

Lecture 14: Dimensionality Reduction – Kernel PCA

Textbook: Chapter 5

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Kernel PCA

- In real applications, we may need to deal with nonlinear problems.
- For nonlinear problems, linear transformation techniques for dimensionality reduction (e.g., PCA and LDA) may not be the best choice.
- Kernelized version of PCA (or KPCA): transform data that is not linearly separable onto a new, higher-dimensional subspace that is suitable for linear classifiers. Or, it extends conventional principal component analysis (PCA) to a high dimensional feature space using the “kernel trick”.

Mapping function

- A nonlinear mapping function $\phi: \mathbb{R}^d \rightarrow \mathbb{R}^k (k \gg d)$
- ϕ can be thought of as a function that creates **nonlinear combinations** of the features in the original d -dimensional dataset onto a larger k -dimensional feature space.
E.g., $\mathbf{x} = [x_1, x_2]$ can be transformed to $\mathbf{z} = [x_1^2, \sqrt{2x_1x_2}, x_2^2]$.
- New feature space has k dimensions. It is denoted as \mathcal{Z} space or Φ space.

How to use Kernel PCA

- Perform a **nonlinear mapping** via kernel PCA that transforms data to a higher dimensional space.
- Use **standard PCA** in this higher dimensional space to project the data back onto a lower-dimensional space where the samples can be separated by a **linear classifier**.
 - This lower-dimensional space (after running PCA) is different from the original X space.

Procedure

- PCA analysis: calculate covariance matrix

$$\Sigma = \frac{1}{n} \sum_{i=1}^n (\mathbf{x}^{(i)}) (\mathbf{x}^{(i)})^T$$

- In the \mathcal{Z} space

$$\Sigma = \frac{1}{n} \sum_{i=1}^n \phi(\mathbf{x}^{(i)}) \phi(\mathbf{x}^{(i)})^T$$

- Then, we need to obtain the eigen vector of Σ . This process is computationally very expensive. (Detailed analysis see the extra slides provided.)
- This is where we use the **kernel trick**. Using the kernel trick, we can compute the similarity between two high-dimension feature vectors in the original feature space.

Kernel trick

- Using the kernel trick, we can avoid calculating the pairwise dot products of the samples \mathbf{x} under ϕ . Instead, we use a kernel function κ

$$\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \phi(\mathbf{x}^{(i)})^T \phi(\mathbf{x}^{(j)})$$

- Different types of kernel functions

- Polynomial kernel: $\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \left((\mathbf{x}^{(i)})^T \mathbf{x}^{(j)} + \theta \right)^p$
 - θ is a threshold. p is the power specified by the user.
- Hyperbolic tangent (sigmoid) kernel: $\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \tanh \left(\eta (\mathbf{x}^{(i)})^T \mathbf{x}^{(j)} + \theta \right)$
- **Radial Basis Function (or Gaussian kernel):** $\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \exp \left(-\frac{\|\mathbf{x}^{(i)} - \mathbf{x}^{(j)}\|^2}{2\sigma^2} \right)$ or $\kappa(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \exp(-\gamma \|\mathbf{x}^{(i)} - \mathbf{x}^{(j)}\|^2)$ where $\gamma = \frac{1}{2\sigma^2}$

Scikit-learn kernel PCA

- **KernelPCA** function in **sklearn.decomposition** module
- Parameter “n_components”: Number of components. If None, all non-zero components are kept.
- Parameter “kernel”.

