Implementing Principal Component Analysis from scratch with Python in OOP

Principial Component Analysis (PCA) is an unsupervised machine learning method that is often used to reduce the dimentionsality of the dataset by transforming a large set into a lower dimensional set that still contains most of the information of the large set.

PCA finds a new set of dimensions such that all the dimensions are orthogonal (and hence linearly independent) and ranked according to the variance of data along them.

- The transformed features should be linearly independent
- Dimensionality can be reduced by taking only the dimensions with the highest importance
- Dimensions should minimize projection error

def transform(self,X):

• Projected points should have maximum spread (i.e., maximum variance)

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In [4]: import numpy as np
        import matplotlib.pyplot as plt
        from sklearn import datasets
In [5]: class PCA:
            def __init__(self, n_components):
                self.n_components = n_components
                self.components = None
                self.mean = None
            def fit(self, X): # only need X, no class labels because unsupervised
                # Mean centering
                self.mean = np.mean(X, axis=0)
                X = X - self.mean
                # Covariance, function needs samples as columns
                cov = np.cov(X.T)
                # Calculate eigenvectors and eigenvalues
                eigenvectors, eigenvalues = np.linalg.eig(cov)
                # Transpose eigenvectors, v = [:,i] column vector
                eigenvectors = eigenvectors.T
                # Sort the eigenvectors according to eigenvalues
                indxs = np.argsort(eigenvalues)[::-1] # in decreasing order
                eigenvalues = eigenvalues[indxs]
                eigenvectors = eigenvectors[indxs]
                # Only save first k components
                self.components = eigenvectors[:self.n_components]
```

```
X = X - self.mean
# Project the data
return np.dot(X, self.components.T)
```

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In [7]: # Testing
        if __name__ == "__main__":
            # Import data
            data = datasets.load_iris()
            X = data.data
            y = data.target
            # Project the data onto the 2 primary principal components
            pca = PCA(2)
            pca.fit(X)
            X_projected = pca.transform(X)
            print(f"Shape of X: {X.shape}")
            print(f"Shape of transformed X: {X_projected.shape}")
            # Extract first 2 dimensions of projected data and plot
            x1 = X_projected[:,0]
            x2 = X projected[:,1]
            plt.scatter(x1, x2, c=y, edgecolor="none", alpha=0.8, cmap=plt.cm.get_cm
            plt.xlabel("Principal Component 1")
            plt.ylabel("Principal Component 2")
            plt.colorbar()
            plt.show()
       Shape of X: (150, 4)
```

Shape of transformed X: (150, 2)

/var/folders/cx/jsmdsr392b16bs81k3s1s4n00000gn/T/ipykernel_89303/2975043885. py:20: MatplotlibDeprecationWarning: The get_cmap function was deprecated in Matplotlib 3.7 and will be removed in 3.11. Use ``matplotlib.colormaps[name] `` or ``matplotlib.colormaps.get_cmap()`` or ``pyplot.get_cmap()`` instead. plt.scatter(x1, x2, c=y, edgecolor="none", alpha=0.8, cmap=plt.cm.get_cmap ("viridis",3))

