

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

```
1 data = np.random.randn(50000) * 20 + 20
2
3 def find_anomalies(data):
4     anomalies=[]
5
6     data_std = np.std(data)
7     data_mean = np.mean(data)
8     anomaly_cut = data_std * 3
9
10    low_limit = data_mean - anomaly_cut
11    upp_limit = data_mean + anomaly_cut
12    print(low_limit, ' - ',upp_limit)
13    for datum in data:
14        if datum > upp_limit or datum < low_limit:
15            anomalies.append(datum)
16
17    return anomalies
```

```
1 anomalies = find_anomalies(data)
2 len(anomalies)

-39.69992174203004 - 79.82009753093006
147
```

```
1 import pandas as pd
2 stocks = pd.read_csv('http://dicyg.fi-c.unam.mx:8080/lalo/pypcd/presentaciones/st
3                      header='infer')
4 stocks.index = stocks.Date
5 stocks = stocks.drop(['Date'],axis=1)
6 stocks.head(3)
```

```
1 from matplotlib import projections
2 from mpl_toolkits.mplot3d import Axes3D
3 import matplotlib.pyplot as plt
4
```

```
5 fig = plt.figure(figsize=(8,5)).gca(projection='3d')
6 fig.scatter(stocks.MSFT,stocks.F,stocks.BAC)
7 fig.set_xlabel('Microsoft')
8 fig.set_ylabel('Ford')
9 fig.set_zlabel('Bank of America')
10 plt.show()
```

```
1 # Boxplot
2 data = np.random.randn(50000)*20+20
3 data = pd.DataFrame(data,columns=['random'])
4 #data
```

```
1 import seaborn as sns
2 sns.boxplot(data=data)
```

```
1 sns.boxplot(data=stocks)
```

```
1  from pandas.core.dtypes.missing import na_value_for_dtype
2  fig,ax = plt.subplots(nrows=1,ncols=3,figsize=(11,3))
3  sns.distplot(stocks.MSFT, ax=ax[0],color='darkblue')
4  sns.distplot(stocks.F, ax=ax[1], color='red')
5  sns.distplot(stocks.BAC, ax=ax[2], color='green')
```

```
1 # DBSCAN
2 import pandas as pd
3 import matplotlib.pyplot as plt
4 import numpy as np
5 import seaborn as sns
6 from pylab import rcParams
7 rcParams['figure.figsize'] = 5,4
8
9 from sklearn.cluster import DBSCAN
10 from collections import Counter
```

```
1 df = pd.read_csv("https://bit.ly/3arouNg")
2 df.shape

(6463, 13)
```

```
1 df.head(1)
```

```
1 data = df.iloc[:,1:3]
2 data.head(1)
```

```
1 model = DBSCAN(eps=0.2, min_samples=20).fit(data)
2 print(model)
3 model.get_params()
```

```
DBSCAN(eps=0.2, min_samples=20)
{'algorithm': 'auto',
 'eps': 0.2,
 'leaf_size': 30,
 'metric': 'euclidean',
 'metric_params': None,
 'min_samples': 20,
 'n_jobs': None,
 'p': None}
```

```
1 print(Counter(model.labels_))
2 outliers_df = pd.DataFrame(data)
3 #print(outliers_df[model.labels_==-1])

Counter({0: 6281, -1: 117, 1: 40, 2: 25})
```

```
1 fig = plt.figure()
2 ax = fig.add_axes([0.1,0.1,2,2])
3 colors = model.labels_
4
5 ax.scatter(data.iloc[:,0].values, data.iloc[:,1].values, c=colors, s=100)
```

```
1 # KrenelPCA "desde cero"
```

```
1 import scipy
2 from scipy.spatial.distance import pdist, squareform
3 from scipy.linalg import eigh
4 import numpy as np
5 import matplotlib.pyplot as plt
```

```
1 def rbf_kpca(X, gamma, n_components):
2     # distancias cuadradas de cada pareja
3     # del conjunto de datos, es MxN-dimensional
4     sq_dists = pdist(X, 'sqeuclidean')
5     #print(sq_dists.shape)
6     # convertirla en matriz
7     mat_sq_dists = squareform(sq_dists)
```

```

8  #print(mat_sq_dists.shape)
9  # Matriz de núcleos
10 K = np.exp(-gamma*mat_sq_dists)
11 # Centrar la matriz de núcleos
12 N = K.shape[0]
13 one_n = np.ones((N,N)) / N
14 K = K - one_n.dot(K) - K.dot(one_n) + one_n.dot(K).dot(one_n)
15 # Eigenpares; scipy.linalg.eigh devuelve los eigenpares ordenados ascendentemen
16 eigenvals,eigenvecs = eigh(K)
17 eigenvals,eigenvecs = eigenvals[::-1], eigenvecs[:,::-1] #invertir el orden
18 # Elegir los k primeros eigenvecs
19 alphas = np.column_stack([eigenvecs[:,i] for i in range(n_components)])
20 # Eigenvals
21 lambdas = np.column_stack([eigenvals[i] for i in range(n_components)])
22 return alphas,lambdas

