# dim red

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TD1 Machine Learning - MMVAI

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Import

```
[]: import numpy as np
import matplotlib.pyplot as plt

from sklearn.datasets import make_moons
from sklearn.metrics import accuracy_score
from sklearn.model_selection import train_test_split
```

#### Part 1: PCA

The goal of this TD is to understand the difference between PCA and Kernel based PCA

To perform a PCA based dimension reduction, we need:

Compute the covariance matrix C of the original data X.

Perform the eigendecomposition of the computed matrix.

Sort the eigenvalues in decreasing order.

Construct the projection matrix W with the first (k) eigenvalues.

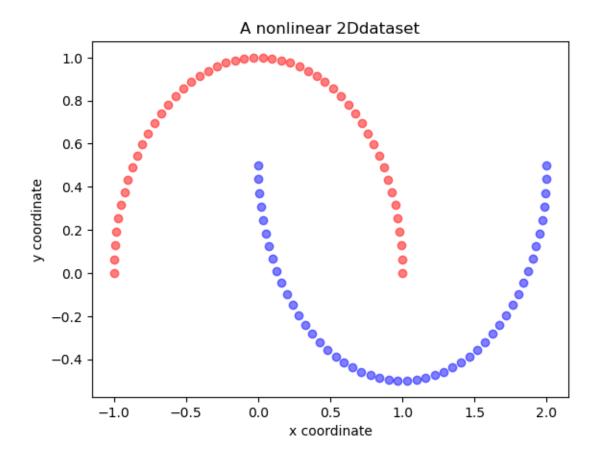
Transform the data into the projected space  $X_{pca} = W^T X$ .

Generate the data

```
[]: X, y = make_moons(n_samples=100, random_state=123)

plt.scatter(X[y==0, 0], X[y==0, 1], color='red', alpha=0.5)
plt.scatter(X[y==1, 0], X[y==1, 1], color='blue', alpha=0.5)

plt.title('A nonlinear 2Ddataset')
plt.ylabel('y coordinate')
plt.xlabel('x coordinate')
y = y.reshape((y.shape[0], 1))
```



## TODO

Perform PCA based reduction on this data

Keep only the first leading eigen value

Plot the resulting points

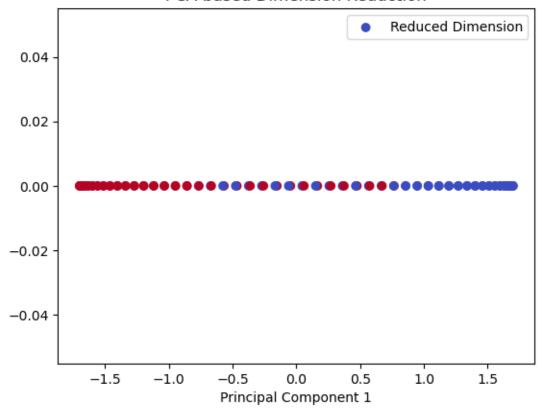
```
[]: # Step 1: Compute the covariance matrix C
X_centered = (X - np.mean(X, axis=0)) / np.std(X, axis = 0)
C = np.cov(X_centered, rowvar=False)

# Step 2: Perform eigendecomposition
eigenvalues, eigenvectors = np.linalg.eigh(C)

# Step 3: Sort eigenvalues and eigenvectors
sorted_indices = np.argsort(eigenvalues)[::-1]
eigenvalues = eigenvalues[sorted_indices]
eigenvectors = eigenvectors[:, sorted_indices]

# Step 4: Construct projection matrix W using the first leading eigenvector
k = 1
```





### TODO

Perform linear classification using your favorite linear classifier

Implementation of the Linear Classifier

```
[]: def initialisation(X):
    W = np.random.randn(X.shape[1], 1)
```

```
b = np.random.randn(1)
    return (W, b)
def model(X, W, b):
    Z = X.dot(W + b)
    A = 1 / (1 + np.exp(-Z))
    return A
def log loss(A, y):
   return 1 / len(y) * np.sum(-y * np.log(A) - (1 - y) * np.log(1 - A))
def gradients(A, X, y):
    dW = 1 / len(y) * np.dot(X.T, A - y)
    db = 1 / len(y) * np.sum(A - y)
    return (dW, db)
def update(dW, db, W, b, learning_rate):
    W -= learning_rate * dW
    b -= learning_rate * db
    return (W, b)
def predict(X, W, b):
    A = model(X, W, b)
    return A >= 0.5
def perceptron(X, y, learning_rate = 0.1, n_iter = 100, loss = False):
    W, b = initialisation(X)
    Loss = []
    for i in range(n_iter):
        A = model(X, W, b)
        Loss.append(log_loss(A, y))
        dW, db = gradients(A, X, y)
        W, b = update(dW, db, W, b, learning_rate)
    y_pred = predict(X, W, b)
    if loss:
        return (W, b, Loss)
    return (W, b)
```

