

## Imports and Data Loading

```
In [1]: import pandas as pd
import numpy as np
from sklearn.preprocessing import StandardScaler

# Load datasets
cancer_df = pd.read_csv('cancer.csv')
bankrupt_df = pd.read_csv('bankrupt.csv')

# Show column names and info
print("Cancer dataset columns:\n", cancer_df.columns)
print("Bankruptcy dataset columns:\n", bankrupt_df.columns)
```

Cancer dataset columns:

```
Index(['Patient_ID', 'Age', 'Gender', 'Country_Region', 'Year', 'Genetic_Risk',  
      'Air_Pollution', 'Alcohol_Use', 'Smoking', 'Obesity_Level',  
      'Cancer_Type', 'Cancer_Stage', 'Treatment_Cost_USD', 'Survival_Years',  
      'Target_Severity_Score'],  
      dtype='object')
```

Bankruptcy dataset columns:

```
Index(['Bankrupt?', 'ROA(C) before interest and depreciation before interest',  
      'ROA(A) before interest and % after tax',  
      'ROA(B) before interest and depreciation after tax',  
      'Operating Gross Margin', 'Realized Sales Gross Margin',  
      'Operating Profit Rate', 'Pre-tax net Interest Rate',  
      'After-tax net Interest Rate',  
      'Non-industry income and expenditure/revenue',  
      'Continuous interest rate (after tax)', 'Operating Expense Rate',  
      'Research and development expense rate', 'Cash flow rate',  
      'Interest-bearing debt interest rate', 'Tax rate (A)',  
      'Net Value Per Share (B)', 'Net Value Per Share (A)',  
      'Net Value Per Share (C)', 'Persistent EPS in the Last Four Seasons',  
      'Cash Flow Per Share', 'Revenue Per Share (Yuan ¥)',  
      'Operating Profit Per Share (Yuan ¥)',  
      'Per Share Net profit before tax (Yuan ¥)',  
      'Realized Sales Gross Profit Growth Rate',  
      'Operating Profit Growth Rate', 'After-tax Net Profit Growth Rate',  
      'Regular Net Profit Growth Rate', 'Continuous Net Profit Growth Rate',  
      'Total Asset Growth Rate', 'Net Value Growth Rate',  
      'Total Asset Return Growth Rate Ratio', 'Cash Reinvestment %',  
      'Current Ratio', 'Quick Ratio', 'Interest Expense Ratio',  
      'Total debt/Total net worth', 'Debt ratio %', 'Net worth/Assets',  
      'Long-term fund suitability ratio (A)', 'Borrowing dependency',  
      'Contingent liabilities/Net worth',  
      'Operating profit/Paid-in capital',  
      'Net profit before tax/Paid-in capital',  
      'Inventory and accounts receivable/Net value', 'Total Asset Turnover',  
      'Accounts Receivable Turnover', 'Average Collection Days',  
      'Inventory Turnover Rate (times)', 'Fixed Assets Turnover Frequency',  
      'Net Worth Turnover Rate (times)', 'Revenue per person',  
      'Operating profit per person', 'Allocation rate per person',  
      'Working Capital to Total Assets', 'Quick Assets/Total Assets',  
      'Current Assets/Total Assets', 'Cash/Total Assets',  
      'Quick Assets/Current Liability', 'Cash/Current Liability',  
      'Current Liability to Assets', 'Operating Funds to Liability',  
      'Inventory/Working Capital', 'Inventory/Current Liability',  
      'Current Liabilities/Liability', 'Working Capital/Equity',  
      'Current Liabilities/Equity', 'Long-term Liability to Current Assets',  
      'Retained Earnings to Total Assets', 'Total income/Total expense',  
      'Total expense/Assets', 'Current Asset Turnover Rate',  
      'Quick Asset Turnover Rate', 'Working capital Turnover Rate',
```

```

        ' Cash Turnover Rate', ' Cash Flow to Sales', ' Fixed Assets to Asset
s',
        ' Current Liability to Liability', ' Current Liability to Equity',
        ' Equity to Long-term Liability', ' Cash Flow to Total Assets',
        ' Cash Flow to Liability', ' CFO to Assets', ' Cash Flow to Equity',
        ' Current Liability to Current Assets', ' Liability-Assets Flag',
        ' Net Income to Total Assets', ' Total assets to GNP price',
        ' No-credit Interval', ' Gross Profit to Sales',
        ' Net Income to Stockholder's Equity', ' Liability to Equity',
        ' Degree of Financial Leverage (DFL)',
        ' Interest Coverage Ratio (Interest expense to EBIT)',
        ' Net Income Flag', ' Equity to Liability'],
dtype='object')