```

```

1 # Medias lunas
2 from sklearn.datasets import make_moons
3 X,y = make_moons(n_samples=100, random_state=123)
4 plt.scatter(X[y==0,0],X[y==0,1],color='red',marker='^',alpha=0.5)
5 plt.scatter(X[y==1,0],X[y==1,1],color='blue',marker='o',alpha=0.5)

```

```

1 from sklearn.decomposition import PCA
2 pca = PCA(n_components=2)
3 X_pca = pca.fit_transform(X)

```

```

1 from IPython.core.pylabtools import figsize
2 fig,ax = plt.subplots(nrows=1,ncols=2,figsize=(10,4))
3 ax[0].scatter(X_pca[y==0,0],X_pca[y==0,1],color='red',marker='^',alpha=0.5)
4 ax[0].scatter(X_pca[y==1,0],X_pca[y==1,1],color='blue',marker='o',alpha=0.5)
5 ax[1].scatter(X_pca[y==0,0],np.zeros((50,1))+0.02,color='red',marker='^',alpha=0.
6 ax[1].scatter(X_pca[y==1,0],np.zeros((50,1))-0.02,color='blue',marker='o',alpha=0
7 ax[1].set_ylim([-1,1])
8 plt.show()

```

```

1 X_kpca,l = rbf_kpca(X, gamma=15, n_components=2)
2 fig,ax = plt.subplots(nrows=1,ncols=2,figsize=(10,4))
3 ax[0].scatter(X_kpca[y==0,0],X_kpca[y==0,1],color='red',marker='^',alpha=0.5)
4 ax[0].scatter(X_kpca[y==1,0],X_kpca[y==1,1],color='blue',marker='o',alpha=0.5)
5 ax[1].scatter(X_kpca[y==0,0],np.zeros((50,1))+0.02,color='red',marker='^',alpha=0)
6 ax[1].scatter(X_kpca[y==1,0],np.zeros((50,1))-0.02,color='blue',marker='o',alpha=0)
7 ax[1].set_ylim([-1,1])
8 plt.show()

```

```

1 # KPCA de sklearn
2 from sklearn.decomposition import KernelPCA
3 kpca = KernelPCA(n_components=2, kernel='rbf', gamma=15)
4 X_kpca = kpca.fit_transform(X)
5 fig,ax = plt.subplots(nrows=1,ncols=2,figsize=(10,4))
6 ax[0].scatter(X_kpca[y==0,0],X_kpca[y==0,1],color='red',marker='^',alpha=0.5)
7 ax[0].scatter(X_kpca[y==1,0],X_kpca[y==1,1],color='blue',marker='o',alpha=0.5)
8 ax[1].scatter(X_kpca[y==0,0],np.zeros((50,1))+0.02,color='red',marker='^',alpha=0)
9 ax[1].scatter(X_kpca[y==1,0],np.zeros((50,1))-0.02,color='blue',marker='o',alpha=0)
10 ax[1].set_ylim([-1,1])
11 plt.show()

```

```

1 from sklearn.decomposition import KernelPCA
2 kpca = KernelPCA(n_components=2, kernel='rbf', gamma=5)
3 X_kpca = kpca.fit_transform(X)
4 fig, ax = plt.subplots(nrows=1, ncols=2, figsize=(10, 4))
5 ax[0].scatter(X_kpca[y==0, 0], X_kpca[y==0, 1], color='red', marker='^', alpha=0.5)
6 ax[0].scatter(X_kpca[y==1, 0], X_kpca[y==1, 1], color='blue', marker='o', alpha=0.5)
7 ax[1].scatter(X_kpca[y==0, 0], np.zeros((50, 1)) + 0.02, color='red', marker='^', alpha=0.5)
8 ax[1].scatter(X_kpca[y==1, 0], np.zeros((50, 1)) - 0.02, color='blue', marker='o', alpha=0.5)
9 ax[1].set_ylim([-1, 1])
10 plt.show()