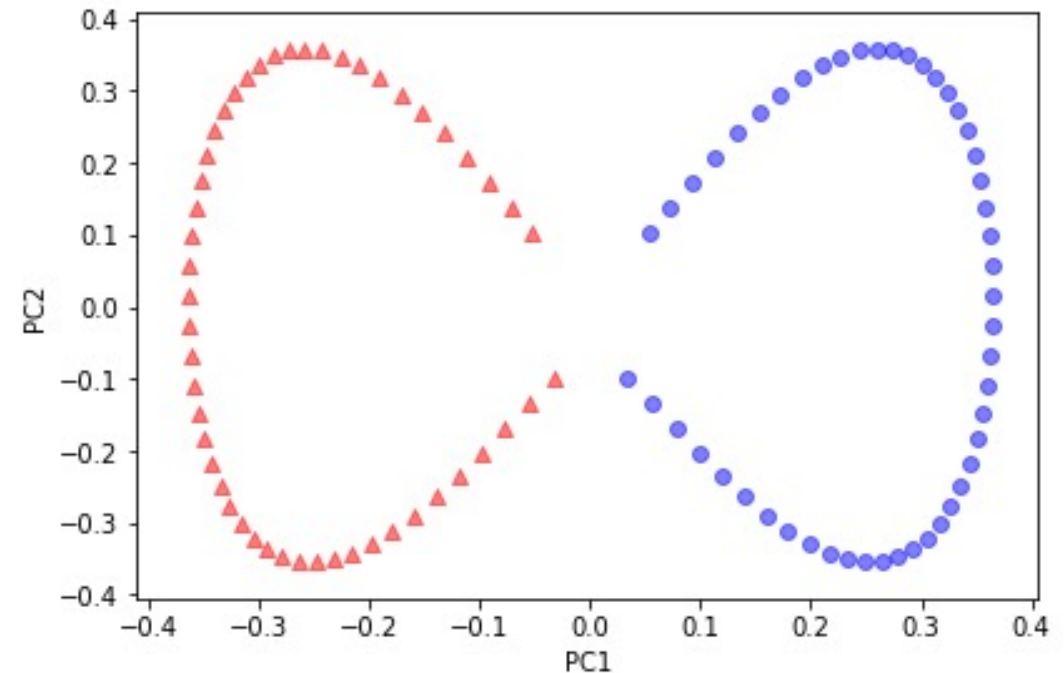
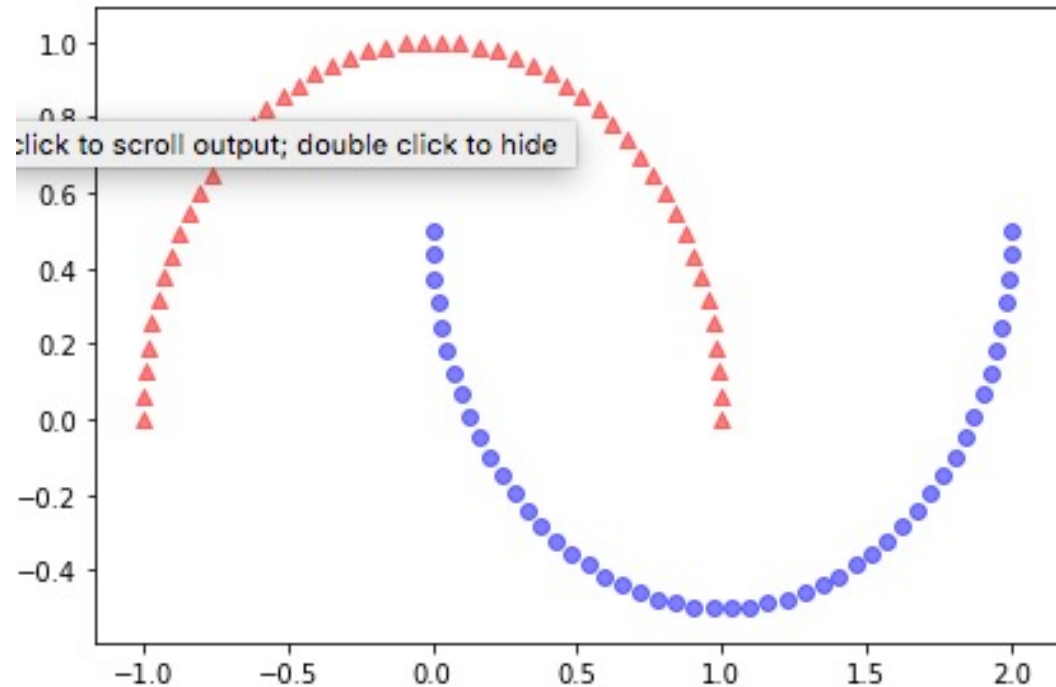
```
from sklearn.decomposition import KernelPCA
```

```
scikit_kpca = KernelPCA(n_components=2, kernel='rbf', gamma=15)  
X_kpca = scikit_kpca.fit_transform(X)
```