```

## Preprocessing (Drop NAs, Keep Numeric, Scale)

```

In [2]: # Drop rows with missing values
cancer_df_clean = cancer_df.dropna()
bankrupt_df_clean = bankrupt_df.dropna()

# Select only numerical columns
cancer_X = cancer_df_clean.select_dtypes(include=[np.number])
bankrupt_X = bankrupt_df_clean.select_dtypes(include=[np.number])

# Standardize features
scaler_cancer = StandardScaler()
cancer_X_scaled = scaler_cancer.fit_transform(cancer_X)

scaler_bankrupt = StandardScaler()
bankrupt_X_scaled = scaler_bankrupt.fit_transform(bankrupt_X)

```

```

In [3]: from sklearn.cluster import KMeans
from sklearn.mixture import GaussianMixture
from sklearn.metrics import silhouette_score

# You can change n_clusters if you wish (try 2-5)
n_clusters = 3

# K-Means
kmeans_cancer = KMeans(n_clusters=n_clusters, random_state=42)
kmeans_cancer_labels = kmeans_cancer.fit_predict(cancer_X_scaled)

kmeans_bankrupt = KMeans(n_clusters=n_clusters, random_state=42)
kmeans_bankrupt_labels = kmeans_bankrupt.fit_predict(bankrupt_X_scaled)

# Gaussian Mixture Model (EM)
gmm_cancer = GaussianMixture(n_components=n_clusters, random_state=42)
gmm_cancer_labels = gmm_cancer.fit_predict(cancer_X_scaled)

gmm_bankrupt = GaussianMixture(n_components=n_clusters, random_state=42)
gmm_bankrupt_labels = gmm_bankrupt.fit_predict(bankrupt_X_scaled)

# Silhouette Scores
print("Silhouette (KMeans, Cancer):", silhouette_score(cancer_X_scaled, kmeans_cancer_labels))
print("Silhouette (GMM, Cancer):", silhouette_score(cancer_X_scaled, gmm_cancer_labels))

```

```
print("Silhouette (KMeans, Bankrupt):", silhouette_score(bankrupt_X_scaled,
print("Silhouette (GMM, Bankrupt):", silhouette_score(bankrupt_X_scaled, gmm
```

```
/Users/maxmatkovski/Library/Python/3.9/lib/python/site-packages/sklearn/cluster/_kmeans.py:1416: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning
```

```
super()._check_params_vs_input(X, default_n_init=10)
/Users/maxmatkovski/Library/Python/3.9/lib/python/site-packages/sklearn/cluster/_kmeans.py:1416: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning
```

```
super()._check_params_vs_input(X, default_n_init=10)
```

```
Silhouette (KMeans, Cancer): 0.08681559132410345
```

```
Silhouette (GMM, Cancer): 0.07222682244912479
```

```
Silhouette (KMeans, Bankrupt): 0.10741175195746241
```

```
Silhouette (GMM, Bankrupt): 0.03379523488318877
```

```
In [4]: from sklearn.decomposition import PCA, FastICA
from sklearn.random_projection import GaussianRandomProjection

n_components = 2

# PCA
pca_cancer = PCA(n_components=n_components)
cancer_pca = pca_cancer.fit_transform(cancer_X_scaled)

pca_bankrupt = PCA(n_components=n_components)
bankrupt_pca = pca_bankrupt.fit_transform(bankrupt_X_scaled)

# ICA
ica_cancer = FastICA(n_components=n_components, random_state=42)
cancer_ica = ica_cancer.fit_transform(cancer_X_scaled)

ica_bankrupt = FastICA(n_components=n_components, random_state=42)
bankrupt_ica = ica_bankrupt.fit_transform(bankrupt_X_scaled)

# Randomized Projections
rp_cancer = GaussianRandomProjection(n_components=n_components, random_state=42)
cancer_rp = rp_cancer.fit_transform(cancer_X_scaled)

rp_bankrupt = GaussianRandomProjection(n_components=n_components, random_state=42)
bankrupt_rp = rp_bankrupt.fit_transform(bankrupt_X_scaled)
```

```
In [5]: # Defining function to run on
def run_clustering(X, n_clusters=3):
    kmeans = KMeans(n_clusters=n_clusters, random_state=42).fit_predict(X)
    gmm = GaussianMixture(n_components=n_clusters, random_state=42).fit_predict(X)
    return kmeans, gmm

# Cancer dataset
km_pca_cancer, gm_pca_cancer = run_clustering(cancer_pca)
km_ica_cancer, gm_ica_cancer = run_clustering(cancer_ica)
km_rp_cancer, gm_rp_cancer = run_clustering(cancer_rp)

# Bankruptcy dataset
```

```

km_pca_bankrupt, gm_pca_bankrupt = run_clustering(bankrupt_pca)
km_ica_bankrupt, gm_ica_bankrupt = run_clustering(bankrupt_ica)
km_rp_bankrupt, gm_rp_bankrupt = run_clustering(bankrupt_rp)