Training of the perceptron

```
[]: # Split the data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X_pca, y, test_size=0.2, □ → random_state=42)

# Create a linear classifier (Perceptron)
```

```
clf = perceptron(X_pca, y, loss = True)
```

Extract the Parameters

```
[ ]: W, b, Loss = clf
```

Accuracy of the Perceptron

```
[]: # Make predictions on the test set
y_pred = predict(X_test, W, b)

# Evaluate accuracy
accuracy = accuracy_score(y_test, y_pred)
print(f"Accuracy: {accuracy * 100:.2f}%")
```

Accuracy: 80.00%

Plot the result

```
[]: plt.figure()
  plt.scatter(X_pca, y, c=y, cmap="coolwarm")
  plt.xlabel('X_pca')
  plt.title('PCA-based Dimension Reduction')
```

[]: Text(0.5, 1.0, 'PCA-based Dimension Reduction')



0.0

X\_pca

0.5

1.0

1.5

PCA-based Dimension Reduction

```
[]: # Learning Curve
plt.figure()
plt.plot(Loss)
plt.ylabel("loss")
plt.xlabel("n_iter")
plt.title("Learning Curve")
```

-0.5

```
[]: Text(0.5, 1.0, 'Learning Curve')
```

1.0

0.8

0.6

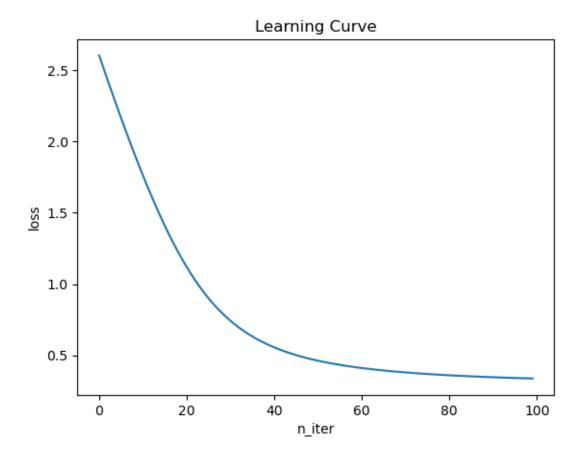
0.4

0.2

0.0

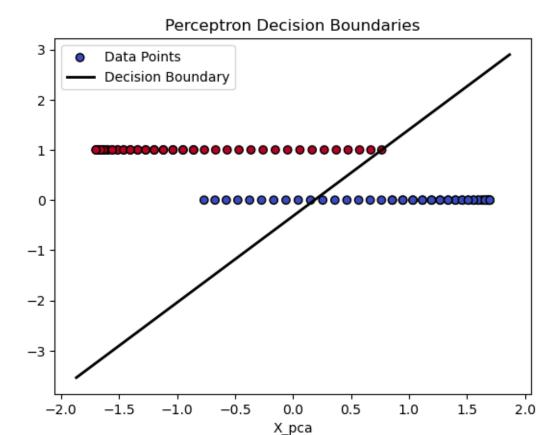
-1.5

-1.0



```
[]: # Plot decision boundaries
     plt.figure()
     # Scatter plot of the data points
     plt.scatter(X_pca, y, c=y, cmap="coolwarm", edgecolors='k', marker='o', u
      ⇔label='Data Points')
     # Plot decision boundaries
     ax = plt.gca()
     xlim = ax.get_xlim()
     xx = np.linspace(xlim[0], xlim[1], 100)
     yy = (-W[0] * xx - b)
    plt.plot(xx, yy, color='black', linestyle='-', linewidth=2, label='Decision_

→Boundary')
    plt.xlabel('X_pca')
     plt.title('Perceptron Decision Boundaries')
     plt.legend()
     plt.show()
```



Part 2: Kernel PCA

Kernel functions and the kernel trick:

The basic idea to deal with inseparable data using linear classifiers is to project it onto a higher dimensional space where it becomes linearly separable. To do so, we:

Compute the kernel matrix K using RBF kernel for instance  $exp(-\gamma||x_i-x_j||^2)$  (exp(-g \* abs(x\_i - x\_j) ^2))

Eigendecompose of the kernel matrix.  ${\bf K}$  .

to transform the data into the projected space.