Example 1

```
from sklearn.datasets import make_moons
X, y = make_moons(n_samples=100, random_state=123)

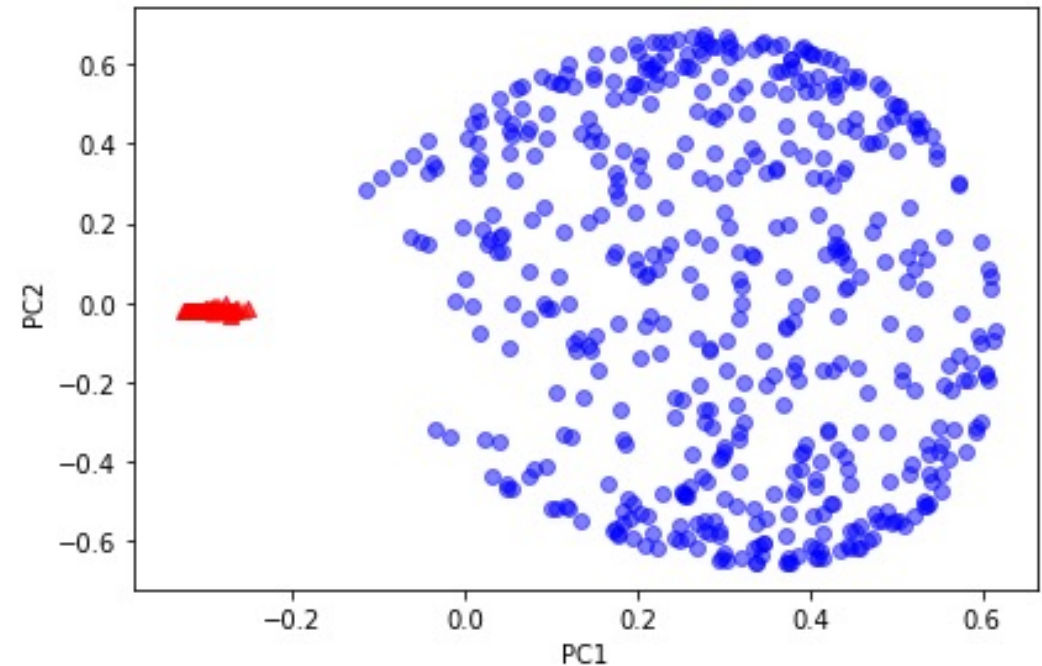
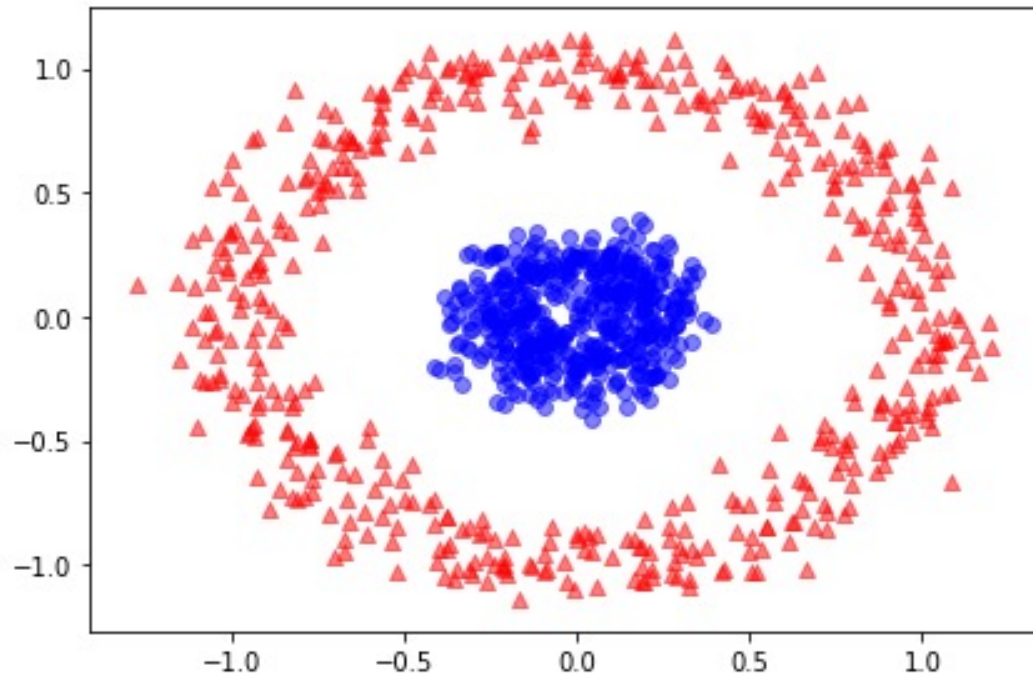
# plotting X with different colors for different class labels
# X_kpca = apply KPCA operation on X with n_components=2, kernel='rbf', gamma=15
# plotting X_kpca with different colors for different class labels
```



Example 2

```
from sklearn.datasets import make_circles
X, y = make_circles(n_samples=1000, random_state=123, noise=0.1, factor=0.2)

# plotting X with different colors for different class labels
# X_skernpca = apply KPCA operation on X with n_components=2, kernel='rbf', gamma=15
# plotting X_skernpca with different colors for different class labels
```



Use Kernel PCA

```
from sklearn.decomposition import KernelPCA
```

```
X, y = make_circles(n_samples=1000, random_state=123, noise=0.1, factor=0.2)
```

```
X_train, X_test, y_train, y_test = train_test_split(X,y,test_size=0.3,stratify=y,random_state=0)
```

```
kpca = KernelPCA(n_components=2, kernel='rbf', gamma=15)
```

```
X_train_kpca = kpca.fit_transform(X_train)
```

```
X_test_kpca = kpca.transform(X_test)
```

```
ppn1 = Perceptron(eta0=0.1, random_state=1)
```

```
ppn1.fit(X_train, y_train)
```

```
y_pred1 = ppn1.predict(X_test)
```

```
print("accuracy (no kpca) = ", sum(y_test==y_pred1)/y_test.shape[0])
```

```
ppn2 = Perceptron(eta0=0.1, random_state=1)
```

```
ppn2.fit(X_train_kpca, y_train)
```

```
y_pred2 = ppn2.predict(X_test_kpca)
```

```
print("accuracy (with kpca) ", sum(y_test==y_pred2)/y_test.shape[0])
```

```
accuracy (no kpca) = 0.6433333333333333
```

```
accuracy (with kpca) 1.0
```

Discussions

- Run linear classification algorithms after Kernel PCA process
- Transform the data **only once**, but can utilize it multiple times in different classification algorithms.