```

```

1 from sklearn.decomposition import KernelPCA
2 kpca = KernelPCA(n_components=2, kernel='rbf', gamma=10)
3 X_kpca = kpca.fit_transform(X)
4 fig, ax = plt.subplots(nrows=1, ncols=2, figsize=(10, 4))
5 ax[0].scatter(X_kpca[y==0, 0], X_kpca[y==0, 1], color='red', marker='^', alpha=0.5)
6 ax[0].scatter(X_kpca[y==1, 0], X_kpca[y==1, 1], color='blue', marker='o', alpha=0.5)
7 ax[1].scatter(X_kpca[y==0, 0], np.zeros((50, 1)) + 0.02, color='red', marker='^', alpha=0.5)
8 ax[1].scatter(X_kpca[y==1, 0], np.zeros((50, 1)) - 0.02, color='blue', marker='o', alpha=0.5)
9 ax[1].set_ylim([-1, 1])
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```



```

1 from sklearn.decomposition import KernelPCA
2 kpca = KernelPCA(n_components=2, kernel='rbf', gamma=20)
3 X_kpca = kpca.fit_transform(X)
4 fig, ax = plt.subplots(nrows=1, ncols=2, figsize=(10, 4))
5 ax[0].scatter(X_kpca[y==0, 0], X_kpca[y==0, 1], color='red', marker='^', alpha=0.5)
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8 ax[1].scatter(X_kpca[y==1, 0], np.zeros((50, 1)) - 0.02, color='blue', marker='o', alpha=0.5)
9 ax[1].set_ylim([-1, 1])
10 plt.show()

```

```

1 from sklearn.datasets import make_circles
2 X, y = make_circles(n_samples=1000, random_state=123, noise=0.1, factor=0.2)
3 plt.scatter(X[y==0, 0], X[y==0, 1], color='red', marker='^', alpha=0.5)
4 plt.scatter(X[y==1, 0], X[y==1, 1], color='blue', marker='o', alpha=0.5)
5 plt.show()

```

```

1 # Con PCA
2 pca = PCA(n_components=2)
3 X_pca = pca.fit_transform(X)
4 fig, ax = plt.subplots(nrows=1, ncols=2, figsize=(10,4))
5 ax[0].scatter(X_pca[y==0,0],X_pca[y==0,1],color='red',marker='^',alpha=0.5)
6 ax[0].scatter(X_pca[y==1,0],X_pca[y==1,1],color='blue',marker='o',alpha=0.5)
7 ax[1].scatter(X_pca[y==0,0],np.zeros((500,1))+0.02,color='red',marker='^',alpha=0.5)
8 ax[1].scatter(X_pca[y==1,0],np.zeros((500,1))-0.02,color='blue',marker='o',alpha=0.5)
9 ax[1].set_ylim([-1,1])
10 plt.show()

```

```

1 X_kpca,l = rbf_kpca(X, gamma=15, n_components=2)
2 fig, ax = plt.subplots(nrows=1, ncols=2, figsize=(10,4))
3 ax[0].scatter(X_kpca[y==0,0],X_kpca[y==0,1],color='red',marker='^',alpha=0.5)
4 ax[0].scatter(X_kpca[y==1,0],X_kpca[y==1,1],color='blue',marker='o',alpha=0.5)
5 ax[1].scatter(X_kpca[y==0,0],np.zeros((500,1))+0.02,color='red',marker='^',alpha=0.5)
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7 ax[1].set_ylim([-1,1])
8 plt.show()

```

```

1  from sklearn.decomposition import KernelPCA
2  kpca = KernelPCA(n_components=2, kernel='rbf', gamma=15)
3  X_kpca = kpca.fit_transform(X)
4  fig,ax = plt.subplots(nrows=1,ncols=2,figsize=(10,4))
5  ax[0].scatter(X_kpca[y==0,0],X_kpca[y==0,1],color='red',marker='^',alpha=0.5)
6  ax[0].scatter(X_kpca[y==1,0],X_kpca[y==1,1],color='blue',marker='o',alpha=0.5)
7  ax[1].scatter(X_kpca[y==0,0],np.zeros((500,1))+0.02,color='red',marker='^',alpha=
8  ax[1].scatter(X_kpca[y==1,0],np.zeros((500,1))-0.02,color='blue',marker='o',alpha=
9  ax[1].set_ylim([-1,1])
10 plt.show()

```

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