```

```

/Users/maxmatkovski/Library/Python/3.9/lib/python/site-packages/sklearn/cluster/_kmeans.py:1416: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning

```

```

    super()._check_params_vs_input(X, default_n_init=10)

```

```

/Users/maxmatkovski/Library/Python/3.9/lib/python/site-packages/sklearn/cluster/_kmeans.py:1416: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning

```

```

    super()._check_params_vs_input(X, default_n_init=10)

```

```

/Users/maxmatkovski/Library/Python/3.9/lib/python/site-packages/sklearn/cluster/_kmeans.py:1416: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning

```

```

    super()._check_params_vs_input(X, default_n_init=10)

```

```

/Users/maxmatkovski/Library/Python/3.9/lib/python/site-packages/sklearn/cluster/_kmeans.py:1416: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning

```

```

    super()._check_params_vs_input(X, default_n_init=10)

```

```

/Users/maxmatkovski/Library/Python/3.9/lib/python/site-packages/sklearn/cluster/_kmeans.py:1416: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning

```

```

    super()._check_params_vs_input(X, default_n_init=10)

```

```

/Users/maxmatkovski/Library/Python/3.9/lib/python/site-packages/sklearn/cluster/_kmeans.py:1416: FutureWarning: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of `n_init` explicitly to suppress the warning

```

```

    super()._check_params_vs_input(X, default_n_init=10)

```

In [6]: `import matplotlib.pyplot as plt`

```

# Visualize clustering for Cancer dataset (PCA)

```

```

plt.figure(figsize=(6,4))

```

```

plt.scatter(cancer_pca[:, 0], cancer_pca[:, 1], c=km_pca_cancer, cmap='rainbow')

```

```

plt.title("KMeans Clusters (Cancer PCA)")

```

```

plt.xlabel("PC1")

```

```

plt.ylabel("PC2")

```

```

plt.show()

```

```

# Visualize GMM for Cancer dataset (PCA)

```

```

plt.figure(figsize=(6,4))

```

```

plt.scatter(cancer_pca[:, 0], cancer_pca[:, 1], c=gm_pca_cancer, cmap='rainbow')

```

```

plt.title("GMM Clusters (Cancer PCA)")

```

```

plt.xlabel("PC1")

```

```

plt.ylabel("PC2")

