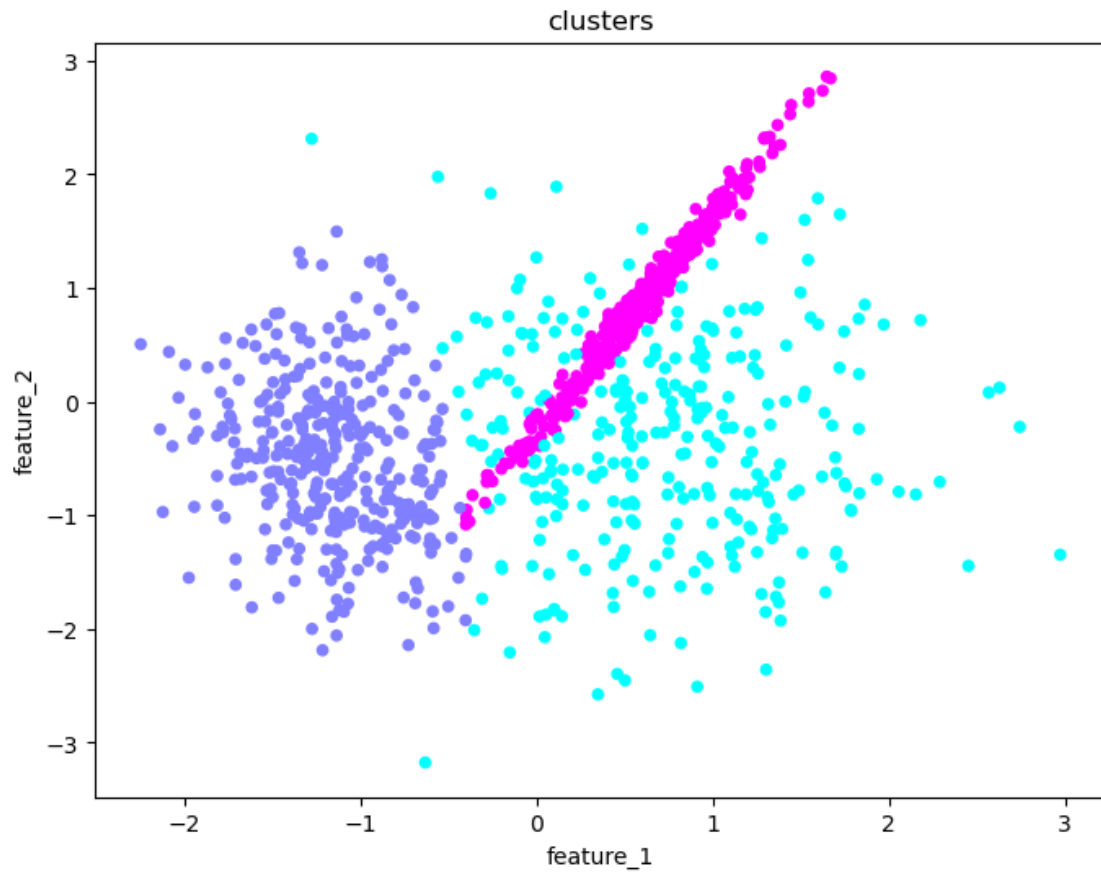
```
[177]: gmm = GaussianMixture(n_components=3, random_state=42)
      gmm.fit(df_scaled[["feature_1", "feature_2"]])
```

```
[177]: GaussianMixture(n_components=3, random_state=42)
```

```
[178]: pca = PCA(n_components=2)
      X_pca = pca.fit_transform(df_scaled[["feature_1", "feature_2"]])
```

```
[179]: df['cluster'] = gmm.predict(df_scaled[["feature_1", "feature_2"]])
```

```
[180]: df_scaled.plot.scatter("feature_1", "feature_2", c=df["cluster"], cmap=pylab.cm.  
      ↪ cool, figsize=(8,6), title="clusters", colorbar=False)  
plt.show();
```



```
[ ]:
```