### TODO

Perform KPCA based reduction on this data

Keep only the first leading eigen value

Plot the resulting points

Function for KPCA

```
[]: def rbf_kernel(X, gamma):
         # Extract the number of samples
         n = X.shape[0]
         # Define the different matrix
         xxt = X @ X.T
         A = X @ X.T
         B = np.repeat(np.diag(xxt), n).reshape(n, n)
         # The return is an n x n matrix
         return np.exp(-(B.T - 2*A + B) * gamma)
     def kernel_pca(X, n_components, gamma):
         # Compute the RBF kernel matrix
         K = rbf_kernel(X, gamma)
         # Center the kernel matrix
         n_samples = K.shape[0]
         one_n = np.ones((n_samples, n_samples)) / n_samples
         K_centered = K - one_n @ K - K @ one_n + one_n @ K @ one_n
         # Perform eigendecomposition
         eigenvalues, eigenvectors = np.linalg.eigh(K_centered)
         # Sort eigenvalues and corresponding eigenvectors in descending order
         sorted indices = np.argsort(eigenvalues)[::-1]
         eigenvalues = eigenvalues[sorted_indices]
         eigenvectors = eigenvectors[:, sorted_indices]
         # Keep only the first leading eigenvalues and corresponding eigenvectors
         K_pca_components = eigenvectors[:, :n_components]
         # Project the data into the reduced space
         X_kpca = K_pca_components.T @ K_centered.T
         return X_kpca.T
```

```
[]: # Set kernel parameters
gamma_value = 0.5
n_components = 3

# Perform Kernel PCA
X_kpca = kernel_pca(X, n_components, gamma_value)
```

#### TODO

Perform linear classification using your favorite linear classifier

Training the model

Extract the parameters

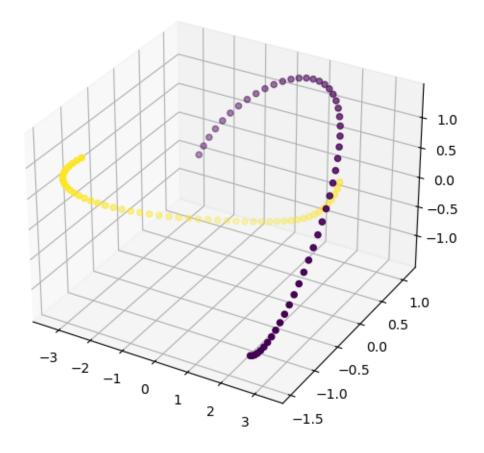
```
[ ]: W_kpca, b_kpca, Loss_kpca = clf_kpca
```

Compute the accuracy

```
[]: # Make predictions on the test set
y_pred_kpca = predict(X_test_kpca, W_kpca, b_kpca)

# Evaluate accuracy
accuracy = accuracy_score(y_test_kpca, y_pred_kpca)
print(f"Accuracy: {accuracy * 100:.2f}%")
```

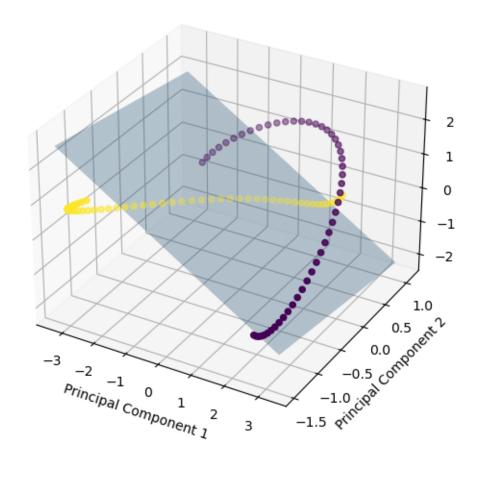
Accuracy: 75.00%



Plot

```
ax.set_zlabel('Principal Component 3')
ax.set_title('Kernel PCA Projection with Decision Boundary in 3D')
plt.show()
```

# Kernel PCA Projection with Decision Boundary in 3D



```
[]: plt.figure()
  plt.plot(Loss_kpca)
  plt.ylabel("loss")
  plt.xlabel("n_iter")
  plt.title("Learning Curve")
```

[]: Text(0.5, 1.0, 'Learning Curve')

