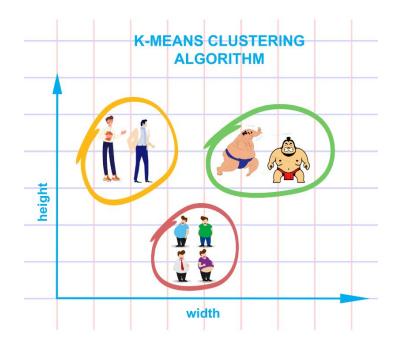
UNSUPERVISED LEARNING

Clustering Algorithms

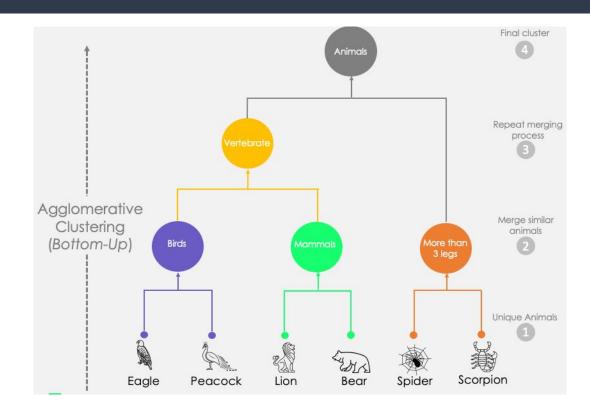
K-Means Clustering

```
from sklearn.cluster import
KMeans
data = list(zip(x, y))
inertias = []
for i in range (1,11):
    kmeans =
KMeans(n clusters=i)
    kmeans.fit(data)
inertias.append(kmeans.inertia )
plt.plot(range(1,11), inertias,
marker='o')
plt.title('Elbow method')
plt.xlabel('Number of clusters')
plt.ylabel('Inertia')
plt.show()
```



Hierarchical Clustering

```
import numpy as np
import matplotlib.pyplot as plt
from scipy.cluster.hierarchy
import dendrogram, linkage
x = [4, 5, 10, 4, 3, 11, 14, 6,
10, 12]
y = [21, 19, 24, 17, 16, 25, 24,
22, 21, 21]
data = list(zip(x, y))
linkage data = linkage(data,
method='ward',
metric='euclidean')
dendrogram(linkage data)
plt.show()
```



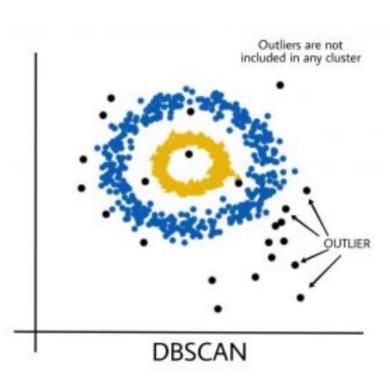
DBSCAN

```
import numpy as np
from sklearn import metrics
from sklearn.cluster import DBSCAN

db = DBSCAN(eps=0.3, min_samples=10).fit(X)
labels = db.labels_

# Number of clusters in labels, ignoring noise if present.
n_clusters_ = len(set(labels)) - (1 if -1 in labels else 0)
n_noise_ = list(labels).count(-1)

print("Estimated number of clusters: %d" % n_clusters_)
print("Estimated number of noise points: %d" % n_noise_)
```



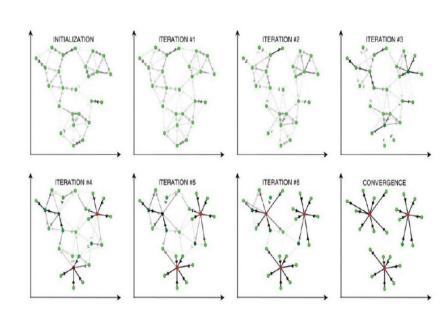
Affinity Propegation

from sklearn.cluster import AffinityPropagation from sklearn import metrics from sklearn.datasets.samples_generator import make_blobs

```
centers = [[1, 1], [-1, -1], [1, -1], [-1, -1]]
X, labels_true = make_blobs(n_samples = 400, centers
= centers, cluster_std = 0.5, random_state = 0)
```

af = AffinityPropagation(preference =-50).fit(X)
cluster_centers_indices = af.cluster_centers_indices_
labels = af.labels_

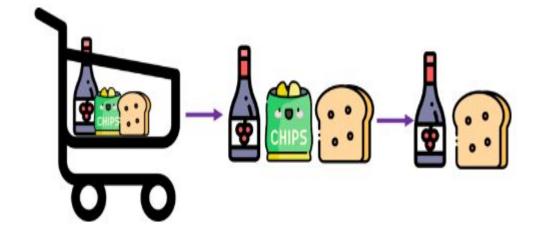
n_clusters_ = len(cluster_centers_indices)