```

```

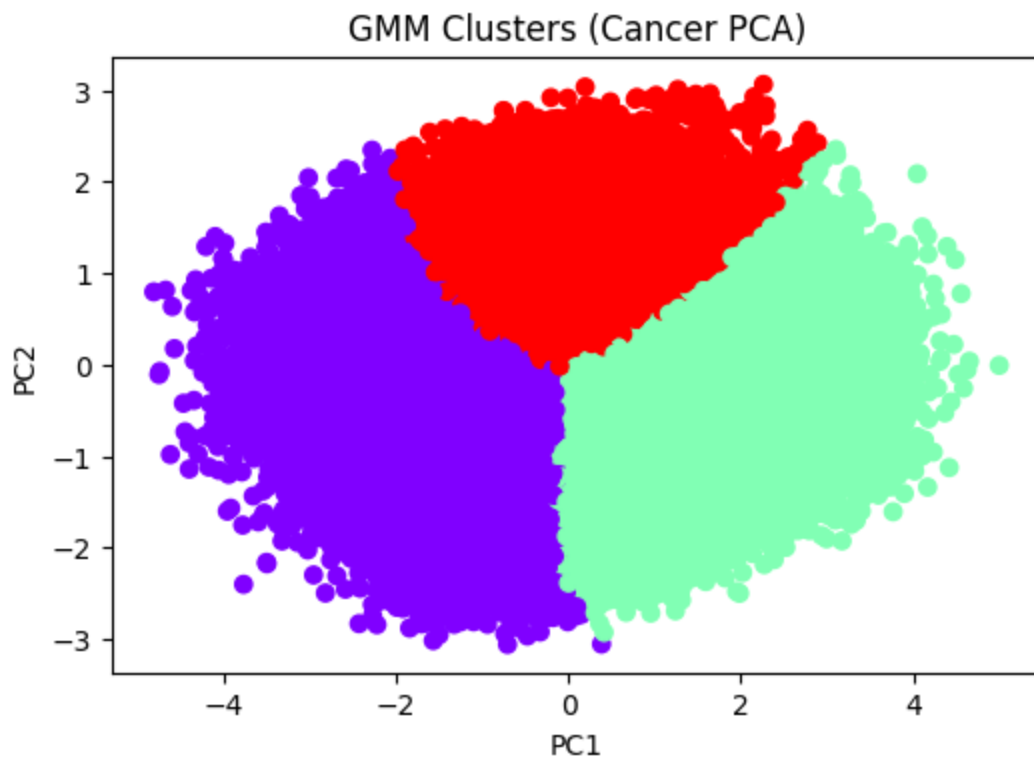
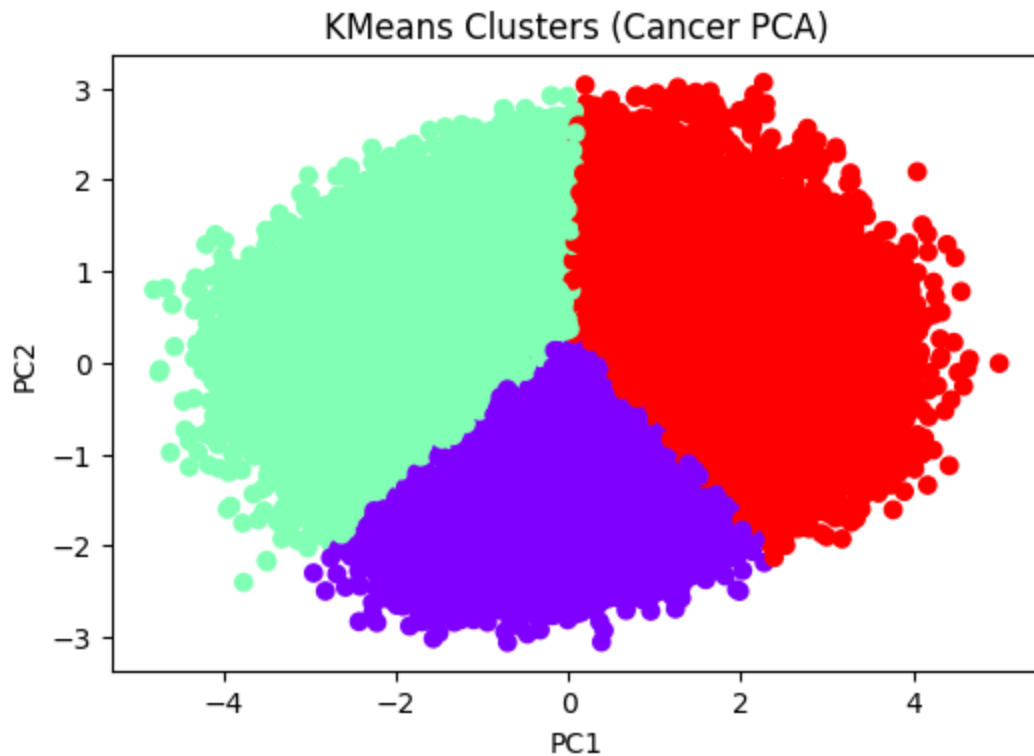
plt.show()

```

```

# Repeat for ICA and RP, and for bankruptcy dataset as desired

```



```
In [ ]: # Example: Print silhouette scores
methods = {
    'Original': (cancer_X_scaled, kmeans_cancer_labels, gmm_cancer_labels),
    'PCA':      (cancer_pca, km_pca_cancer, gm_pca_cancer),
    'ICA':      (cancer_ica, km_ica_cancer, gm_ica_cancer),
    'RP':       (cancer_rp, km_rp_cancer, gm_rp_cancer)
}
```

```

for name, (X, kmeans_labels, gmm_labels) in methods.items():
    print(f"Cancer {name} - KMeans Silhouette: {silhouette_score(X, kmeans_labels)}")
    print(f"Cancer {name} - GMM Silhouette: {silhouette_score(X, gmm_labels)}")

```

Cancer Original - KMeans Silhouette: 0.087  
 Cancer Original - GMM Silhouette: 0.072  
 Cancer PCA - KMeans Silhouette: 0.329  
 Cancer PCA - GMM Silhouette: 0.330  
 Cancer ICA - KMeans Silhouette: 0.338

```

In [ ]: import pandas as pd
import numpy as np

# Load and clean
cancer_df = pd.read_csv('cancer.csv').dropna()
# Keep only numeric columns
cancer_X = cancer_df.select_dtypes(include=[np.number])

