

ML - Experiment 7 - PCA

```
import pandas as pd
import numpy as np
from numpy.linalg import eig
```

DATASET 1 - CODE

```
data = np.array([[4, 6], [8, 2], [13, 3], [7, 15]])

def PCA(df):
    centered_data = df - df.mean()

    cov_matrix = np.cov(centered_data, rowvar=False)

    eigenvalues, eigenvectors = np.linalg.eig(cov_matrix)

    sorted_indices = np.argsort(eigenvalues)[::-1]
    eigenvalues = eigenvalues[sorted_indices]
    eigenvectors = eigenvectors[:, sorted_indices]

    new_values = np.dot(centered_data, eigenvectors)[:,:0]

    print("Centered Data:")
    print(centered_data)
    print("\nCovariance Matrix:")
    print(cov_matrix)
    print("\nEigenvalues:")
    print(eigenvalues)
    print("\nEigenvectors:")
    print(eigenvectors)
    print("\nNew Values:")
    print(new_values)

df2 = pd.DataFrame(data)
PCA(df2)
```

OUTPUT

```
Centered Data:
  0  1
0 -4.0 -0.5
1  0.0 -4.5
2  5.0 -3.5
3 -1.0  8.5

Covariance Matrix:
[[14. -8.]
 [-8. 35.]]

Eigenvalues:
[37.70037878 11.29962122]

Eigenvectors:
[[ 0.31981892 -0.94747869]
 [-0.94747869 -0.31981892]]

New Values:
[-0.80553633  4.26365409  4.91526999 -8.37338775]
```

DATASET 2 - CODE

```
df = pd.read_csv('/content/gdrive/MyDrive/ML/salary_data.csv')
```



	YearsExperience	Salary	
0	1.1	39343.0	
1	1.3	46205.0	
2	1.5	37731.0	
3	2.0	43525.0	
4	2.2	39891.0	

```
import pandas as pd
from sklearn.decomposition import PCA
```

```
def apply_pca(data, n_components):
    pca = PCA(n_components=n_components)
    principalComponents = pca.fit_transform(data)
    principalDf = pd.DataFrame(data=principalComponents, columns=[f'New Values'
    for i in range(n_components)])
```

```
# Calculate additional PCA components
centered_data = data - data.mean()
cov_matrix = pca.get_covariance()
```

```
eigenvalues = pca.explained_variance_  
eigenvectors = pca.components_
```

```
print("Centered Data:")  
print(centered_data)  
print("\nCovariance Matrix:")  
print(cov_matrix)  
print("\nEigenvalues:")  
print(eigenvalues)  
print("\nEigenvectors:")  
print(eigenvectors)  
print("\n")
```

```
return principalDf
```

```
n_components = 1  
result = apply_pca(df2, n_components)  
print(result)
```

OUTPUT

```
Centered Data:  
   0    1  
0 -4.0 -0.5  
1  0.0 -4.5  
2  5.0 -3.5  
3 -1.0  8.5  
  
Covariance Matrix:  
[[14. -8.]  
 [-8. 35.]]  
  
Eigenvalues:  
[37.70037878]  
  
Eigenvectors:  
[[-0.31981892  0.94747869]]  
  
New Values  
0    0.805536  
1   -4.263654  
2   -4.915270  
3    8.373388
```

Conclusion : PCA, a powerful dimensionality reduction technique, simplifies large datasets by transforming variables into a smaller set while retaining essential information. It can be used for summarizing complex datasets, uncovering relationships between variables, and simplifying data analysis processes effectively.