Associative Algorithms

Apriori Algorithm

- 1. # Build the model
- frq_items1 = AP(basket1_France, min_support = 0.05, use_colnames = True)
- 3.
- 4. # Collect the inferred rules in a dataframe
- 5. rules1 = AR(frq_items1, metric = "lift", min_threshold = 1)
- f. rules1 = rules1.sort_values(['confidence', 'lift'], ascending = [False, False])
- 7. **print**(rules1.head())



Eclat Algorithm

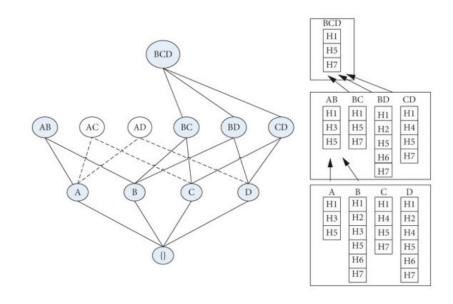
```
from pyECLAT import ECLAT
```

```
# create an instance of eclat
my_eclat = ECLAT(data=data, verbose=True)
```

fit the algorithm
rule_indices, rule_supports =
my_eclat.fit(min_support=min_support,

min_combination=min_n_products,

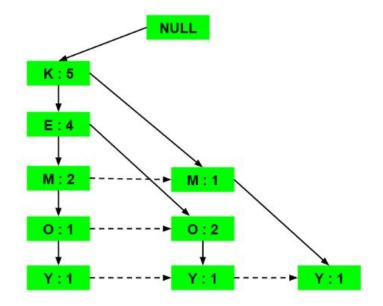
max_combination=max_length)



FP Growth

print(frequent itemsets)

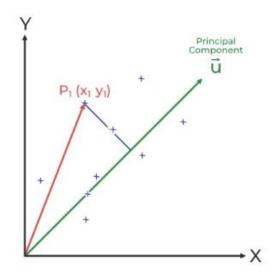
```
from mlxtend.preprocessing import TransactionEncoder
from mlxtend.frequent patterns import fpgrowth
# Sample transaction dataset
dataset = [['Apple', 'Banana', 'Egg'],
          ['Banana', 'Eqq', 'Milk'],
          ['Apple', 'Banana'],
          ['Banana', 'Milk']]
# Convert dataset to one-hot encoded format
te = TransactionEncoder()
te ary = te.fit(dataset).transform(dataset)
df = pd.DataFrame(te ary, columns=te.columns)
# Mining frequent itemsets using FP-Growth
frequent itemsets = fpgrowth(df, min support=0.3,
use colnames=True)
# Display the frequent itemsets
```



Dimensionality Reduction

Principal Component Analysis

```
# Matrix multiplication or
dot Product
Z_pca = Z @ pca_component
# Rename the columns name
Z_pca.rename({ 'PC1':
   'PCA1', 'PC2': 'PCA2'},
   axis=1, inplace=True)
# Print the Principal
Component values
print(Z pca)
```



$$Proj_{P_1}\vec{u} = \frac{P_1 \cdot \vec{u}}{|u|}$$
$$= P_1 \cdot \vec{u} \quad\vec{u} - Unit Vector$$

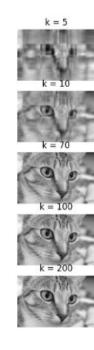
Singular Value Decomposition

```
# Imports
```

curr fig = 0

```
from skimage.color import rgb2gray
from skimage import data
import matplotlib.pyplot as plt
import numpy as np
from scipy.linalg import svd
SVD on image compression
cat = data.chelsea()
plt.imshow(cat)
# convert to grayscale
gray cat = rgb2gray(cat)
# calculate the SVD and plot the image
U, S, V T = svd(gray cat, full matrices=False)
S = np.diag(S)
```

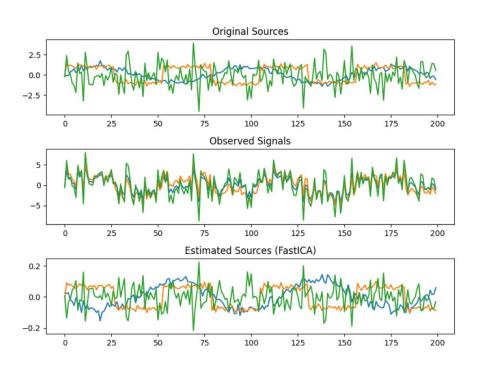
fig, ax = plt.subplots(5, 2, figsize=(8, 20))