```

```

In [ ]: n_cancer, d_cancer = cancer_X.shape
print(f"n_cancer: {n_cancer}, d_cancer: {d_cancer}")

```

```

In [ ]: cancer_X.shape

```

```

In [ ]: cancer_X

```

```

In [ ]: import numpy as np
import pandas as pd

bankrupt_df = pd.read_csv('bankrupt.csv').dropna()
bankrupt_X = bankrupt_df.select_dtypes(include=[np.number])

# sample and feature counts:
n_bankrupt, d_bankrupt = bankrupt_X.shape
print(f"n_bankrupt: {n_bankrupt}, d_bankrupt: {d_bankrupt}")

```

```

In [ ]: n_bankrupt, d_bankrupt = bankrupt_X.shape
print(f"n_bankrupt: {n_bankrupt}, d_bankrupt: {d_bankrupt}")

```

```

In [ ]: import pandas as pd
from sklearn.preprocessing import StandardScaler

# Load data
cancer_df = pd.read_csv('cancer.csv')

# Drop rows with missing values
cancer_df_clean = cancer_df.dropna()

# Select only numerical columns
cancer_X = cancer_df_clean.select_dtypes(include=['float64', 'int64'])

# Standardize features (important for PCA/KMeans)
scaler = StandardScaler()
cancer_X_scaled = scaler.fit_transform(cancer_X)

```

```
In [ ]: from sklearn.decomposition import PCA

# Fit PCA with 2 components
pca = PCA(n_components=2)
cancer_pca = pca.fit_transform(cancer_X_scaled)

# Explained variance
explained_var = pca.explained_variance_ratio_
total_explained = explained_var.sum() * 100
print(f"Total variance explained by first two PCs: {total_explained:.1f}%")
```

```
In [ ]:
```

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In [ ]:
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In [ ]:
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```
In [ ]: # Neural Network
```

```
In [ ]: import pandas as pd
import numpy as np
from sklearn.preprocessing import StandardScaler, LabelEncoder
from sklearn.model_selection import train_test_split
from sklearn.decomposition import PCA, FastICA
from sklearn.random_projection import GaussianRandomProjection
from sklearn.neural_network import MLPClassifier
from sklearn.metrics import accuracy_score, f1_score

# Load and clean data
cancer_df = pd.read_csv('cancer.csv')
cancer_df_clean = cancer_df.dropna()
cancer_X = cancer_df_clean.select_dtypes(include=[np.number])

# Use Cancer_Type as label
y = cancer_df_clean['Cancer_Type'].values
le = LabelEncoder()
y_encoded = le.fit_transform(y)

# Standardize features
scaler = StandardScaler()
cancer_X_scaled = scaler.fit_transform(cancer_X)

# Dimensionality reduction
pca = PCA(n_components=2)
cancer_pca = pca.fit_transform(cancer_X_scaled)

ica = FastICA(n_components=2, random_state=42)
cancer_ica = ica.fit_transform(cancer_X_scaled)

rp = GaussianRandomProjection(n_components=2, random_state=42)
cancer_rp = rp.fit_transform(cancer_X_scaled)

# Train-test split
X_train, X_test, y_train, y_test = train_test_split(cancer_X_scaled, y_encoded,
```



```

X_train_pca, X_test_pca, _, _ = train_test_split(cancer_pca, y_encoded, test_size=0.2)
X_train_ica, X_test_ica, _, _ = train_test_split(cancer_ica, y_encoded, test_size=0.2)
X_train_rp, X_test_rp, _, _ = train_test_split(cancer_rp, y_encoded, test_size=0.2)

# Neural network
def run_nn(X_train, X_test, y_train, y_test, name=""):
    clf = MLPClassifier(hidden_layer_sizes=(32,), max_iter=300, random_state=0)
    clf.fit(X_train, y_train)
    y_pred = clf.predict(X_test)
    acc = accuracy_score(y_test, y_pred)
    f1 = f1_score(y_test, y_pred, average='weighted')
    print(f"{name} - Accuracy: {acc:.3f}, F1: {f1:.3f}")

run_nn(X_train, X_test, y_train, y_test, "Original")
run_nn(X_train_pca, X_test_pca, y_train, y_test, "PCA")
run_nn(X_train_ica, X_test_ica, y_train, y_test, "ICA")
run_nn(X_train_rp, X_test_rp, y_train, y_test, "RP")

```

In [ ]:

In [ ]: