

## Lecture 14

### Nonlinear PCA and Manifold Learning

Reading: Izenman 2008: Chapter 16.1-16.2; 16.5-16.6

DSA 8070 Multivariate Analysis

Whitney Huang  
Clemson University

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#### Agenda



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#### ① Nonlinear PCA



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#### Motivation

- High-dimensional data often lie on low-dimensional manifolds
- Linear methods like PCA may fail to capture intrinsic geometry
- Goal: find lower-dimensional embedding that preserves structure



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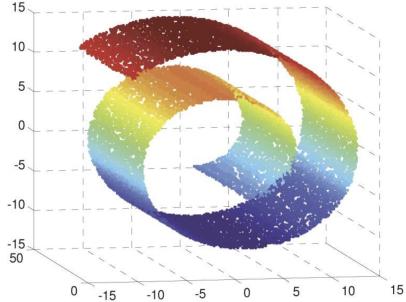
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## Swiss Roll Example: Motivation for Manifold Learning

- A classic synthetic example for nonlinear dimensionality reduction.
- Data lie on a 2-dimensional manifold smoothly embedded in  $\mathbb{R}^3$ .
- Structure: a flat rectangle “rolled up” into a spiral surface.



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## Learning Objectives

By the end of this lecture, you will be able to:

- Understand the motivation behind manifold learning
- Describe key nonlinear dimensionality reduction techniques
- Implement and interpret, [Nonlinear PCA methods](#), [Isomap](#), [LLE](#), and [t-SNE](#) in R
- Compare manifold methods to [Linear PCA](#)



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## Why Nonlinear PCA?

- Classical PCA assumes that the main structure of the data lies in a [linear subspace](#) of  $\mathbb{R}^p$ .
- For many data sets, structure is [nonlinear](#):
  - Curved manifolds (e.g., Swiss roll)
  - Nonlinear functional relationships (e.g., quadratic curves)
- Idea: generalize PCA so that we still
  - Reduce dimension
  - Capture major modes of variation
  - Allow [nonlinear](#) transformations of the original variables
- There is no single unique “nonlinear PCA” – different methods generalize different properties of PCA



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### Polynomial PCA: Basic Idea

- Suppose  $X \in \mathbb{R}^p$  is the original data vector
- Construct an **expanded feature vector**  $X'$  by adding polynomial terms, e.g. for quadratic PCA:  
$$X' = (X_1, \dots, X_p, X_1^2, \dots, X_p^2, X_1 X_2, \dots, X_{p-1} X_p)$$
- Apply **ordinary PCA** to the expanded data  $X'$ :
  - Center  $X'$
  - Compute covariance matrix of  $X'$
  - Eigenvalues/eigenvectors give “principal components” in the polynomial feature space
- If the true relationship is approximately polynomial, small eigenvalues can correspond to **approximate nonlinear constraints**



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### Polynomial PCA: Comments

- Advantages:**
  - Simple conceptually: just augment features and run PCA
  - Can capture simple nonlinear structure (quadratic, cubic, etc.)
- Disadvantages:**
  - Dimension of  $X'$  grows quickly with polynomial degree  $d$  and number of variables  $p$
  - Risk of overfitting and numerical instability



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### Kernel PCA: PCA in Feature Space

- Instead of explicitly constructing polynomial (or other) features, map  $x \in \mathbb{R}^p$  to a feature space  $H$ :  
$$\Phi : \mathbb{R}^p \rightarrow H,$$
 and perform **linear PCA** on  $\Phi(x_1), \dots, \Phi(x_n)$
- We never need  $\Phi(x)$  explicitly, only inner products  
$$K(x_i, x_j) = \langle \Phi(x_i), \Phi(x_j) \rangle_H$$
- A **kernel**  $K(x_i, x_j)$  replaces the explicit feature map
- Kernel PCA = PCA in feature space defined by  $K$



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## Kernel PCA: Algorithm Sketch

- Choose a kernel  $K$ , e.g.

$$K(x, y) = \exp\left(-\frac{\|x - y\|^2}{2\sigma^2}\right) \quad (\text{Gaussian / RBF kernel}).$$

- Form the  $n \times n$  Gram matrix  $K_{ij} = K(x_i, x_j)$

- Center  $K$  in feature space:

$$\tilde{K} = K - \mathbf{1}K/n - K\mathbf{1}/n + \mathbf{1}\mathbf{1}^T/n^2,$$

where  $\mathbf{1}$  is the all-ones matrix

- Compute eigenvalues/eigenvectors of  $\tilde{K}$ :

$$\tilde{K}\alpha^{(\ell)} = n\lambda_\ell\alpha^{(\ell)}$$

- The coordinates of data point  $x_i$  on kernel PC  $\ell$  are proportional to the  $i$ th component of  $\alpha^{(\ell)}$



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## Kernel PCA: Interpretation

- Kernel PCA finds directions of maximal variance in a **nonlinear feature space**.

- For suitable kernels, kernel PCA can:

- capture curved manifolds,
- partially “unfold” structures like the Swiss roll,
- behave similarly to metric MDS based on a kernel-induced distance.

- The kernel choice (e.g., bandwidth  $\sigma$  in RBF kernel) controls the balance between:

- local vs. global structure,
- smooth vs. noisy embeddings.



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## Kernel PCA vs Linear PCA (Swiss Roll)

Compare 2D linear PCA and RBF kernel PCA on the same Swiss roll dataset (coloured by height  $h$ )

### Linear PCA

- Projects data onto a best linear 2D subspace
- Swiss roll remains curved and “folded”
- Colour gradient (in  $h$ ) is not monotone along any axis

### Kernel PCA (RBF)

- Applies PCA in a nonlinear feature space defined by the kernel
- Effectively unrolls the manifold into 2D
- Colour gradient in  $h$  varies smoothly across the embedding

**Key message:** Linear PCA is limited to global linear structure, whereas kernel PCA can recover nonlinear manifold structure when the kernel is chosen appropriately



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## Takeaways: Nonlinear PCA

- Linear PCA can fail when the data lie on a nonlinear manifold
- Polynomial PCA:
  - PCA on explicitly expanded polynomial features
  - Simple but can be high-dimensional
- Kernel PCA:
  - PCA in an implicit feature space defined by a kernel
  - More flexible and scalable than explicit polynomial expansion



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## Manifold Learning: Core Idea

- Data points lie on a nonlinear subspace (manifold)
- Manifold learning tries to “unroll” this space
- Preserves local or global geometric relationships



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## Isomap

- Combines classical MDS with geodesic distance
- Steps:
  - Construct neighborhood graph (k-nearest or epsilon)
  - Compute shortest path distances (Floyd-Warshall)
  - Apply MDS on the geodesic distance matrix
- Captures global geometry



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## Locally Linear Embedding (LLE)

- Preserves local linear relationships between neighbors
- Steps:
  - ① Identify k-nearest neighbors
  - ② Compute weights to reconstruct each point from neighbors
  - ③ Find embedding that preserves weights
- Sensitive to noise and parameter choice



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## t-SNE (t-distributed Stochastic Neighbor Embedding)

- Probabilistic approach to preserve pairwise similarities
- Converts distances to probabilities in high and low dimensions
- Optimizes Kullback-Leibler divergence between distributions
- Good for visualization (not true metric embedding)



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## UMAP (Uniform Manifold Approximation and Projection)

- Balances local and global structure preservation
- Faster than t-SNE and scales better
- Based on fuzzy simplicial sets and topological data analysis



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## Comparison with PCA

- PCA is linear and global
- Manifold methods are nonlinear, and often local
- Use PCA when interpretability and simplicity matter
- Use t-SNE, Isomap, or UMAP for visualizing structure



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## Nonlinear PCA vs. Manifold Methods

- Polynomial PCA and kernel PCA:
  - Extend PCA using richer feature spaces
  - Remain **variance-based** methods
- Methods like Isomap, LLE:
  - Focus on preserving **geodesic distances** or **local neighborhoods**
  - Explicitly target manifold geometry



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## Summary

- Manifold learning methods uncover low-dimensional structures
- Isomap and LLE preserve distances or local geometry
- t-SNE and UMAP are widely used for high-dimensional data visualization



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