Independent Component Analysis

```
import numpy as np
from sklearn.decomposition import FastICA
import matplotlib.pyplot as plt
ica = FastICA(n components=3)
S = ica.fit transform(X) # Estimated sources
# Plot the results
plt.figure(figsize=(8, 6))
plt.subplot(3, 1, 1)
plt.title('Original Sources')
plt.plot(S)
plt.subplot(3, 1, 2)
plt.title('Observed Signals')
plt.plot(X)
plt.subplot(3, 1, 3)
plt.title('Estimated Sources (FastICA)')
plt.plot(S)
plt.tight_layout()
plt.show()
```



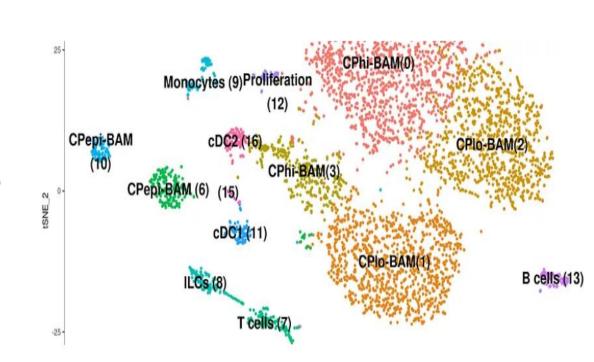
t-SNE

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.manifold import TSNE
from sklearn.preprocessing import StandardScaler
```

```
data_1000 = standardized_data[0:1000, :]
labels_1000 = labels[0:1000]
```

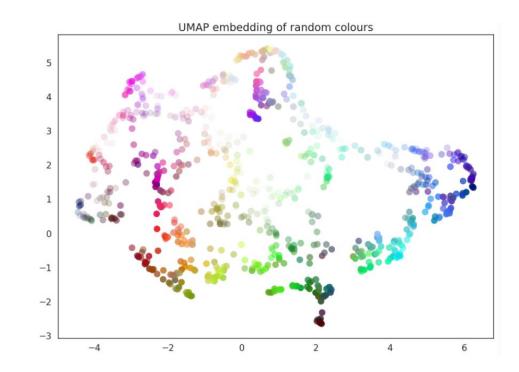
```
model = TSNE(n_components = 2, random_state = 0)
tsne_data = model.fit_transform(data_1000)
tsne_data = np.vstack((tsne_data.T, labels_1000)).T
tsne_df = pd.DataFrame(data = tsne_data,columns
=("Dim_1", "Dim_2", "label"))
```

```
sn.scatterplot(data=tsne_df, x='Dim_1', y='Dim_2', hue='label', palette="bright") plt.show()
```



UMAP

```
import numpy as np
import matplotlib.pyplot as plt
from mpl_toolkits.mplot3d import Axes3D
import seaborn as sns
import umap
%matplotlib inline
fit = umap.UMAP()
%time u = fit.fit_transform(data)
CPU times: user 7.73 s, sys: 211 ms, total: 7.94 s
Wall time: 6.8 s
plt
plt.sc after(u[:,0],c=data)
plt.title('UMAP embedding of random colours');
```

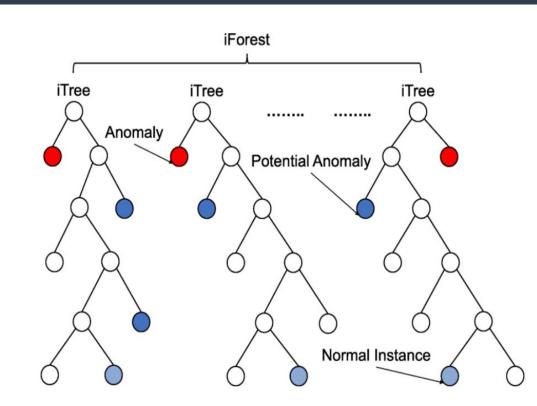


Anomaly Detection

Isolation Forest

import numpy as np import pandas as pd import seaborn as sns import matplotlib.pyplot as plt from sklearn.ensemble import IsolationForest

df = pd.read_csv('salary.csv')
df.head(10)
model=IsolationForest(n_estimators=50,
max_samples='auto',
contamination=float(0.1),max_features=1.0)
model.fit(df[['salary']])



One - Class SVM

import numpy as np
import matplotlib.pyplot as plt
import matplotlib.font_manager
from sklearn import svm
%matplotlib inline

clf = svm.OneClassSVM()
clf.fit(X_train)